Linux-Bioinformatica

October 1, 2018

1 Comandos Básicos de Linux

1.1 Conceptos

- La mayor parte de comandos que va a utilizar son para el manejo de archivos
- Los archivos se ubican en **directorios** o carpetas.
- Cuando inicia seción siempre va a estar en su directorio raíz (su cuenta)

```
In [34]: ## Ejemplo
         pwd

/home/lg
```

- En cada directorio siempre están presenta dos directorios:
    - El directorio padre: **..**
    - El directorio actual: **.**

```
In [43]: ## Ejemplo
         mkdir p1
         cd p1
         ll

total 8.0K
drwxrwxr-x  2 4.0K 10-01 20:46 .
drwxr-xr-x 47 4.0K 10-01 20:46 ..
```

```
In [44]: ## Ejemplo
         mkdir taller1
         ll

total 12K
drwxrwxr-x  3 4.0K 10-01 20:48 .
drwxr-xr-x 47 4.0K 10-01 20:46 ..
drwxrwxr-x  2 4.0K 10-01 20:48 taller1
```

2 Resumen de Comandos

2.1 Manejo de directorios

- pwd : Muestra el directorio actual donde estoy ubicado
- mkdir : Crea (make dir) un directorio
- cd : Se cambia (change dir) a un directorio
- rmdir : Borra un directorio (que esté vacio)

2.2 Manejo de archivos

- ls : Lista (list) los archivos y directorios
- ll : Lista (long list) de forma detallada (permisos, tamaño, hora)
- cp : Copia (copy) un archivo a otro lugar o hace una copia del archivo
- mv : Mueve (move) un archivo a otro lugar o le cambia de nombre
- rm : Borra (remove) un archivo
- touch: Crea un archivo vacio

2.3 Descompresión archivos

- gunzip : Descomprime **archivo** con extension **.gz**
- tar : Descomprime **archivo** con extension **.tgz** o **.tar.gz**

2.4 Descarga de archivos

- wget : Descarga un archivo de un URL (http:/....)

2.5 Pantalla/Visualización de archivos

- clear : Limpia la pantalla de comandos o resultados anteriores
- cat : Muestra el contenido de un archivo
- head : Muestra las primeras 10 líneas de un archivo
- tail : Muestra las últimas 10 liínes de un archivo

2.6 Filtrado de archivos

- wget : Filtra un archivo dado un patron de búsqueda (muestra las líneas que solo contengan ese patron.

2.7 Editores tipo consola

- nano : Editor de texto modo consola

3 Uso y Ejemplos de los comandos

3.1 pwd : Muestra el directorio actual donde estoy ubicado

```
In [45]: ## Ejemplo
         pwd
```

```
/home/lg/p1
```

3.2 mkdir : Crea (make dir) un directorio

- USO: mkdir < nombreDirectorioNuevo >

```
In [46]: ## Ejemplo
         mkdir taller2
         ll

total 16K
drwxrwxr-x  4 4.0K 10-01 20:50 .
drwxr-xr-x 47 4.0K 10-01 20:46 ..
drwxrwxr-x  2 4.0K 10-01 20:48 taller1
drwxrwxr-x  2 4.0K 10-01 20:50 taller2
```

3.3 cd : Se cambia (change dir) a un directorio

- USO: cd < nombreDirectorio >
- Si se ejecuta solito se cambia a mi directorio raíz (directorio inicial)

```
In [47]: ## Ejemplo
         pwd

/home/lg/p1


In [48]: cd
         pwd

/home/lg
```

- Si se agrega un nombre de directorio se ubica en ese directorio

```
In [56]: ## Ejemplo
         cd p1
         pwd

/home/lg/p1
```

3.4 rmdir : Borra un directorio (que esté vacio)

```
In [57]: ## Ejemplo:
         ll
```

```
total 16K
drwxrwxr-x  4 4.0K 10-01 20:50 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
drwxrwxr-x  2 4.0K 10-01 20:48 taller1
drwxrwxr-x  2 4.0K 10-01 20:50 taller2
```

```
In [58]: rmdir taller1
         ll

total 12K
drwxrwxr-x  3 4.0K 10-01 20:58 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
drwxrwxr-x  2 4.0K 10-01 20:50 taller2
```

3.5 ls y ll: Lista (list) los archivos y directorios

```
In [59]: ## Ejemplo
         ls

.  ..  taller2
```

```
In [60]: ll

total 12K
drwxrwxr-x  3 4.0K 10-01 20:58 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
drwxrwxr-x  2 4.0K 10-01 20:50 taller2
```

3.6 Manejo de archivos

- cp : Copia (copy) un archivo a otro lugar o hace una copia del archivo
- mv : Mueve (move) un archivo a otro lugar o le cambia de nombre
- rm : Borra (remove) un archivo

3.6.1 cp : Copia (copy) un archivo a otro lugar o hace una copia del archivo

- USO: cp < archivoOrigen > < archivoDestino >

```
In [61]: ## Ejemplo: crea un archivo "hola.txt" vacio
         touch hola.txt
         ll

total 12K
drwxrwxr-x  3 4.0K 10-01 21:05 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
-rw-rw-r--  1    0 10-01 21:05 hola.txt
drwxrwxr-x  2 4.0K 10-01 20:50 taller2
```

```
In [62]: ## Hace una copia de hola.txt
         cp hola.txt mundo.txt
         ll

total 12K
drwxrwxr-x  3 4.0K 10-01 21:06 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
-rw-rw-r--  1    0 10-01 21:05 hola.txt
-rw-rw-r--  1    0 10-01 21:06 mundo.txt
drwxrwxr-x  2 4.0K 10-01 20:50 taller2


In [64]: ## Hace una copia en el dir taller2
         cp hola.txt taller2

In [65]: ## Lista lo que hay en taller2
         ll taller2

total 8.0K
drwxrwxr-x 2 4.0K 10-01 21:07 .
drwxrwxr-x 3 4.0K 10-01 21:06 ..
-rw-rw-r-- 1    0 10-01 21:07 hola.txt
```

3.6.2   mv : Mueve (move) un archivo a otro lugar o le cambia de nombre

- USO1: mv < archivoOrigen> < directorioDestino >
- USO2: mv < archivoOrigen> < archivoDestino >

```
In [67]: ## Ejemplo
         ll

total 12K
drwxrwxr-x  3 4.0K 10-01 21:06 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
-rw-rw-r--  1    0 10-01 21:05 hola.txt
-rw-rw-r--  1    0 10-01 21:06 mundo.txt
drwxrwxr-x  2 4.0K 10-01 21:07 taller2


In [68]: mkdir taller3
         ll

total 16K
drwxrwxr-x  4 4.0K 10-01 21:11 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
-rw-rw-r--  1    0 10-01 21:05 hola.txt
-rw-rw-r--  1    0 10-01 21:06 mundo.txt
drwxrwxr-x  2 4.0K 10-01 21:07 taller2
drwxrwxr-x  2 4.0K 10-01 21:11 taller3
```

```
In [69]: ## Mueve todos los archivos que terminan en .txt al dir taller3
         mv *.txt taller3

In [70]: ll

total 16K
drwxrwxr-x  4 4.0K 10-01 21:12 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
drwxrwxr-x  2 4.0K 10-01 21:07 taller2
drwxrwxr-x  2 4.0K 10-01 21:12 taller3


In [71]: ## Lista que hay en taller3
         ll taller3

total 8.0K
drwxrwxr-x 2 4.0K 10-01 21:12 .
drwxrwxr-x 4 4.0K 10-01 21:12 ..
-rw-rw-r-- 1    0 10-01 21:05 hola.txt
-rw-rw-r-- 1    0 10-01 21:06 mundo.txt
```

3.6.3   rm : Borra (remove) un archivo

- USO: rm < nombreArchivo >
- Eliminar el archivo

```
In [73]: ### Ejemplo
         ll

total 16K
drwxrwxr-x  4 4.0K 10-01 21:12 .
drwxr-xr-x 47 4.0K 10-01 20:55 ..
drwxrwxr-x  2 4.0K 10-01 21:07 taller2
drwxrwxr-x  2 4.0K 10-01 21:12 taller3


In [74]: cd taller3
         ll

total 8.0K
drwxrwxr-x 2 4.0K 10-01 21:12 .
drwxrwxr-x 4 4.0K 10-01 21:12 ..
-rw-rw-r-- 1    0 10-01 21:05 hola.txt
-rw-rw-r-- 1    0 10-01 21:06 mundo.txt


In [16]: ## borra hola.txt
         rm hola.txt
         ll
```

```
total 0
-rw-rw-r-- 1 0 10-01 21:06 mundo.txt
```

```
In [17]: ll
```

```
total 0
-rw-rw-r-- 1 0 10-01 21:06 mundo.txt
```

3.7 Pantalla/Visualización de archivos

- clear : Limpia la pantalla de comandos o resultados anteriores
- cat : Muestra el contenido de un archivo

    – USO: cat < nombreArchivo >

- head : Muestra las primeras 10 líneas de un archivo

    – USO: head < nombreArchivo >

- tail : Muestra las últimas 10 liínes de un archivo

    – USO: tail < nombreArchivo >

```
In [18]: ### Ejemplos: copio un archivo a mi directorio y veo que tiene
         cp /tmp/genoma.fna .
```

```
In [19]: ll
```

```
total 4.5M
-rwxrwxr-x 1 4.5M 10-01 21:22 genoma.fna
-rw-rw-r-- 1    0 10-01 21:06 mundo.txt
```

```
In [20]: cat genoma.fna
```

```
>NC_000913.3 Escherichia coli str. K-12 substr. MG1655, complete genome
AGCTTTTCATTCTGACTGCAACGGGCAATATGTCTCTGTGTGGATTAAAAAAAGAGTGTCTGATAGCAGCTTCTGAACTG
GTTACCTGCCGTGAGTAAATTAAAATTTTATTGACTTAGGTCACTAAATACTTTAACCAATATAGGCATAGCGCACAGAC
AGATAAAAATTACAGAGTACACAACATCCATGAAACGCATTAGCACCACCATTACCACCACCATCACCATTACCACAGGT
AACGGTGCGGGCTGACGCGTACAGGAAACACAGAAAAAAGCCCGCACCTGACAGTGCGGGCTTTTTTTTTCGACCAAAGG
TAACGAGGTAACAACCATGCGAGTGTTGAAGTTCGGCGGTACATCAGTGGCAAATGCAGAACGTTTTCTGCGTGTTGCCG
ATATTCTGGAAAGCAATGCCAGGCAGGGGCAGGTGGCCACCGTCCTCTCTGCCCCCGCCAAAATCACCAACCACCTGGTG
GCGATGATTGAAAAAACCATTAGCGGCCAGGATGCTTTACCCAATATCAGCGATGCCGAACGTATTTTTGCCGAACTTTT
GACGGGACTCGCCGCCGCCCAGCCGGGGTTCCCGCTGGCGCAATTGAAAACTTTCGTCGATCAGGAATTTGCCCAAATAA
AACATGTCCTGCATGGCATTAGTTTGTTGGGGCAGTGCCCGGATAGCATCAACGCTGCGCTGATTTGCCGTGGCGAGAAA
ATGTCGATCGCCATTATGGCCGGCGTATTAGAAGCGCGCGGTCACAACGTTACTGTTATCGATCCGGTCGAAAAACTGCT
GGCAGTGGGGCATTACCTCGAATCTACCGTCGATATTGCTGAGTCCACCCGCCGTATTGCGGCAAGCCGCATTCCGGCTG
ATCACATGGTGCTGATGGCAGGTTTCACCGCCGGTAATGAAAAAGGCGAACTGGTGGTGCTTGGACGCAACGGTTCCGAC
TACTCTGCTGCGGTGCTGGCTGCCTGTTTACGCGCCGATTGTTGCGAGATTTGGACGGACGTTGACGGGGTCTATACCTG
```

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CGACCCGCGTCAGGTGCCCGATGCGAGGTTGTTGAAGTCGATGTCCTACCAGGAAGCGATGGAGCTTTCCTACTTCGGCG
CTAAAGTTCTTCACCCCCGCACCATTACCCCCATCGCCCAGTTCCAGATCCCCTTGCCTGATTAAAAATACCGGAAATCCT
CAAGCACCAGGTACGCTCATTGGTGCCAGCCGTGATGAAGACGAATTACCGGTCAAGGGCATTTCCAATCTGAATAACAT
GGCAATGTTCAGCGTTTCTGGTCCGGGGATGAAAGGGATGGTCGGCATGGCGGCGCGCGTCTTTGCAGCGATGTCACGCG
CCCGTATTTCCGTGGTGCTGATTACGCAATCATCTTCCGAATACAGCATCAGTTTCTGCGTTCCACAAAGCGACTGTGTG
CGAGCTGAACGGGCAATGCAGGAAGAGTTCTACCTGGAACTGAAAGAAGGCTTACTGGAGCCGCTGGCAGTGACGGAACG
GCTGGCCATTATCTCGGTGGTAGGTGATGGTATGCGCACCTTGCGTGGGATCTCGGCGAAATTCTTTGCCGCACTGGCCC
GCGCCAATATCAACATTGTCGCCATTGCTCAGGGATCTTCTGAACGCTCAATCTCTGTCGTGGTAAATAACGATGATGCG
ACCACTGGCGTGCGCGTTACTCATCAGATGCTGTTCAATACCGATCAGGTTATCGAAGTGTTTGTGATTGGCGTCGGTGG
CGTTGGCGGTGCGCTGCTGGAGCAACTGAAGCGTCAGCAAAGCTGGCTGAAGAATAAACATATCGACTTACGTGTCTGCG
GTGGTTGCCAACTCGAAGGCTCTGCTCACCAATGTACATGGCCTTAATCTGGAAAACTGGCAGGAAGAACTGGCGCAAGCC
AAAGAGCCGTTTAATCTCGGGCGCTTAATTCGCCTCGTGAAAGAATATCATCTGCTGAACCCGGTCATTGTTGACTGCAC
TTCCAGCCAGGCAGTGGCGGATCAATATGCCGACTTCCTGCGCGAAGGTTTCCACGTTGTCACGCCGAACAAAAAGGCCA
ACACCTCGTCGATGGATTACTACCATCAGTTGCGTTATGCGGCGGAAAAATCGCGGCGTAAATTCCTCTATGACACCAAC
GTTGGGGCTGGATTACCGGTTATTGAGAACCTGCAAAATCTGCTCAATGCAGGTGATGAATTGATGAAGTTCTCCGGCAT
TCTTTCTGGTTCGCTTTCTTATATCTTCGGCAAGTTAGACGAAGGCATGAGTTTCTCCGAGGCGACCACGCTGGCGCGGG
AAATGGGTTATACCGAACCGGACCCGCGAGATGATCTTTCTGGTATGGATGTGGCGCGTAAACTATTGATTCTCGCTCGT
GAAACGGGACGTGAACTGGAGCTGGCGGATATTGAAATTGAACCTGTGCTGCCCGCAGAGTTTAACGCCGAGGGTGATGT
TGCCGCTTTTATGGCGAATCTGTCACAACTCGACGATCTCTTTGCCGCGCGCGTGGCGAAGGCCCGTGATGAAGGAAAAG
TTTTGCGCTATGTTGGCAATATTGATGAAGATGGCGTCTGCCGCGTGAAGATTGCCGAAGTGGATGGTAATGATCCGCTG
TTCAAAGTGAAAAATGGCGAAAACGCCCTGGCCTTCTATAGCCACTATTATCAGCCGCTGCCGTTGGTACTGCGCGGATA
TGGTGCGGGCAATGACGTTACAGCTGCCGGTGTCTTTGCTGATCTGCTACGTACCCTCTCATGGAAGTTAGGAGTCTGAC
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TGATGGTGCATTGCTCGGAGATGTAGTCACGGTTGAGGCGGCAGAGACATTCAGTCTCAACAACCTCGGACGCTTTGCCG
ATAAGCTGCCGTCAGAACCACGGGAAAATATCGTTTATCAGTGCTGGGAGCGTTTTTGCCAGGAACTGGGTAAGCAAATT
CCAGTGGCGATGACCCTGGAAAAGAATATGCCGATCGGTTCGGGCTTAGGCTCCAGTGCCTGTTCGGTGGTCGCGGCGCT
GATGGCGATGAATGAACACTGCGGCAAGCCGCTTAATGACACTCGTTTGCTGGCTTTGATGGGCGAGCTGGAAGGCCGTA
TCTCCGGCAGCATTCATTACGACAACGTGGCACCGTGTTTTCTCGGTGGTATGCAGTTGATGATCGAAGAAAACGACATC
ATCAGCCAGCAAGTGCCAGGGTTTGATGAGTGGCTGTGGGTGCTGGCGTATCCGGGGATTAAAGTCTCGACGGCAGAAGC
CAGGGCTATTTTACCGGCGCAGTATCGCCGCCAGGATTGCATTGCGCACGGGCGACATCTGGCAGGCTTCATTCACGCCT
GCTATTCCCGTCAGCCTGAGCTTGCCGCGAAGCTGATGAAAGATGTTATCGCTGAACCCTACCGTGAACGGTTACTGCCA
GGCTTCCGGCAGGCGCGGCAGGCGGTCGCGGAAATCGGCGCGGTAGCGAGCGGTATCTCCGGCTCCGGCCCGACCTTGTT
CGCTCTGTGTGACAAGCCGGAAACCGCCCAGCGCGTTGCCGACTGGTTGGGTAAGAACTACCTGCAAAATCAGGAAGGTT
TTGTTCATATTTGCCGGCTGGATACGGCGGGCGCACGAGTACTGGAAAACTAAATGAAACTCTACAATCTGAAAGATCAC
AACGAGCAGGTCAGCTTTGCGCAAGCCGTAACCCAGGGGTTGGGCAAAAATCAGGGGCTGTTTTTTCCGCACGACCTGCC
GGAATTCAGCCTGACTGAAATTGATGAGATGCTGAAGCTGGATTTTGTCACCCGCAGTGCGAAGATCCTCTCGGCGTTTA
TTGGTGATGAAATCCCACAGGAAATCCTGGAAGAGCGCGTGCGCGCGGCGTTTGCCTTCCCGGCTCCGGTCGCCAATGTT
GAAAGCGATGTCGGTTGTCTGGAATTGTTCCACGGGCCAACGCTGGCATTTAAAGATTTCGGCGGTCGCTTTATGGCACA
AATGCTGACCCATATTGCGGGTGATAAGCCAGTGACCATTCTGACCGCGACCTCCGGTGATACCGGAGCGGCAGTGGCTC
ATGCTTTCTACGGTTTACCGAATGTGAAAGTGGTTATCCTCTATCCACGAGGCAAAATCAGTCCACTGCAAGAAAAACTG
TTCTGTACATTGGGCGGCAATATCGAAACTGTTGCCATCGACGGCGATTTCGATGCCTGTCAGGCGCTGGTGAAGCAGGC
GTTTGATGATGAAGAACTGAAAGTGGCGCTAGGGTTAAACTCGGCTAACTCGATTAACATCAGCCGTTTGCTGGCGCAGA
TTTGCTACTACTTTGAAGCTGTTGCGCAGCTGCCGCAGGAGACGCGCAACCAGCTGGTTGTCTCGGTGCCAAGCGGAAAC
TTCGGCGATTTGACGGCGGGTCTGCTGGCGAAGTCACTCGGTCTGCCGGTGAAACGTTTTATTGCTGCGACCAACGTGAA
CGATACCGTGCCACGTTTCCTGCACGACGGTCAGTGGTCACCCAAAGCGACTCAGGCGACGTTATCCAACGCGATGGACG
TGAGTCAGCCGAACAACTGGCCGCGTGTGGAAGAGTTGTTCCGCCGCAAAATCTGGCAACTGAAAGAGCTGGGTTATGCA
GCCGTGGATGATGAAACCACGCAACAGACAATGCGTGAGTTAAAAGAACTGGGCTACACTTCGGAGCCGCACGCTGCCGT
AGCTTATCGTGCGCTGCGTGATCAGTTGAATCCAGGCGAATATGGCTTGTTCCTCGGCACCGCGCATCCGGCGAAATTTA
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8

```
AAGAGAGCGTGGAAGCGATTCTCGGTGAAACGTTGGATCTGCCAAAAGAGCTGGCAGAACGTGCTGATTTACCCTTGCTT
TCACATAATCTGCCCGCCGATTTTGCTGCGTTGCGTAAATTGATGATGAATCATCAGTAAAATCTATTCATTATCTCAAT
CAGGCCGGGTTTGCTTTTTATGCAGCCCGGCTTTTTTATGAAGAAATTATGGAGAAAAATGACAGGGAAAAAGGAGAAATT
CTCAATAAATGCGGTAACTTAGAGATTAGGATTGCGGAGAATAACAACCGCCGTTCTCATCGAGTAATCTCCGGATATCG
ACCCATAACGGGCAATGATAAAAGGAGTAACCTGTGAAAAAGATGCAATCTATCGTACTCGCACTTTCCCTGGTTCTGGT
CGCTCCCATGGCAGCACAGGCTGCGGAAATTACGTTAGTCCCGTCAGTAAAATTACAGATAGGCGATCGTGATAATCGTG
GCTATTACTGGGATGGAGGTCACTGGCGCGACCACGGCTGGTGGAAACAACATTATGAATGGCGAGGCAATCGCTGGCAC
CTACACGGACCGCCGCCACCGCCGCGCCACCATAAGAAAGCTCCTCATGATCATCACGGCGGTCATGGTCCAGGCAAACA
TCACCGCTAAATGACAAATGCCGGGTAACAATCCGGCATTCAGCGCCTGATGCGACGCTGGCGCGTCTTATCAGGCCTAC
GTTAATTCTGCAATATATTGAATCTGCATGCTTTTGTAGGCAGGATAAGGCGTTCACGCCGCATCCGGCATTGACTGCAA
ACTTAACGCTGCTCGTAGCGTTTAAACACCAGTTCGCCATTGCTGGAGGAATCTTCATCAAAGAAGTAACCTTCGCTATT
AAAACCAGTCAGTTGCTCTGGTTTGGTCAGCCGATTTTCAATAATGAAACGACTCATCAGACCGCGTGCTTTCTTAGCGT
AGAAGCTGATGATCTTAAATTTGCCGTTCTTCTCATCGAGGAACACCGGCTTGATAATCTCGGCATTCAATTTCTTCGGC
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TCGTCTTCGCTGAAGGTTTCGGCCTGCAAGCCGGTGTAGACATCACCTTTAAACGCCAGAATCGCCTGGCGGGCATTCGC
CGGCGTGAAATCTGGCTGCCAGTCATGAAAGCGAGCGGCGTTGATACCCGCCAGTTTGTCGCTGATGCGCATCAGCGTGC
TAATCTGCGGAGGCGTCAGTTTCCGCGCCTCATGGATCAACTGCTGGGAATTGTCTAACAGCTCCGGCAGCGTATAGCGC
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TATGCCGACTTTAGCAAAAAATGAGAATGAGTTGATCGATAGTTGTGATTACTCCTGCGAAACATCATCCCACGCGTCCG
GAGAAAGCTGGCGACCGATATCCGGATAACGCAATGGATCAAACACCGGGCGCACGCCGAGTTTACGCTGGCGTAGATAA
TCACTGGCAATGGTATGAACCACAGGCGAGAGCAGTAAAATGGCGGTCAAATTGGTAATAGCCATGCAGGCCATTATGAT
ATCTGCCAGTTGCCACATCAGCGGAAGGCTTAGCAAGGTGCCGCCGATGACCGTTGCGAAGGTGCAGATCCGCAAACACC
AGATCGCTTTAGGGTTGTTCAGGCGTAAAAAGAAGAGATTGTTTTCGGCATAAATGTAGTTGGCAACGATGGAGCTGAAG
GCAAACAGAATAACCACAAGGGTAACAAACTCAGCACCCCAGGAACCCATTAGCACCCGCATCGCCTTCTGGATAAGCTG
AATACCTTCCAGCGGCATGTAGGTTGTGCCGTTACCCGCCAGTAATATCAGCATGGCGCTTGCCGTACAGATGACCAGGG
TGTCGATAAAAATGCCAATCATCTGGACAATCCCTTGCGCTGCCGGATGCGGAGGCCAGGACGCCGCTGCCGCTGCCGCG
TTTGGCGTCGAACCCATTCCCGCCTCATTGGAAAACATACTGCGCTGAAAACCGTTAGTAATCGCCTGGCTTAAGGTATA
TCCCGCCGCGCCGCCTGCCGCTTCCTGCCAGCCAAAAGCACTCTCAAAAATAGACCAAATGACGTGGGGAAGTTGCCCGA
TATTCATTACGCAAATTACCAGGCTGGTCAGTACCCAGATTATCGCCATCAACGGGACAAAGCCCTGCATGAGCCGGGCG
ACGCCATGAAGACCGCGAGTGATTGCCAGCAGAGTAAAGACAGCGAGAATAATGCCTGTCACCAGCGGGGGAAAATCAAA
AGAAAAACTCAGGGCGCGGGCAACGGCGTTCGCTTGAACTCCGCTGAAAATTATGCCATAGGCGATGAGCAAAAAGACGG
CGAACAGAACGCCCATCCAGCGCATCCCCAGCCCGCGCGCCATATACCATGCCGGTCCGCCACGAAACTGCCCATTGACG
TCACGTTCTTTATAAAGTTGTGCCAGAGAACATTCGGCAAACGAGGTCGCCATGCCGATAAACGCGGCAACCCACATCCA
AAAGACGGCTCCAGGTCCACCGGCGGTAATAGCCAGCGCAACGCCGGCCAGGTTGCCGCTACCCACGCGCGCCGCAAGAC
TGGTACACAATGACTGAAATGAGGTTAAACCGCCTGGCTGTGGATGAATGCTATTTTTAAGACTTTTGCCAAACTGGCGG
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CCAAAGGACGCTGTTAATGAAGGAGAAAAAATCTGGCATGCATATCCCTCTTATTGCCGGTCGCGATGACTTTCCTGTGT
AAACGTTACCAATTGTTTAAGAAGTATATACGCTACGAGGTACTTGATAACTTCTGCGTAGCATACATGAGGTTTTGTAT
AAAAATGGCGGGCGATATCAACGCAGTGTCAGAAATCCGAAACAGTCTCGCCTGGCGATAACCGTCTTGTCGGCGGTTGC
GCTGACGTTGCGTCGTGATATCATCAGGGCAGACCGGTTACATCCCCCTAACAAGCTGTTTAAAGAGAAATACTATCATG
ACGGACAAATTGACCTCCCTTCGTCAGTACACCACCGTAGTGGCCGACACTGGGGACATCGCGGCAATGAAGCTGTATCA
ACCGCAGGATGCCACAACCAACCCTTCTCTCATTCTTAACGCAGCGCAGATTCCGGAATACCGTAAGTTGATTGATGATG
CTGTCGCCTGGGCGAAACAGCAGAGCAACGATCGCGCGCAGCAGATCGTGGACGCGACCGACAAACTGGCAGTAAATATT
GGTCTGGAAATCCTGAAACTGGTTCCGGGCCGTATCTCAACTGAAGTTGATGCGCGTCTTTCCTATGACACCGAAGCGTC
AATTGCGAAAGCAAAACGCCTGATCAAACTCTACAACGATGCTGGTATTAGCAACGATCGTATTCTGATCAAACTGGCTT
CTACCTGGCAGGGTATCCGTGCTGCAGAACAGCTGGAAAAAGAAGGCATCAACTGTAACCTGACCCTGCTGTTCTCCTTC
```

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GCTCAGGCTCGTGCTTGTGCGGAAGCGGGCGTGTTCCTGATCTCGCCGTTTGTTGGCCGTATTCTTGACTGGTACAAAGC
GAATACCGATAAGAAAGAGTACGCTCCGGCAGAAGATCCGGGCGTGGTTTCTGTATCTGAAATCTACCAGTACTACAAAG
AGCACGGTTATGAAACCGTGGTTATGGGCGCAAGCTTCCGTAACATCGGCGAAATTCTGGAACTGGCAGGCTGCGACCGT
CTGACCATCGCACCGGCACTGCTGAAAGAGCTGGCGGAGAGCGAAGGGGCTATCGAACGTAAACTGTCTTACACCGGCGA
AGTGAAAGCGCGTCCGGCGCGTATCACTGAGTCCGAGTTCCTGTGGCAGCACAACCAGGATCCAATGGCAGTAGATAAAC
TGGCGGAAGGTATCCGTAAGTTTGCTATTGACCAGGAAAAACTGGAAAAAATGATCGGCGATCTGCTGTAATCATTCTTA
GCGTGACCGGGAAGTCGGTCACGCTACCTCTTCTGAAGCCTGTCTGTCACTCCCTTCGCAGTGTATCATTCTGTTTAACG
AGACTGTTTAAACGGAAAAATCTTGATGAATACTTTACGTATTGGCTTAGTTTCCATCTCTGATCGCGCATCCAGCGGCG
TTTATCAGGATAAAGGCATCCCTGCGCTGGAAGAATGGCTGACATCGGCGCTAACCACGCCGTTTGAACTGGAAACCCGC
TTAATCCCCGATGAGCAGGCGATCATCGAGCAAACGTTGTGTGAGCTGGTGGATGAAATGAGTTGCCATCTGGTGCTCAC
CACGGGCGGAACTGGCCCGGCGCGTCGTGACGTAACGCCCGATGCGACGCTGGCAGTAGCGGACCGCGAGATGCCTGGCT
TTGGTGAACAGATGCGCCAGATCAGCCTGCATTTTGTACCAACTGCGATCCTTTCGCGTCAGGTGGGCGTGATTCGCAAA
CAGGCGCTGATCCTTAACTTACCCGGTCAGCCGAAGTCTATTAAAGAGACGCTGGAAGGTGTGAAGGACGCTGAGGGTAA
CGTTGTGGTACACGGTATTTTTGCCAGCGTACCGTACTGCATTCAGTTGCTGGAAGGGCCATACGTTGAAACGGCACCGG
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TATTCCCTGGCGATAACAGTCGGGCCGGGGTGCGGCGGCAAAAAACCGTGTGCGACCAGCAAACCAGAAAGCATCGGCAC
ACACATAAACATCGGTGATATTTTTGCTTCACGGGCAATAGCGAATAAAATAGGTACCAGAAGAATTAAACCGACTTCGA
AAAAAAGTGCGATACCGACAATAAACGCCGAACAGACCACTGCCCAGTCAAGTTTATTTTTCCCGAAATAATTCAGCATG
```

In [26]: head genoma.fna

>NC_000913.3 Escherichia coli str. K-12 substr. MG1655, complete genome
```
AGCTTTTCATTCTGACTGCAACGGGCAATATGTCTCTGTGTGGATTAAAAAAAGAGTGTCTGATAGCAGCTTCTGAACTG
GTTACCTGCCGTGAGTAAATTAAAATTTTATTGACTTAGGTCACTAAATACTTTAACCAATATAGGCATAGCGCACAGAC
AGATAAAAATTACAGAGTACACAACATCCATGAAACGCATTAGCACCACCATTACCACCACCATCACCATTACCACAGGT
AACGGTGCGGGCTGACGCGTACAGGAAACACAGAAAAAAGCCCGCACCTGACAGTGCGGGCTTTTTTTTTCGACCAAAGG
TAACGAGGTAACAACCATGCGAGTGTTGAAGTTCGGCGGTACATCAGTGGCAAATGCAGAACGTTTTCTGCGTGTTGCCG
ATATTCTGGAAAGCAATGCCAGGCAGGGGCAGGTGGCCACCGTCCTCTCTGCCCCCGCCAAAATCACCAACCACCTGGTG
GCGATGATTGAAAAAACCATTAGCGGCCAGGATGCTTTACCCAATATCAGCGATGCCGAACGTATTTTTGCCGAACTTTT
GACGGGACTCGCCGCCGCCCAGCCGGGGTTCCCGCTGGCGCAATTGAAAACTTTCGTCGATCAGGAATTTGCCCAAATAA
AACATGTCCTGCATGGCATTAGTTTGTTGGGGCAGTGCCCGGATAGCATCAACGCTGCGCTGATTTGCCGTGGCGAGAAA
```

In [27]: tail genoma.fna
```
CGCGTAATAAGTTGACAGTGATCACCCGGTTCGCGGTTATTTGATCAAGAAGAGTGGCAATATGCGTATAACGATTATTC
TGGTCGCACCCGCCAGAGCAGAAAATATTGGGGCAGCGGCGCGGGCAATGAAAACGATGGGGTTTAGCGATCTGCGGATT
```

```
GTCGATAGTCAGGCACACCTGGAGCCAGCCACCCGCTGGGTCGCACATGGATCTGGTGATATTATTGATAATATTAAAGT
TTTCCCGACATTGGCTGAATCGTTACACGATGTCGATTTCACTGTCGCCACCACTGCGCGCAGTCGGGCGAAATATCATT
ACTACGCCACGCCAGTTGAACTGGTGCCGCTGTTAGAGGAAAAATCTTCATGGATGAGCCATGCCGCGCTGGTGTTTGGT
CGCGAAGATTCCGGGTTGACTAACGAAGAGTTAGCGTTGGCTGACGTTCTTACTGGTGTGCCGATGGTGGCGGATTATCC
TTCGCTCAATCTGGGGCAGGCGGTGATGGTCTATTGCTATCAATTAGCAACATTAATACAACAACCGGCGAAAAGTGATG
CAACGGCAGACCAACATCAACTGCAAGCTTTACGCGAACGAGCCATGACATTGCTGACGACTCTGGCAGTGGCAGATGAC
ATAAAACTGGTCGACTGGTTACAACAACGCCTGGGGCTTTTAGAGCAACGAGACACGGCAATGTTGCACCGTTTGCTGCA
TGATATTGAAAAAAATATCACCAAATAAAAAACGCCTTAGTAAGTATTTTTC
```

3.8 Descarga de archivos

- wget : Descarga un archivo de un URL (http:/....)

```
In [28]: ### Ejemplos: Descarga del archivo ftp://ftp.ncbi.nlm.nih.gov/genomes/Drosophila_mira
         wget ftp://ftp.ncbi.nlm.nih.gov/genomes/Drosophila_miranda/protein/protein.fa.gz
         ll


--2018-10-01 21:31:53--  ftp://ftp.ncbi.nlm.nih.gov/genomes/Drosophila_miranda/protein/protein
           => protein.fa.gz
Resolving ftp.ncbi.nlm.nih.gov (ftp.ncbi.nlm.nih.gov)... 130.14.250.10, 2607:f220:41e:250::7
Connecting to ftp.ncbi.nlm.nih.gov (ftp.ncbi.nlm.nih.gov)|130.14.250.10|:21... connected.
Logging in as anonymous ... Logged in!
==> SYST ... done.    ==> PWD ... done.
==> TYPE I ... done.  ==> CWD (1) /genomes/Drosophila_miranda/protein ... done.
==> SIZE protein.fa.gz ... 5512360
==> PASV ... done.    ==> RETR protein.fa.gz ... done.
Length: 5512360 (5.3M) (unauthoritative)

protein.fa.gz       100%[====================>]   5.26M   954KB/s    in 5.9s

2018-10-01 21:32:01 (912 KB/s) - protein.fa.gz saved [5512360]

total 9.8M
-rwxrwxr-x 1 4.5M 10-01 21:22 genoma.fna
-rw-rw-r-- 1    0 10-01 21:06 mundo.txt
-rw-rw-r-- 1 5.3M 10-01 21:32 protein.fa.gz
```

3.9 Descompresión archivos

- gunzip : Descomprime **archivo** con extension **.gz**
- tar : Descomprime **archivo** con extension **.tgz** o **.tar.gz**

```
In [29]: ### Descompresión del archivo "protein.fa.gz"
         gunzip protein.fa.gz
         ll
```

```
total 23M
-rwxrwxr-x 1 4.5M 10-01 21:22 genoma.fna
-rw-rw-r-- 1    0 10-01 21:06 mundo.txt
-rw-rw-r-- 1  18M 10-01 21:32 protein.fa
```

In [34]: *### Despliegue 10 primeras líneas*
         head protein.fa

```
>gi|1036701783|ref|XP_017155988.1| PREDICTED: LOW QUALITY PROTEIN: ATP-dependent RNA helicase
MKVLIATSVASRGLDIKNVKHVVNYDMPKTIDDYVHRIGRTGRVXNYGRATSFFDPEQDSGVATDLVKIL
EGSEQVVPDFLRVLSGVGGHGGCGAQFGGYDVRGSGNVIQEASTVEDEQEWD
>gi|1036701631|ref|XP_017155883.1| PREDICTED: LOW QUALITY PROTEIN: probable cytochrome P450 31
MPCWVLLLAGLGLLLLLQKWALLRLGSSLPGPWAFPLLGNAQMVGKLRPEFIFLVFTELRDRFGATYRLW
LGPQLWVFLHTAEETREALHDPTLRKAKTFLQLEPLIGNGLLISHGSHWTQQRRLLTPAFRPILLRSFAP
SVAKHADRLIQRLEETRGSFLEVTDPLFACLLDAIVDTSMGEQLETQSVAHSPIIKAFHRSSQLLFKRMI
NPLLXSDWVFKQTRLWRELSEQLQVIHTLMDSLIXRRSAQLQQQGNEPRKTLNLLDTLLLAKVDGEPLSR
KEIRDEINTFVFAGVDTTTAAMSFVLYALAMNPEAQDRLQEELRQLPPGLSTDLDTLNELPYLDXLLKEV
LRLYTIVPTTGRQTTRSTVIGGRTYCAGVTLWINMYGLAHDASYFPDPYVFKPERWMQETELQTEVPFSY
```

In [35]: *### Despliegue 10 ultimas líneas*
         tail protein.fa

```
LLSEGEGVDYHTFRLRSERGMRIVIRHLHRSTPTDWVREQLGRLGYKVRHIANIHKRFTKEPLDLFEVEL
EPQESNNGIYELTAICSQRIKVEKPIRAPGVPQCHKCQMFGHSKNYCSRAVVCLKCGEKHDPKDCPKKRE
DVPKCANCNGPHVASYRGCIKYKEYCNRRKNGLEQTNRMLSKLSMPTLPAAQGLRRPRQPPVMNQQGFPA
LQQPSRRRARSGARAPAPPAQMSSYADALRTRQPASHSQPLPGRQVWQSVPPTATTIGFQPTVSSQSVRP
SDELSQKLDYLITLLTQERIQHAAAAAESTQKLEGKIDVLIAQMTKLTDCLYASIQANSYASNV
>gi|1037106843|ref|XP_017155477.1| PREDICTED: uncharacterized protein LOC108164382 [Drosophila
MAVVPLPFAAANFILSAAAIALSCWPLLLLVLAWRRSIIRDRLHRLDPLVLAPPGRWLGMYWIIVVVWSS
VFGTVRNRTALDERVCKVRQETWTMHSAVQSGWSDRRGPPGGPGFGQICVAMVASSHEVQENHPCVDSDA
PCVQRKRVQICIHGQPAAVMTLVRGALLEAGCRGDAHEEGSEEQLGVSPGVASVFVHASGTTTMSTTGIL
DLLRPDCGVPIIGPCCTKVRSDCGGVCNVPAGIIVVVVVVVI
```

3.10   Filtrado de un archivo con wget

USO: wget < patronParaBuscar > < nombreArchivo >

In [36]: *### Ejemplo: Muestre solo los encabezados del archivo genoma.fna*
         grep ">" protein.fa

```
>gi|1036701783|ref|XP_017155988.1| PREDICTED: LOW QUALITY PROTEIN: ATP-dependent RNA helicase
>gi|1036701631|ref|XP_017155883.1| PREDICTED: LOW QUALITY PROTEIN: probable cytochrome P450 31
>gi|1036701733|ref|XP_017155947.1| PREDICTED: uncharacterized protein LOC108164616 [Drosophila
>gi|1036701800|ref|XP_017155998.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036701338|ref|XP_017155122.1| PREDICTED: serine/threonine-protein kinase par-1 [Drosophila
>gi|1036701601|ref|XP_017155861.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
```

```
>gi|1036701488|ref|XP_017155806.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701509|ref|XP_017155814.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701526|ref|XP_017155824.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701545|ref|XP_017155832.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701562|ref|XP_017155842.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701582|ref|XP_017155851.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701768|ref|XP_017155968.1| PREDICTED: E3 ubiquitin-protein ligase RNF128 [Drosophila m
>gi|1036701422|ref|XP_017155494.1| PREDICTED: alpha-actinin, sarcomeric isoform X4 [Drosophila
>gi|1036701616|ref|XP_017155871.1| PREDICTED: protein ultraspiracle [Drosophila miranda]
>gi|1036701355|ref|XP_017155231.1| PREDICTED: alpha-actinin, sarcomeric isoform X1 [Drosophila
>gi|1036701374|ref|XP_017155306.1| PREDICTED: alpha-actinin, sarcomeric isoform X1 [Drosophila
>gi|1036701390|ref|XP_017155378.1| PREDICTED: alpha-actinin, sarcomeric isoform X2 [Drosophila
>gi|1036701406|ref|XP_017155453.1| PREDICTED: alpha-actinin, sarcomeric isoform X3 [Drosophila
>gi|1036701287|ref|XP_017154817.1| PREDICTED: uncharacterized protein LOC108163645 isoform X4
>gi|1036701272|ref|XP_017154742.1| PREDICTED: uncharacterized protein LOC108163645 isoform X3
>gi|1036701257|ref|XP_017154665.1| PREDICTED: uncharacterized protein LOC108163645 isoform X2
>gi|1036701243|ref|XP_017154584.1| PREDICTED: uncharacterized protein LOC108163645 isoform X1
>gi|1036701701|ref|XP_017155930.1| PREDICTED: putative GPI-anchor transamidase [Drosophila mira
>gi|1036701717|ref|XP_017155937.1| PREDICTED: putative GPI-anchor transamidase [Drosophila mira
>gi|1036698287|ref|XP_017139785.1| PREDICTED: AF4/FMR2 family member 4-like [Drosophila miranda
>gi|1036701323|ref|XP_017155006.1| PREDICTED: hormone receptor 4-like isoform X2 [Drosophila m
>gi|1036701305|ref|XP_017154930.1| PREDICTED: hormone receptor 4-like isoform X1 [Drosophila m
>gi|1036701752|ref|XP_017155958.1| PREDICTED: protein KTI12 homolog [Drosophila miranda]
>gi|1036701647|ref|XP_017155896.1| PREDICTED: protein SMG9 [Drosophila miranda]
>gi|1036701680|ref|XP_017155916.1| PREDICTED: carnosine N-methyltransferase isoform X2 [Drosoph
>gi|1036701663|ref|XP_017155907.1| PREDICTED: carnosine N-methyltransferase isoform X1 [Drosoph
>gi|1036701439|ref|XP_017155626.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila miran
>gi|1036701454|ref|XP_017155715.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila miran
>gi|1036701471|ref|XP_017155795.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila miran
>gi|1036711564|ref|XP_017139547.1| PREDICTED: dehydrodolichyl diphosphate synthase complex subu
>gi|1036711597|ref|XP_017139567.1| PREDICTED: 40S ribosomal protein S14 [Drosophila miranda]
>gi|1036711581|ref|XP_017139558.1| PREDICTED: 40S ribosomal protein S14-like [Drosophila miran
>gi|1036711500|ref|XP_017139511.1| PREDICTED: E3 ubiquitin-protein ligase UBR3 [Drosophila mira
>gi|1036711515|ref|XP_017139518.1| PREDICTED: E3 ubiquitin-protein ligase UBR3 [Drosophila mira
>gi|1036711534|ref|XP_017139529.1| PREDICTED: uncharacterized protein LOC108153940 isoform X1
>gi|1036711550|ref|XP_017139537.1| PREDICTED: uncharacterized protein LOC108153940 isoform X2
>gi|1036751956|ref|XP_017156406.1| PREDICTED: DM7 family protein GG17591 [Drosophila miranda]
>gi|1036729081|ref|XP_017149347.1| PREDICTED: neuropathy target esterase sws isoform X4 [Drosop
>gi|1036729050|ref|XP_017149329.1| PREDICTED: neuropathy target esterase sws isoform X2 [Drosop
>gi|1036729021|ref|XP_017149318.1| PREDICTED: neuropathy target esterase sws isoform X1 [Drosop
>gi|1036729063|ref|XP_017149338.1| PREDICTED: neuropathy target esterase sws isoform X3 [Drosop
>gi|1036737636|ref|XP_017154022.1| PREDICTED: protein singed [Drosophila miranda]
>gi|1036702955|ref|XP_017156697.1| PREDICTED: protein arginine N-methyltransferase 1-like [Dros
>gi|1036707540|ref|XP_017137034.1| PREDICTED: E3 ubiquitin ligase complex SCF subunit sconC-li
>gi|1036702891|ref|XP_017156653.1| PREDICTED: uncharacterized protein LOC108165104 [Drosophila
>gi|1036756219|ref|XP_017156657.1| PREDICTED: tyramine beta-hydroxylase [Drosophila miranda]
>gi|1036756236|ref|XP_017156658.1| PREDICTED: tyramine beta-hydroxylase [Drosophila miranda]
>gi|1036756255|ref|XP_017156659.1| PREDICTED: tyramine beta-hydroxylase [Drosophila miranda]
```

```
>gi|1036756728|ref|XP_017156687.1| PREDICTED: uncharacterized protein LOC108165131 isoform X3
>gi|1036756709|ref|XP_017156686.1| PREDICTED: uncharacterized protein LOC108165131 isoform X2
>gi|1036756690|ref|XP_017156685.1| PREDICTED: uncharacterized protein LOC108165131 isoform X1
>gi|1036709552|ref|XP_017138295.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709468|ref|XP_017138254.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709484|ref|XP_017138263.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709501|ref|XP_017138272.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709518|ref|XP_017138278.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709535|ref|XP_017138286.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709568|ref|XP_017138303.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709612|ref|XP_017138327.1| PREDICTED: uncharacterized protein LOC108153062 [Drosophila
>gi|1036709344|ref|XP_017138193.1| PREDICTED: alpha-(1,6)-fucosyltransferase [Drosophila mirand
>gi|1036709365|ref|XP_017138200.1| PREDICTED: alpha-(1,6)-fucosyltransferase [Drosophila mirand
>gi|1036709381|ref|XP_017138209.1| PREDICTED: alpha-(1,6)-fucosyltransferase [Drosophila mirand
>gi|1036709430|ref|XP_017138234.1| PREDICTED: uncharacterized protein LOC108153001 [Drosophila
>gi|1036709449|ref|XP_017138244.1| PREDICTED: uncharacterized protein LOC108153001 [Drosophila
>gi|1036709597|ref|XP_017138315.1| PREDICTED: thioredoxin domain-containing protein 5 [Drosophi
>gi|1036709398|ref|XP_017138222.1| PREDICTED: BTB/POZ domain-containing protein 17 [Drosophila
>gi|1036752027|ref|XP_017156411.1| PREDICTED: putative transporter svop-1 [Drosophila miranda]
>gi|1036752042|ref|XP_017156412.1| PREDICTED: uncharacterized protein LOC108164934 [Drosophila
>gi|1036698181|ref|XP_017138639.1| PREDICTED: uncharacterized protein LOC108150747 [Drosophila
>gi|1036752060|ref|XP_017156413.1| PREDICTED: uncharacterized protein LOC108164935 [Drosophila
>gi|1036698196|ref|XP_017150822.1| PREDICTED: uncharacterized protein LOC108158651 [Drosophila
>gi|1036756638|ref|XP_017156682.1| PREDICTED: uncharacterized protein LOC108165128 [Drosophila
>gi|1036756653|ref|XP_017156683.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036706619|ref|XP_017136424.1| PREDICTED: uncharacterized protein LOC108151975 [Drosophila
>gi|1036755818|ref|XP_017156635.1| PREDICTED: uncharacterized protein LOC108165094 isoform X1
>gi|1036755837|ref|XP_017156636.1| PREDICTED: uncharacterized protein LOC108165094 isoform X2
>gi|1036755645|ref|XP_017156624.1| PREDICTED: insecticyanin-A [Drosophila miranda]
>gi|1036729511|ref|XP_017149608.1| PREDICTED: protein wings apart-like [Drosophila miranda]
>gi|1036698304|ref|XP_017140963.1| PREDICTED: uncharacterized protein LOC108154713 [Drosophila
>gi|1036755698|ref|XP_017156628.1| PREDICTED: uncharacterized protein LOC108165087 [Drosophila
>gi|1036755716|ref|XP_017156629.1| PREDICTED: uncharacterized protein LOC108165088 [Drosophila
>gi|1036730248|ref|XP_017150076.1| PREDICTED: LOW QUALITY PROTEIN: histone H2B-like [Drosophila
>gi|1036730189|ref|XP_017150041.1| PREDICTED: histone H1-like isoform X1 [Drosophila miranda]
>gi|1036730208|ref|XP_017150051.1| PREDICTED: histone H1-like isoform X2 [Drosophila miranda]
>gi|1036730229|ref|XP_017150062.1| PREDICTED: protein lethal(2)denticleless-like [Drosophila mi
>gi|1036729548|ref|XP_017149634.1| PREDICTED: centrosomal protein of 162 kDa isoform X1 [Drosop
>gi|1036729567|ref|XP_017149643.1| PREDICTED: centrosomal protein of 162 kDa isoform X2 [Drosop
>gi|1036729580|ref|XP_017149655.1| PREDICTED: carboxypeptidase B isoform X1 [Drosophila miranda
>gi|1036729600|ref|XP_017149664.1| PREDICTED: carboxypeptidase B isoform X2 [Drosophila miranda
>gi|1036698208|ref|XP_017156463.1| PREDICTED: zinc carboxypeptidase A 1 [Drosophila miranda]
>gi|1036729529|ref|XP_017149621.1| PREDICTED: histone acetyltransferase KAT6A [Drosophila miran
>gi|1036755785|ref|XP_017156633.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036703982|ref|XP_017134622.1| PREDICTED: proteasome subunit beta type-7 [Drosophila mirand
>gi|1036703732|ref|XP_017157117.1| PREDICTED: dopamine N-acetyltransferase [Drosophila miranda]
>gi|1036747149|ref|XP_017156152.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036704091|ref|XP_017134680.1| PREDICTED: bax inhibitor 1-like [Drosophila miranda]
```

```
>gi|1036698319|ref|XP_017142208.1| PREDICTED: uncharacterized protein LOC108155475 [Drosophila
>gi|1036741943|ref|XP_017155857.1| PREDICTED: glycosylphosphatidylinositol anchor attachment 1
>gi|1036713747|ref|XP_017140860.1| PREDICTED: uncharacterized protein LOC108154906 isoform X2
>gi|1036713732|ref|XP_017140851.1| PREDICTED: uncharacterized protein LOC108154906 isoform X1
>gi|1036713793|ref|XP_017140898.1| PREDICTED: G2/M phase-specific E3 ubiquitin-protein ligase
>gi|1036713764|ref|XP_017140871.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 30 homolog
>gi|1036713827|ref|XP_017140923.1| PREDICTED: LOW QUALITY PROTEIN: T-complex protein 11-like p
>gi|1036713779|ref|XP_017140883.1| PREDICTED: DDB1- and CUL4-associated factor 10 homolog [Dros
>gi|1036713810|ref|XP_017140913.1| PREDICTED: patatin-like phospholipase domain-containing prot
>gi|1036741536|ref|XP_017155836.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036741517|ref|XP_017155835.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036741577|ref|XP_017155838.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036741558|ref|XP_017155837.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036706037|ref|XP_017136019.1| PREDICTED: tubulin-specific chaperone A-like [Drosophila mi
>gi|1036739823|ref|XP_017155212.1| PREDICTED: microsomal glutathione S-transferase 1 [Drosophil
>gi|1036739857|ref|XP_017155232.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X2
>gi|1036739839|ref|XP_017155224.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X
>gi|1036739875|ref|XP_017155242.1| PREDICTED: cytochrome c oxidase copper chaperone [Drosophila
>gi|1036739893|ref|XP_017155248.1| PREDICTED: cytochrome c oxidase copper chaperone [Drosophila
>gi|1036739805|ref|XP_017155202.1| PREDICTED: translation machinery-associated protein 16 homol
>gi|1036739787|ref|XP_017155190.1| PREDICTED: LOW QUALITY PROTEIN: lipoamide acyltransferase co
>gi|1036738684|ref|XP_017154541.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738702|ref|XP_017154547.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738720|ref|XP_017154556.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738740|ref|XP_017154564.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738758|ref|XP_017154576.1| PREDICTED: tubulin-specific chaperone A [Drosophila miranda]
>gi|1036752119|ref|XP_017156416.1| PREDICTED: probable cytochrome P450 4s3 [Drosophila miranda]
>gi|1036740350|ref|XP_017155473.1| PREDICTED: uncharacterized protein LOC108164335 isoform X1
>gi|1036740366|ref|XP_017155474.1| PREDICTED: uncharacterized protein LOC108164335 isoform X2
>gi|1036740384|ref|XP_017155475.1| PREDICTED: probable 60S ribosomal protein L37-A [Drosophila
>gi|1036743313|ref|XP_017155936.1| PREDICTED: uncharacterized protein LOC108164608 [Drosophila
>gi|1036743295|ref|XP_017155935.1| PREDICTED: uncharacterized protein LOC108164607 [Drosophila
>gi|1036705018|ref|XP_017135315.1| PREDICTED: uncharacterized protein LOC108151289 [Drosophila
>gi|1036753611|ref|XP_017156508.1| PREDICTED: uncharacterized protein LOC108164994 isoform X1
>gi|1036753631|ref|XP_017156510.1| PREDICTED: uncharacterized protein LOC108164994 isoform X2
>gi|1036748586|ref|XP_017156220.1| PREDICTED: uncharacterized protein LOC108164812 [Drosophila
>gi|1036698334|ref|XP_017143452.1| PREDICTED: uncharacterized protein LOC108156218 [Drosophila
>gi|1036751287|ref|XP_017156370.1| PREDICTED: uncharacterized protein LOC108164904 [Drosophila
>gi|1036751308|ref|XP_017156371.1| PREDICTED: uncharacterized protein LOC108164904 [Drosophila
>gi|1036751329|ref|XP_017156372.1| PREDICTED: uncharacterized protein LOC108164905 [Drosophila
>gi|1036705899|ref|XP_017135911.1| PREDICTED: uncharacterized protein LOC108151676 [Drosophila
>gi|1036739354|ref|XP_017154947.1| PREDICTED: uncharacterized protein LOC108163921 [Drosophila
>gi|1036738971|ref|XP_017154717.1| PREDICTED: uncharacterized protein LOC108163755 [Drosophila
>gi|1036738989|ref|XP_017154725.1| PREDICTED: uncharacterized protein LOC108163755 [Drosophila
>gi|1036739007|ref|XP_017154733.1| PREDICTED: uncharacterized protein LOC108163755 [Drosophila
>gi|1036715019|ref|XP_017141618.1| PREDICTED: lymphokine-activated killer T-cell-originated pro
>gi|1036715000|ref|XP_017141607.1| PREDICTED: partitioning defective protein 6 [Drosophila mira
>gi|1036715036|ref|XP_017141631.1| PREDICTED: ubiquitin-conjugating enzyme E2 S isoform X1 [Dro
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>gi|1036715053|ref|XP_017141639.1| PREDICTED: ubiquitin-conjugating enzyme E2 S isoform X2 [Dro
>gi|1036715091|ref|XP_017141660.1| PREDICTED: uncharacterized protein LOC108155384 isoform X2
>gi|1036715072|ref|XP_017141652.1| PREDICTED: uncharacterized protein LOC108155384 isoform X1
>gi|1036715109|ref|XP_017141670.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036756166|ref|XP_017156654.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036709204|ref|XP_017138102.1| PREDICTED: spectrin beta chain isoform X2 [Drosophila miran
>gi|1036709154|ref|XP_017138077.1| PREDICTED: spectrin beta chain isoform X1 [Drosophila miran
>gi|1036709171|ref|XP_017138085.1| PREDICTED: spectrin beta chain isoform X1 [Drosophila miran
>gi|1036709188|ref|XP_017138093.1| PREDICTED: spectrin beta chain isoform X1 [Drosophila miran
>gi|1036709219|ref|XP_017138111.1| PREDICTED: spectrin beta chain isoform X3 [Drosophila miran
>gi|1036709232|ref|XP_017138123.1| PREDICTED: integrator complex subunit 2 [Drosophila miranda]
>gi|1036709251|ref|XP_017138135.1| PREDICTED: rho GTPase-activating protein 190 isoform X1 [Dro
>gi|1036709267|ref|XP_017138143.1| PREDICTED: rho GTPase-activating protein 190 isoform X1 [Dro
>gi|1036709281|ref|XP_017138151.1| PREDICTED: rho GTPase-activating protein 190 isoform X2 [Dro
>gi|1036709296|ref|XP_017138161.1| PREDICTED: rho GTPase-activating protein 190 isoform X3 [Dro
>gi|1036709312|ref|XP_017138170.1| PREDICTED: rho GTPase-activating protein 190 isoform X4 [Dro
>gi|1036709327|ref|XP_017138181.1| PREDICTED: rho GTPase-activating protein 190 isoform X5 [Dro
>gi|1036742108|ref|XP_017155865.1| PREDICTED: probable ATP-dependent RNA helicase DDX10 [Drosop
>gi|1036736749|ref|XP_017153521.1| PREDICTED: calpain-D isoform X1 [Drosophila miranda]
>gi|1036736770|ref|XP_017153530.1| PREDICTED: calpain-D isoform X2 [Drosophila miranda]
>gi|1036736789|ref|XP_017153539.1| PREDICTED: calpain-D isoform X3 [Drosophila miranda]
>gi|1036736807|ref|XP_017153548.1| PREDICTED: calpain-D isoform X4 [Drosophila miranda]
>gi|1036738006|ref|XP_017154223.1| PREDICTED: septin-1 isoform X2 [Drosophila miranda]
>gi|1036737988|ref|XP_017154215.1| PREDICTED: septin-1 isoform X1 [Drosophila miranda]
>gi|1036718395|ref|XP_017143507.1| PREDICTED: translation factor waclaw, mitochondrial [Drosoph
>gi|1036718226|ref|XP_017143423.1| PREDICTED: uncharacterized protein LOC108156467 [Drosophila
>gi|1036718246|ref|XP_017143430.1| PREDICTED: uncharacterized protein LOC108156467 [Drosophila
>gi|1036718264|ref|XP_017143439.1| PREDICTED: uncharacterized protein LOC108156467 [Drosophila
>gi|1036718303|ref|XP_017143462.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X2
>gi|1036718377|ref|XP_017143496.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X4
>gi|1036718283|ref|XP_017143454.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X1
>gi|1036718322|ref|XP_017143472.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X1
>gi|1036718343|ref|XP_017143479.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X1
>gi|1036718360|ref|XP_017143487.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X3
>gi|1036718410|ref|XP_017143518.1| PREDICTED: probable ATP-dependent RNA helicase DDX56 [Drosop
>gi|1036751607|ref|XP_017156387.1| PREDICTED: uncharacterized protein LOC108164912 [Drosophila
>gi|1036742446|ref|XP_017155886.1| PREDICTED: mpv17-like protein [Drosophila miranda]
>gi|1036742427|ref|XP_017155885.1| PREDICTED: protein misato [Drosophila miranda]
>gi|1036698349|ref|XP_017144690.1| PREDICTED: myosin heavy chain IB-like [Drosophila miranda]
>gi|1036706022|ref|XP_017136003.1| PREDICTED: proline-rich receptor-like protein kinase PERK9
>gi|1036753255|ref|XP_017156488.1| PREDICTED: neurotrimin isoform X4 [Drosophila miranda]
>gi|1036753219|ref|XP_017156486.1| PREDICTED: neurotrimin isoform X2 [Drosophila miranda]
>gi|1036753238|ref|XP_017156487.1| PREDICTED: neuronal growth regulator 1 isoform X3 [Drosophil
>gi|1036753204|ref|XP_017156485.1| PREDICTED: neuronal growth regulator 1 isoform X1 [Drosophil
>gi|1036706318|ref|XP_017136222.1| PREDICTED: uncharacterized protein LOC108151845 [Drosophila
>gi|1036735067|ref|XP_017152556.1| PREDICTED: uncharacterized protein LOC108162358 [Drosophila
>gi|1036728354|ref|XP_017148921.1| PREDICTED: neutral alpha-glucosidase AB [Drosophila miranda]
>gi|1036728373|ref|XP_017148929.1| PREDICTED: neutral alpha-glucosidase AB [Drosophila miranda]
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>gi|1036728412|ref|XP_017148951.1| PREDICTED: tektin-3-like isoform X2 [Drosophila miranda]
>gi|1036728391|ref|XP_017148943.1| PREDICTED: tektin-3-like isoform X1 [Drosophila miranda]
>gi|1036728433|ref|XP_017148962.1| PREDICTED: NFU1 iron-sulfur cluster scaffold homolog, mitoch
>gi|1036742771|ref|XP_017155904.1| PREDICTED: uncharacterized protein LOC108164590 [Drosophila
>gi|1036725812|ref|XP_017147391.1| PREDICTED: solute carrier family 35 member E1 homolog [Droso
>gi|1036725850|ref|XP_017147422.1| PREDICTED: uncharacterized protein LOC108159019 [Drosophila
>gi|1036725831|ref|XP_017147407.1| PREDICTED: DDB1- and CUL4-associated factor 7 [Drosophila mi
>gi|1036725012|ref|XP_017146926.1| PREDICTED: cell wall protein AWA1 isoform X4 [Drosophila mi
>gi|1036725030|ref|XP_017146937.1| PREDICTED: cell wall protein AWA1 isoform X4 [Drosophila mi
>gi|1036725051|ref|XP_017146946.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 2 isoform
>gi|1036724951|ref|XP_017146896.1| PREDICTED: myb-like protein A isoform X1 [Drosophila miranda
>gi|1036724993|ref|XP_017146916.1| PREDICTED: mucin-12 isoform X3 [Drosophila miranda]
>gi|1036724972|ref|XP_017146906.1| PREDICTED: myb-like protein A isoform X2 [Drosophila miranda
>gi|1036723743|ref|XP_017146122.1| PREDICTED: uncharacterized protein C18orf19 homolog A [Droso
>gi|1036723724|ref|XP_017146111.1| PREDICTED: tRNA methyltransferase 10 homolog A [Drosophila m
>gi|1036723666|ref|XP_017146082.1| PREDICTED: uncharacterized protein LOC108158350 [Drosophila
>gi|1036723685|ref|XP_017146092.1| PREDICTED: uncharacterized protein LOC108158350 [Drosophila
>gi|1036723704|ref|XP_017146101.1| PREDICTED: uncharacterized protein LOC108158350 [Drosophila
>gi|1036722128|ref|XP_017145358.1| PREDICTED: uncharacterized protein LOC108157696 [Drosophila
>gi|1036721934|ref|XP_017145274.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721953|ref|XP_017145283.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722053|ref|XP_017145325.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721970|ref|XP_017145291.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722071|ref|XP_017145333.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721919|ref|XP_017145266.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721830|ref|XP_017145225.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722007|ref|XP_017145309.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722025|ref|XP_017145317.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721864|ref|XP_017145241.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722090|ref|XP_017145340.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721883|ref|XP_017145249.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722111|ref|XP_017145346.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721847|ref|XP_017145232.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721902|ref|XP_017145258.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721989|ref|XP_017145300.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036727249|ref|XP_017148291.1| PREDICTED: uncharacterized protein LOC108159474 isoform X1
>gi|1036727232|ref|XP_017148278.1| PREDICTED: ribosomal protein S6 kinase 2 beta [Drosophila m
>gi|1036727279|ref|XP_017148311.1| PREDICTED: protein phosphatase 1H [Drosophila miranda]
>gi|1036727262|ref|XP_017148299.1| PREDICTED: formin BNI1 isoform X2 [Drosophila miranda]
>gi|1036704459|ref|XP_017134908.1| PREDICTED: clusterin-associated protein 1 [Drosophila mirand
>gi|1036736279|ref|XP_017153243.1| PREDICTED: adenylate cyclase type 9 isoform X1 [Drosophila m
>gi|1036736297|ref|XP_017153251.1| PREDICTED: adenylate cyclase type 9 isoform X2 [Drosophila m
>gi|1036736315|ref|XP_017153260.1| PREDICTED: adenylate cyclase type 9 isoform X3 [Drosophila m
>gi|1036715889|ref|XP_017142115.1| PREDICTED: thyroid receptor-interacting protein 11 [Drosophi
>gi|1036715959|ref|XP_017142159.1| PREDICTED: F-box-like/WD repeat-containing protein TBL1XR1
>gi|1036715941|ref|XP_017142147.1| PREDICTED: arginine/serine-rich coiled-coil protein 2 isofor
>gi|1036715923|ref|XP_017142139.1| PREDICTED: arginine/serine-rich coiled-coil protein 2 isofor
>gi|1036715907|ref|XP_017142126.1| PREDICTED: uncharacterized protein LOC108155670 [Drosophila
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>gi|1036742178|ref|XP_017155869.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036742197|ref|XP_017155870.1| PREDICTED: UDP-N-acetylglucosamine transferase subunit ALG14
>gi|1036724525|ref|XP_017146626.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036724579|ref|XP_017146667.1| PREDICTED: uncharacterized protein LOC108158667 [Drosophila
>gi|1036724560|ref|XP_017146653.1| PREDICTED: glycolipid transfer protein [Drosophila miranda]
>gi|1036698364|ref|XP_017147071.1| PREDICTED: cell surface glycoprotein 1-like [Drosophila mira
>gi|1036724541|ref|XP_017146641.1| PREDICTED: uncharacterized protein LOC108158654 [Drosophila
>gi|1036749432|ref|XP_017156265.1| PREDICTED: putative inositol monophosphatase 3 [Drosophila m
>gi|1036733728|ref|XP_017151782.1| PREDICTED: histone deacetylase 4 isoform X1 [Drosophila mira
>gi|1036733765|ref|XP_017151798.1| PREDICTED: histone deacetylase 5 isoform X3 [Drosophila mira
>gi|1036733747|ref|XP_017151790.1| PREDICTED: histone deacetylase 4 isoform X2 [Drosophila mira
>gi|1036733782|ref|XP_017151808.1| PREDICTED: histone deacetylase 4 isoform X4 [Drosophila mira
>gi|1036713039|ref|XP_017140388.1| PREDICTED: probable peroxisomal acyl-coenzyme A oxidase 1 [D
>gi|1036713072|ref|XP_017140410.1| PREDICTED: leucine-rich repeat-containing protein 57 [Drosop
>gi|1036713055|ref|XP_017140402.1| PREDICTED: proteasome activator complex subunit 3 [Drosophil
>gi|1036713006|ref|XP_017140366.1| PREDICTED: integrin alpha-PS1 isoform X2 [Drosophila miranda
>gi|1036712989|ref|XP_017140358.1| PREDICTED: integrin alpha-PS1 isoform X1 [Drosophila miranda
>gi|1036713088|ref|XP_017140418.1| PREDICTED: uncharacterized protein LOC108154609 [Drosophila
>gi|1036713023|ref|XP_017140377.1| PREDICTED: vesicle-fusing ATPase 1 [Drosophila miranda]
>gi|1036753070|ref|XP_017156476.1| PREDICTED: protein tilB [Drosophila miranda]
>gi|1036735331|ref|XP_017152710.1| PREDICTED: histone-lysine N-methyltransferase setd3 [Drosoph
>gi|1036735282|ref|XP_017152687.1| PREDICTED: uncharacterized protein LOC108162452 isoform X1
>gi|1036735301|ref|XP_017152695.1| PREDICTED: uncharacterized protein LOC108162452 isoform X2
>gi|1036702908|ref|XP_017156665.1| PREDICTED: uncharacterized protein LOC108165114 [Drosophila
>gi|1036724066|ref|XP_017146325.1| PREDICTED: LOW QUALITY PROTEIN: serine-rich adhesin for plat
>gi|1036754168|ref|XP_017156542.1| PREDICTED: uncharacterized protein LOC108165015 [Drosophila
>gi|1036714661|ref|XP_017141388.1| PREDICTED: uncharacterized protein LOC108155212 [Drosophila
>gi|1036714605|ref|XP_017141352.1| PREDICTED: uncharacterized protein LOC108155188 [Drosophila
>gi|1036714679|ref|XP_017141400.1| PREDICTED: 28 kDa heat- and acid-stable phosphoprotein [Dros
>gi|1036714642|ref|XP_017141375.1| PREDICTED: ribosomal RNA small subunit methyltransferase NEP
>gi|1036714698|ref|XP_017141412.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036714510|ref|XP_017141305.1| PREDICTED: probable cytochrome P450 4d14 [Drosophila miranda
>gi|1036714473|ref|XP_017141284.1| PREDICTED: cytochrome P450 4d2 [Drosophila miranda]
>gi|1036714492|ref|XP_017141293.1| PREDICTED: cytochrome P450 4d2 [Drosophila miranda]
>gi|1036714567|ref|XP_017141332.1| PREDICTED: cytochrome P450 4ae1 isoform X3 [Drosophila miran
>gi|1036714586|ref|XP_017141340.1| PREDICTED: cytochrome P450 4ae1 isoform X4 [Drosophila miran
>gi|1036714529|ref|XP_017141317.1| PREDICTED: cytochrome P450 4ae1 isoform X1 [Drosophila miran
>gi|1036714548|ref|XP_017141325.1| PREDICTED: cytochrome P450 4ae1 isoform X2 [Drosophila miran
>gi|1036714624|ref|XP_017141363.1| PREDICTED: protein prune homolog [Drosophila miranda]
>gi|1036714454|ref|XP_017141272.1| PREDICTED: 60S ribosomal export protein NMD3 [Drosophila mi
>gi|1036714718|ref|XP_017141421.1| PREDICTED: uncharacterized protein LOC108155239 [Drosophila
>gi|1036705377|ref|XP_017135563.1| PREDICTED: keratin-associated protein 19-2 [Drosophila miran
>gi|1036702307|ref|XP_017156278.1| PREDICTED: uncharacterized protein LOC108164857 [Drosophila
>gi|1036736396|ref|XP_017153315.1| PREDICTED: monocarboxylate transporter 12 [Drosophila miran
>gi|1036736415|ref|XP_017153326.1| PREDICTED: fatty acyl-CoA reductase 1 [Drosophila miranda]
>gi|1036727562|ref|XP_017148473.1| PREDICTED: protein msta isoform X1 [Drosophila miranda]
>gi|1036727581|ref|XP_017148482.1| PREDICTED: protein msta isoform X2 [Drosophila miranda]
>gi|1036727524|ref|XP_017148448.1| PREDICTED: vinculin [Drosophila miranda]
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>gi|1036727541|ref|XP_017148459.1| PREDICTED: vinculin [Drosophila miranda]
>gi|1036727598|ref|XP_017148494.1| PREDICTED: histidine-rich glycoprotein isoform X1 [Drosophil
>gi|1036727616|ref|XP_017148503.1| PREDICTED: histidine-rich glycoprotein isoform X1 [Drosophil
>gi|1036727634|ref|XP_017148513.1| PREDICTED: histidine-rich glycoprotein isoform X2 [Drosophil
>gi|1036715126|ref|XP_017141684.1| PREDICTED: protein pecanex [Drosophila miranda]
>gi|1036715144|ref|XP_017141695.1| PREDICTED: probable ATP-dependent RNA helicase kurz [Drosoph
>gi|1036744253|ref|XP_017155981.1| PREDICTED: DNA-binding protein K10 [Drosophila miranda]
>gi|1036705611|ref|XP_017135701.1| PREDICTED: odorant receptor 2a [Drosophila miranda]
>gi|1036743910|ref|XP_017155964.1| PREDICTED: protein crooked neck [Drosophila miranda]
>gi|1036698381|ref|XP_017148371.1| PREDICTED: uncharacterized protein LOC108159290 [Drosophila
>gi|1036729414|ref|XP_017149549.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036729322|ref|XP_017149494.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036729341|ref|XP_017149503.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036729357|ref|XP_017149511.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036729395|ref|XP_017149535.1| PREDICTED: protein prenyltransferase alpha subunit repeat-co
>gi|1036729376|ref|XP_017149523.1| PREDICTED: U3 small nucleolar RNA-associated protein 15 homo
>gi|1036732027|ref|XP_017150947.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X2 [Dr
>gi|1036731916|ref|XP_017150904.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Dr
>gi|1036731935|ref|XP_017150911.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Dr
>gi|1036731954|ref|XP_017150918.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Dr
>gi|1036731971|ref|XP_017150926.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Dr
>gi|1036731989|ref|XP_017150933.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Dr
>gi|1036732008|ref|XP_017150940.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Dr
>gi|1036733364|ref|XP_017151590.1| PREDICTED: hypoxia up-regulated protein 1 [Drosophila miran
>gi|1036733383|ref|XP_017151599.1| PREDICTED: hypoxia up-regulated protein 1 [Drosophila miran
>gi|1036733402|ref|XP_017151612.1| PREDICTED: uncharacterized protein LOC108161779 [Drosophila
>gi|1036733417|ref|XP_017151621.1| PREDICTED: uncharacterized protein LOC108161779 [Drosophila
>gi|1036698396|ref|XP_017149616.1| PREDICTED: uncharacterized protein LOC108159997 [Drosophila
>gi|1036757665|ref|XP_017156736.1| PREDICTED: uncharacterized protein LOC108165176 [Drosophila
>gi|1036750976|ref|XP_017156351.1| PREDICTED: probable insulin-like peptide 6 [Drosophila mira
>gi|1036757496|ref|XP_017156730.1| PREDICTED: uncharacterized protein LOC108165167 [Drosophila
>gi|1036724596|ref|XP_017146682.1| PREDICTED: apoptosis-inducing factor 3 isoform X1 [Drosophil
>gi|1036724612|ref|XP_017146693.1| PREDICTED: apoptosis-inducing factor 3 isoform X2 [Drosophil
>gi|1036724692|ref|XP_017146728.1| PREDICTED: uncharacterized protein LOC108158699 [Drosophila
>gi|1036724632|ref|XP_017146701.1| PREDICTED: apoptosis-inducing factor 3 isoform X3 [Drosophil
>gi|1036724671|ref|XP_017146716.1| PREDICTED: vacuolar protein sorting-associated protein 26 [
>gi|1036724711|ref|XP_017146740.1| PREDICTED: ribosomal protein 63, mitochondrial [Drosophila
>gi|1036739662|ref|XP_017155117.1| PREDICTED: required for meiotic nuclear division protein 1
>gi|1036739648|ref|XP_017155106.1| PREDICTED: anaphase-promoting complex subunit 7 [Drosophila
>gi|1036708557|ref|XP_017137723.1| PREDICTED: uncharacterized protein LOC108152708 isoform X1
>gi|1036708577|ref|XP_017137733.1| PREDICTED: uncharacterized protein LOC108152708 isoform X1
>gi|1036708594|ref|XP_017137741.1| PREDICTED: uncharacterized protein LOC108152708 isoform X2
>gi|1036708685|ref|XP_017137795.1| PREDICTED: calcyclin-binding protein [Drosophila miranda]
>gi|1036708668|ref|XP_017137782.1| PREDICTED: protein swallow [Drosophila miranda]
>gi|1036708611|ref|XP_017137753.1| PREDICTED: cell division cycle 7-related protein kinase [Dro
>gi|1036708630|ref|XP_017137762.1| PREDICTED: cell division cycle 7-related protein kinase [Dro
>gi|1036708651|ref|XP_017137769.1| PREDICTED: cell division cycle 7-related protein kinase [Dro
>gi|1036708766|ref|XP_017137847.1| PREDICTED: zinc finger protein 593 homolog [Drosophila mira
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>gi|1036708702|ref|XP_017137808.1| PREDICTED: high affinity copper uptake protein 1 isoform X1
>gi|1036708719|ref|XP_017137816.1| PREDICTED: high affinity copper uptake protein 1 isoform X1
>gi|1036708733|ref|XP_017137826.1| PREDICTED: high affinity copper uptake protein 1 isoform X1
>gi|1036708750|ref|XP_017137834.1| PREDICTED: high affinity copper uptake protein 1 isoform X2
>gi|1036708814|ref|XP_017137869.1| PREDICTED: luc7-like protein 3 [Drosophila miranda]
>gi|1036735172|ref|XP_017152624.1| PREDICTED: DNA replication licensing factor Mcm6 [Drosophila
>gi|1036735189|ref|XP_017152635.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036713484|ref|XP_017140666.1| PREDICTED: fumarate hydratase, mitochondrial isoform X1 [Dro
>gi|1036713500|ref|XP_017140674.1| PREDICTED: fumarate hydratase, mitochondrial isoform X2 [Dro
>gi|1036713472|ref|XP_017140655.1| PREDICTED: fumarate hydratase, mitochondrial-like [Drosophil
>gi|1036713457|ref|XP_017140643.1| PREDICTED: uncharacterized protein LOC108154768 [Drosophila
>gi|1036713537|ref|XP_017140699.1| PREDICTED: uncharacterized protein LOC108154807 [Drosophila
>gi|1036713516|ref|XP_017140687.1| PREDICTED: diphthine methyltransferase [Drosophila miranda]
>gi|1036713440|ref|XP_017140632.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila m
>gi|1036713424|ref|XP_017140622.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila m
>gi|1036713569|ref|XP_017140717.1| PREDICTED: transcription factor Maf [Drosophila miranda]
>gi|1036704626|ref|XP_017135013.1| PREDICTED: uncharacterized protein LOC108151114 [Drosophila
>gi|1036704640|ref|XP_017135023.1| PREDICTED: uncharacterized protein LOC108151114 [Drosophila
>gi|1036698225|ref|XP_017134906.1| PREDICTED: breakpoint cluster region protein [Drosophila mi
>gi|1036704659|ref|XP_017135035.1| PREDICTED: probable RNA-binding protein 19 [Drosophila mira
>gi|1036704714|ref|XP_017135067.1| PREDICTED: uncharacterized protein LOC108151149 [Drosophila
>gi|1036698240|ref|XP_017136144.1| PREDICTED: uncharacterized protein LOC108151567 [Drosophila
>gi|1036704677|ref|XP_017135047.1| PREDICTED: uncharacterized protein LOC108151131 [Drosophila
>gi|1036704697|ref|XP_017135055.1| PREDICTED: beta-1,3-galactosyltransferase brn [Drosophila m
>gi|1036704549|ref|XP_017134967.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704564|ref|XP_017134976.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704579|ref|XP_017134983.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704598|ref|XP_017134990.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704613|ref|XP_017134999.1| PREDICTED: histone-lysine N-methyltransferase EHMT2 isoform
>gi|1036702005|ref|XP_017156117.1| PREDICTED: uncharacterized oxidoreductase SSP0419 [Drosophil
>gi|1036702021|ref|XP_017156125.1| PREDICTED: uncharacterized oxidoreductase SSP0419 [Drosophil
>gi|1036701898|ref|XP_017156051.1| PREDICTED: molybdenum cofactor synthesis protein cinnamon [
>gi|1036702165|ref|XP_017156206.1| PREDICTED: cytochrome c oxidase assembly factor 6 homolog [
>gi|1036702042|ref|XP_017156137.1| PREDICTED: uncharacterized protein LOC108164748 [Drosophila
>gi|1036701865|ref|XP_017156030.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofo
>gi|1036701881|ref|XP_017156038.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofo
>gi|1036701833|ref|XP_017156014.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofo
>gi|1036701848|ref|XP_017156023.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofo
>gi|1036702148|ref|XP_017156196.1| PREDICTED: RING-box protein 1A [Drosophila miranda]
>gi|1036701989|ref|XP_017156104.1| PREDICTED: uncharacterized protein LOC108164722 [Drosophila
>gi|1036701915|ref|XP_017156063.1| PREDICTED: protein SPT2 homolog [Drosophila miranda]
>gi|1036701951|ref|XP_017156084.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X2 [Droso
>gi|1036701932|ref|XP_017156075.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X1 [Droso
>gi|1036701970|ref|XP_017156092.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X3 [Droso
>gi|1036702059|ref|XP_017156150.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702076|ref|XP_017156160.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702093|ref|XP_017156167.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702114|ref|XP_017156176.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
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>gi|1036702131|ref|XP_017156185.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036698410|ref|XP_017150823.1| PREDICTED: uncharacterized protein LOC108160753 [Drosophila
>gi|1036707387|ref|XP_017136928.1| PREDICTED: putative odorant receptor 92a [Drosophila miranda
>gi|1036705458|ref|XP_017135617.1| PREDICTED: uncharacterized protein LOC108151486 [Drosophila
>gi|1036703751|ref|XP_017157128.1| PREDICTED: uncharacterized protein LOC108165562 [Drosophila
>gi|1036713104|ref|XP_017140430.1| PREDICTED: protein sidekick isoform X1 [Drosophila miranda]
>gi|1036713121|ref|XP_017140436.1| PREDICTED: protein sidekick isoform X2 [Drosophila miranda]
>gi|1036713144|ref|XP_017140443.1| PREDICTED: protein sidekick isoform X3 [Drosophila miranda]
>gi|1036713161|ref|XP_017140449.1| PREDICTED: protein sidekick isoform X4 [Drosophila miranda]
>gi|1036713177|ref|XP_017140458.1| PREDICTED: protein sidekick isoform X5 [Drosophila miranda]
>gi|1036713194|ref|XP_017140466.1| PREDICTED: protein sidekick isoform X6 [Drosophila miranda]
>gi|1036713211|ref|XP_017140475.1| PREDICTED: protein sidekick isoform X7 [Drosophila miranda]
>gi|1036704270|ref|XP_017134784.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036698425|ref|XP_017151896.1| PREDICTED: uncharacterized protein LOC108161692 [Drosophila
>gi|1036755177|ref|XP_017156601.1| PREDICTED: mitochondrial 2-oxodicarboxylate carrier [Drosoph
>gi|1036751123|ref|XP_017156360.1| PREDICTED: beta-amyloid-like protein isoform X1 [Drosophila
>gi|1036751141|ref|XP_017156361.1| PREDICTED: beta-amyloid-like protein isoform X1 [Drosophila
>gi|1036751215|ref|XP_017156365.1| PREDICTED: beta-amyloid-like protein isoform X5 [Drosophila
>gi|1036751197|ref|XP_017156364.1| PREDICTED: beta-amyloid-like protein isoform X4 [Drosophila
>gi|1036751179|ref|XP_017156363.1| PREDICTED: beta-amyloid-like protein isoform X3 [Drosophila
>gi|1036751234|ref|XP_017156366.1| PREDICTED: beta-amyloid-like protein isoform X6 [Drosophila
>gi|1036751160|ref|XP_017156362.1| PREDICTED: beta-amyloid-like protein isoform X2 [Drosophila
>gi|1036751251|ref|XP_017156367.1| PREDICTED: beta-amyloid-like protein isoform X7 [Drosophila
>gi|1036751269|ref|XP_017156369.1| PREDICTED: beta-amyloid-like protein isoform X8 [Drosophila
>gi|1036698440|ref|XP_017153096.1| PREDICTED: uncharacterized protein LOC108162461 [Drosophila
>gi|1036745739|ref|XP_017156066.1| PREDICTED: homeobox protein vnd isoform X2 [Drosophila miran
>gi|1036745721|ref|XP_017156065.1| PREDICTED: homeobox protein vnd isoform X1 [Drosophila miran
>gi|1036729959|ref|XP_017149863.1| PREDICTED: cytospin-A isoform X2 [Drosophila miranda]
>gi|1036729943|ref|XP_017149855.1| PREDICTED: cytospin-A isoform X1 [Drosophila miranda]
>gi|1036746629|ref|XP_017156120.1| PREDICTED: uncharacterized protein LOC108164738 [Drosophila
>gi|1036746612|ref|XP_017156119.1| PREDICTED: vacuolar protein sorting-associated protein 37C
>gi|1036753497|ref|XP_017156502.1| PREDICTED: ribokinase [Drosophila miranda]
>gi|1036708830|ref|XP_017137879.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036708846|ref|XP_017137887.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036708862|ref|XP_017137893.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036708879|ref|XP_017137902.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036708896|ref|XP_017137911.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036708913|ref|XP_017137920.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036708949|ref|XP_017137945.1| PREDICTED: histone acetyltransferase Tip60 [Drosophila miran
>gi|1036708964|ref|XP_017137958.1| PREDICTED: uncharacterized protein LOC108152849 [Drosophila
>gi|1036708930|ref|XP_017137932.1| PREDICTED: tRNA (cytosine(34)-C(5))-methyltransferase [Droso
>gi|1036708981|ref|XP_017137971.1| PREDICTED: uncharacterized protein LOC108152857 [Drosophila
>gi|1036749808|ref|XP_017156286.1| PREDICTED: beta-1,3-galactosyltransferase brn [Drosophila m
>gi|1036740489|ref|XP_017155531.1| PREDICTED: DNA repair protein complementing XP-A cells homol
>gi|1036740455|ref|XP_017155506.1| PREDICTED: uncharacterized protein LOC108164395 [Drosophila
>gi|1036740472|ref|XP_017155516.1| PREDICTED: uncharacterized protein LOC108164395 [Drosophila
>gi|1036735086|ref|XP_017152569.1| PREDICTED: T-complex protein 1 subunit zeta-like [Drosophila
>gi|1036735103|ref|XP_017152582.1| PREDICTED: phosphatidylinositol 4,5-bisphosphate 5-phosphata
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>gi|1036750227|ref|XP_017156309.1| PREDICTED: ubiquitin-conjugating enzyme E2 G2 [Drosophila m:
>gi|1036721608|ref|XP_017145191.1| PREDICTED: WD repeat-containing protein 18 [Drosophila mira:
>gi|1036721646|ref|XP_017145215.1| PREDICTED: UPF0184 protein CG14818 [Drosophila miranda]
>gi|1036721625|ref|XP_017145204.1| PREDICTED: APOPT family protein CG14806, mitochondrial [Dros
>gi|1036721591|ref|XP_017145180.1| PREDICTED: histone-lysine N-methyltransferase trr [Drosophil
>gi|1036708117|ref|XP_017137433.1| PREDICTED: 39S ribosomal protein L16, mitochondrial isoform
>gi|1036708134|ref|XP_017137440.1| PREDICTED: 39S ribosomal protein L16, mitochondrial isoform
>gi|1036708032|ref|XP_017137380.1| PREDICTED: armadillo segment polarity protein isoform X1 [D:
>gi|1036708049|ref|XP_017137389.1| PREDICTED: armadillo segment polarity protein isoform X1 [D:
>gi|1036708066|ref|XP_017137398.1| PREDICTED: armadillo segment polarity protein isoform X1 [D:
>gi|1036708083|ref|XP_017137407.1| PREDICTED: armadillo segment polarity protein isoform X2 [D:
>gi|1036708151|ref|XP_017137451.1| PREDICTED: uncharacterized protein LOC108152546 isoform X1
>gi|1036708167|ref|XP_017137461.1| PREDICTED: uncharacterized protein LOC108152546 isoform X2
>gi|1036708182|ref|XP_017137474.1| PREDICTED: uncharacterized protein LOC108152557 isoform X1
>gi|1036708199|ref|XP_017137484.1| PREDICTED: uncharacterized protein LOC108152557 isoform X2
>gi|1036707981|ref|XP_017137343.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036707947|ref|XP_017137326.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036707965|ref|XP_017137335.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036707930|ref|XP_017137314.1| PREDICTED: WD repeat-containing protein 7 [Drosophila miran
>gi|1036707997|ref|XP_017137355.1| PREDICTED: protein lin-9 homolog [Drosophila miranda]
>gi|1036708100|ref|XP_017137419.1| PREDICTED: translation initiation factor eIF-2B subunit eps
>gi|1036708013|ref|XP_017137366.1| PREDICTED: type II inositol 1,4,5-trisphosphate 5-phosphata
>gi|1036717028|ref|XP_017142775.1| PREDICTED: synaptobrevin homolog YKT6 [Drosophila miranda]
>gi|1036716881|ref|XP_017142683.1| PREDICTED: protein hold'em isoform X1 [Drosophila miranda]
>gi|1036716898|ref|XP_017142691.1| PREDICTED: protein hold'em isoform X2 [Drosophila miranda]
>gi|1036716864|ref|XP_017142672.1| PREDICTED: sorting nexin-25 [Drosophila miranda]
>gi|1036716994|ref|XP_017142754.1| PREDICTED: very-long-chain 3-oxoacyl-CoA reductase-B [Droso
>gi|1036717013|ref|XP_017142763.1| PREDICTED: very-long-chain 3-oxoacyl-CoA reductase-B [Droso
>gi|1036716937|ref|XP_017142720.1| PREDICTED: fibroblast growth factor receptor 3 [Drosophila
>gi|1036716956|ref|XP_017142729.1| PREDICTED: fibroblast growth factor receptor 3 [Drosophila
>gi|1036716975|ref|XP_017142739.1| PREDICTED: fibroblast growth factor receptor 3 [Drosophila
>gi|1036716918|ref|XP_017142704.1| PREDICTED: bystin [Drosophila miranda]
>gi|1036749133|ref|XP_017156249.1| PREDICTED: 40S ribosomal protein S6 [Drosophila miranda]
>gi|1036742788|ref|XP_017155905.1| PREDICTED: general vesicular transport factor p115 [Drosoph:
>gi|1036724795|ref|XP_017146802.1| PREDICTED: uncharacterized protein LOC108158725 [Drosophila
>gi|1036724812|ref|XP_017146818.1| PREDICTED: N(G),N(G)-dimethylarginine dimethylaminohydrolase
>gi|1036728008|ref|XP_017148715.1| PREDICTED: cyclin-L1 [Drosophila miranda]
>gi|1036728027|ref|XP_017148722.1| PREDICTED: cyclin-L1 [Drosophila miranda]
>gi|1036728095|ref|XP_017148769.1| PREDICTED: DNA damage-regulated autophagy modulator protein
>gi|1036728040|ref|XP_017148734.1| PREDICTED: RNA 3'-terminal phosphate cyclase [Drosophila mi:
>gi|1036728059|ref|XP_017148748.1| PREDICTED: tRNA (guanine-N(7)-)-methyltransferase [Drosophil
>gi|1036728076|ref|XP_017148757.1| PREDICTED: tRNA (guanine-N(7)-)-methyltransferase [Drosophil
>gi|1036728114|ref|XP_017148782.1| PREDICTED: uncharacterized protein LOC108159749 [Drosophila
>gi|1036742463|ref|XP_017155887.1| PREDICTED: fasciculation and elongation protein zeta-2 [Dros
>gi|1036735539|ref|XP_017152824.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X3
>gi|1036735520|ref|XP_017152816.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X2
>gi|1036735502|ref|XP_017152808.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X1
>gi|1036735558|ref|XP_017152833.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X4
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>gi|1036745758|ref|XP_017156067.1| PREDICTED: uncharacterized protein LOC108164703 [Drosophila
>gi|1036749989|ref|XP_017156297.1| PREDICTED: AT-rich binding protein [Drosophila miranda]
>gi|1036717424|ref|XP_017142996.1| PREDICTED: uncharacterized protein LOC108156174 [Drosophila
>gi|1036717443|ref|XP_017143007.1| PREDICTED: protein suppressor of forked [Drosophila miranda]
>gi|1036717461|ref|XP_017143015.1| PREDICTED: protein suppressor of forked [Drosophila miranda]
>gi|1036698454|ref|XP_017154303.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036717511|ref|XP_017143038.1| PREDICTED: uncharacterized protein LOC108156208 [Drosophila
>gi|1036717530|ref|XP_017143050.1| PREDICTED: uncharacterized protein LOC108156220 [Drosophila
>gi|1036717480|ref|XP_017143027.1| PREDICTED: smad nuclear-interacting protein 1 [Drosophila m
>gi|1036751937|ref|XP_017156405.1| PREDICTED: uncharacterized protein LOC108164928 [Drosophila
>gi|1036756860|ref|XP_017156694.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X1
>gi|1036756876|ref|XP_017156695.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X2
>gi|1036757154|ref|XP_017156708.1| PREDICTED: sodium/potassium/calcium exchanger 4 [Drosophila
>gi|1036707741|ref|XP_017137192.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X1
>gi|1036707777|ref|XP_017137210.1| PREDICTED: sodium/potassium/calcium exchanger 4 isoform X3
>gi|1036707758|ref|XP_017137201.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X2
>gi|1036707843|ref|XP_017137261.1| PREDICTED: cytochrome c oxidase assembly factor 7 homolog [
>gi|1036698468|ref|XP_017155452.1| PREDICTED: NAD-dependent protein deacylase Sirt4 [Drosophila
>gi|1036707671|ref|XP_017137148.1| PREDICTED: RNA-binding protein 25 isoform X1 [Drosophila mi
>gi|1036707688|ref|XP_017137157.1| PREDICTED: RNA-binding protein 25 isoform X2 [Drosophila mi
>gi|1036707870|ref|XP_017137276.1| PREDICTED: ras-related protein Rab-18 [Drosophila miranda]
>gi|1036707889|ref|XP_017137289.1| PREDICTED: 60S ribosomal protein L35 [Drosophila miranda]
>gi|1036707794|ref|XP_017137222.1| PREDICTED: protein trapped in endoderm-1 [Drosophila miranda
>gi|1036698483|ref|XP_017155879.1| PREDICTED: LOW QUALITY PROTEIN: gustatory receptor 5a for t
>gi|1036698498|ref|XP_017155992.1| PREDICTED: sulfated surface glycoprotein 185 [Drosophila mi
>gi|1036707707|ref|XP_017137170.1| PREDICTED: GPI inositol-deacylase [Drosophila miranda]
>gi|1036707826|ref|XP_017137249.1| PREDICTED: protein twisted gastrulation [Drosophila miranda]
>gi|1036707724|ref|XP_017137182.1| PREDICTED: paraplegin [Drosophila miranda]
>gi|1036707809|ref|XP_017137235.1| PREDICTED: circadian clock-controlled protein [Drosophila m
>gi|1036707635|ref|XP_017137127.1| PREDICTED: period circadian protein isoform X2 [Drosophila
>gi|1036707620|ref|XP_017137119.1| PREDICTED: period circadian protein isoform X1 [Drosophila
>gi|1036707654|ref|XP_017137135.1| PREDICTED: period circadian protein isoform X3 [Drosophila
>gi|1036698513|ref|XP_017156109.1| PREDICTED: uncharacterized protein LOC108164705 [Drosophila
>gi|1036756602|ref|XP_017156680.1| PREDICTED: tryptophan 2,3-dioxygenase isoform X1 [Drosophila
>gi|1036756619|ref|XP_017156681.1| PREDICTED: tryptophan 2,3-dioxygenase isoform X2 [Drosophila
>gi|1036751436|ref|XP_017156377.1| PREDICTED: poly(U)-specific endoribonuclease homolog isoform
>gi|1036751455|ref|XP_017156378.1| PREDICTED: poly(U)-specific endoribonuclease homolog isoform
>gi|1036735782|ref|XP_017152955.1| PREDICTED: unconventional myosin-XV isoform X1 [Drosophila
>gi|1036735799|ref|XP_017152964.1| PREDICTED: uncharacterized protein LOC108162639 isoform X2
>gi|1036698528|ref|XP_017156222.1| PREDICTED: tenascin-X [Drosophila miranda]
>gi|1036750155|ref|XP_017156304.1| PREDICTED: uncharacterized protein LOC108164872 [Drosophila
>gi|1036750174|ref|XP_017156306.1| PREDICTED: uncharacterized protein LOC108164872 [Drosophila
>gi|1036750190|ref|XP_017156307.1| PREDICTED: uncharacterized protein LOC108164873 [Drosophila
>gi|1036698545|ref|XP_017156336.1| PREDICTED: uncharacterized protein LOC108164869 [Drosophila
>gi|1036754905|ref|XP_017156586.1| PREDICTED: protein msta isoform X1 [Drosophila miranda]
>gi|1036754924|ref|XP_017156587.1| PREDICTED: protein msta isoform X2 [Drosophila miranda]
>gi|1036698562|ref|XP_017156464.1| PREDICTED: polycystic kidney disease 2-like 2 protein [Dros
>gi|1036698577|ref|XP_017156582.1| PREDICTED: uncharacterized protein LOC108165018 [Drosophila
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>gi|1036704892|ref|XP_017135217.1| PREDICTED: uncharacterized protein CG1552 isoform X2 [Droso
>gi|1036704876|ref|XP_017135209.1| PREDICTED: uncharacterized protein CG1552 isoform X1 [Droso
>gi|1036698592|ref|XP_017156696.1| PREDICTED: surface protein-like [Drosophila miranda]
>gi|1036705118|ref|XP_017135393.1| PREDICTED: uncharacterized protein LOC108151345 [Drosophila
>gi|1036705865|ref|XP_017135889.1| PREDICTED: uncharacterized protein LOC108151664 [Drosophila
>gi|1036703770|ref|XP_017157149.1| PREDICTED: uncharacterized protein LOC108165584 [Drosophila
>gi|1036741499|ref|XP_017155834.1| PREDICTED: protein sevenless isoform X2 [Drosophila miranda]
>gi|1036741462|ref|XP_017155831.1| PREDICTED: protein sevenless isoform X1 [Drosophila miranda]
>gi|1036741481|ref|XP_017155833.1| PREDICTED: protein sevenless isoform X1 [Drosophila miranda]
>gi|1036735870|ref|XP_017153010.1| PREDICTED: uncharacterized protein LOC108162674 isoform X1
>gi|1036735890|ref|XP_017153018.1| PREDICTED: uncharacterized protein LOC108162674 isoform X2
>gi|1036735851|ref|XP_017152998.1| PREDICTED: growth hormone-inducible transmembrane protein [
>gi|1036735815|ref|XP_017152977.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036735832|ref|XP_017152986.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036755417|ref|XP_017156615.1| PREDICTED: apyrase isoform X2 [Drosophila miranda]
>gi|1036755398|ref|XP_017156614.1| PREDICTED: apyrase isoform X1 [Drosophila miranda]
>gi|1036710670|ref|XP_017139039.1| PREDICTED: RNA polymerase II subunit A C-terminal domain pho
>gi|1036710653|ref|XP_017139026.1| PREDICTED: glucose-6-phosphate 1-dehydrogenase [Drosophila r
>gi|1036710620|ref|XP_017139002.1| PREDICTED: cytokine receptor isoform X2 [Drosophila miranda]
>gi|1036710602|ref|XP_017138993.1| PREDICTED: cytokine receptor isoform X1 [Drosophila miranda]
>gi|1036710585|ref|XP_017138981.1| PREDICTED: cytokine receptor [Drosophila miranda]
>gi|1036710636|ref|XP_017139013.1| PREDICTED: ubiquilin-1 [Drosophila miranda]
>gi|1036742409|ref|XP_017155884.1| PREDICTED: moesin/ezrin/radixin homolog 2 [Drosophila miranc
>gi|1036752212|ref|XP_017156422.1| PREDICTED: uncharacterized protein LOC108164944 isoform X2
>gi|1036752191|ref|XP_017156421.1| PREDICTED: uncharacterized protein LOC108164944 isoform X1
>gi|1036744272|ref|XP_017155982.1| PREDICTED: cdc42 homolog [Drosophila miranda]
>gi|1036743892|ref|XP_017155963.1| PREDICTED: probable tRNA N6-adenosine threonylcarbamoyltrans
>gi|1036743875|ref|XP_017155962.1| PREDICTED: Golgi resident protein GCP60 [Drosophila miranda]
>gi|1036736935|ref|XP_017153635.1| PREDICTED: probable RNA-binding protein CG14230 [Drosophila
>gi|1036736951|ref|XP_017153648.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036736968|ref|XP_017153655.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036736986|ref|XP_017153664.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036737004|ref|XP_017153672.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036754508|ref|XP_017156561.1| PREDICTED: high affinity cationic amino acid transporter 1
>gi|1036748701|ref|XP_017156226.1| PREDICTED: chorion peroxidase [Drosophila miranda]
>gi|1036750387|ref|XP_017156316.1| PREDICTED: dipeptidyl aminopeptidase-like protein 6 isoform
>gi|1036750368|ref|XP_017156315.1| PREDICTED: dipeptidyl aminopeptidase-like protein 6 isoform
>gi|1036750407|ref|XP_017156317.1| PREDICTED: uncharacterized protein LOC108164881 [Drosophila
>gi|1036734133|ref|XP_017152045.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X2
>gi|1036734076|ref|XP_017152018.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X1
>gi|1036734092|ref|XP_017152029.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X1
>gi|1036734114|ref|XP_017152037.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X1
>gi|1036749674|ref|XP_017156279.1| PREDICTED: uncharacterized protein LOC108164860 isoform X1
>gi|1036749693|ref|XP_017156280.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036749712|ref|XP_017156281.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036749731|ref|XP_017156282.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036749749|ref|XP_017156283.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036698606|ref|XP_017156772.1| PREDICTED: uncharacterized protein LOC108165203 [Drosophila
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>gi|1036757208|ref|XP_017156711.1| PREDICTED: uncharacterized protein LOC108165156 [Drosophila
>gi|1036733239|ref|XP_017151520.1| PREDICTED: proton-coupled amino acid transporter 4 isoform X
>gi|1036733292|ref|XP_017151545.1| PREDICTED: proton-coupled amino acid transporter 1 [Drosophi
>gi|1036733255|ref|XP_017151527.1| PREDICTED: proton-coupled amino acid transporter 4 isoform X
>gi|1036733273|ref|XP_017151535.1| PREDICTED: proton-coupled amino acid transporter 4 isoform X
>gi|1036733347|ref|XP_017151579.1| PREDICTED: actin-related protein 2/3 complex subunit 3 [Dros
>gi|1036733329|ref|XP_017151569.1| PREDICTED: LOW QUALITY PROTEIN: rab-like protein 3 [Drosophi
>gi|1036733311|ref|XP_017151554.1| PREDICTED: decapping nuclease DXO homolog [Drosophila mirand
>gi|1036744610|ref|XP_017156000.1| PREDICTED: uncharacterized protein LOC108164657 isoform X2
>gi|1036744591|ref|XP_017155999.1| PREDICTED: uncharacterized protein LOC108164657 isoform X1
>gi|1036698621|ref|XP_017156875.1| PREDICTED: uncharacterized protein LOC108165318 [Drosophila
>gi|1036711462|ref|XP_017139492.1| PREDICTED: TM2 domain-containing protein almondex [Drosophil
>gi|1036711445|ref|XP_017139482.1| PREDICTED: la-related protein 7 [Drosophila miranda]
>gi|1036711340|ref|XP_017139437.1| PREDICTED: uncharacterized protein LOC108153846 isoform X1
>gi|1036711357|ref|XP_017139442.1| PREDICTED: uncharacterized protein LOC108153846 isoform X2
>gi|1036711479|ref|XP_017139501.1| PREDICTED: uncharacterized protein LOC108153913 [Drosophila
>gi|1036711374|ref|XP_017139452.1| PREDICTED: protein GPR107 isoform X1 [Drosophila miranda]
>gi|1036711393|ref|XP_017139460.1| PREDICTED: protein GPR107 isoform X2 [Drosophila miranda]
>gi|1036711408|ref|XP_017139468.1| PREDICTED: protein GPR107 isoform X2 [Drosophila miranda]
>gi|1036711428|ref|XP_017139474.1| PREDICTED: protein GPR107 isoform X2 [Drosophila miranda]
>gi|1036698635|ref|XP_017156987.1| PREDICTED: gustatory receptor 8a [Drosophila miranda]
>gi|1036698651|ref|XP_017157107.1| PREDICTED: uncharacterized protein LOC108165520 [Drosophila
>gi|1036752267|ref|XP_017156425.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036752282|ref|XP_017156426.1| PREDICTED: uncharacterized protein LOC108164948 [Drosophila
>gi|1036752300|ref|XP_017156428.1| PREDICTED: uncharacterized protein LOC108164948 [Drosophila
>gi|1036752320|ref|XP_017156429.1| PREDICTED: uncharacterized protein LOC108164948 [Drosophila
>gi|1036754740|ref|XP_017156576.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036754756|ref|XP_017156577.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036754771|ref|XP_017156578.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036754787|ref|XP_017156579.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036743966|ref|XP_017155967.1| PREDICTED: kinesin-associated protein 3 [Drosophila miranda]
>gi|1036730760|ref|XP_017150382.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036730779|ref|XP_017150390.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036730797|ref|XP_017150400.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036730815|ref|XP_017150408.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036698667|ref|XP_017134572.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036740943|ref|XP_017155800.1| PREDICTED: uncharacterized protein C1683.06c [Drosophila mi
>gi|1036740959|ref|XP_017155801.1| PREDICTED: inosine-uridine preferring nucleoside hydrolase
>gi|1036714435|ref|XP_017141260.1| PREDICTED: probable glutamine-dependent NAD(+) synthetase [
>gi|1036714235|ref|XP_017141152.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X1 [Drosc
>gi|1036714273|ref|XP_017141171.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X3 [Drosc
>gi|1036714345|ref|XP_017141205.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X7 [Drosc
>gi|1036714290|ref|XP_017141181.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X4 [Drosc
>gi|1036714309|ref|XP_017141190.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X5 [Drosc
>gi|1036714327|ref|XP_017141199.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X6 [Drosc
>gi|1036714254|ref|XP_017141160.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X2 [Drosc
>gi|1036714363|ref|XP_017141214.1| PREDICTED: uncharacterized protein LOC108155072 isoform X8
>gi|1036714380|ref|XP_017141222.1| PREDICTED: uncharacterized protein LOC108155072 isoform X9
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>gi|1036714416|ref|XP_017141248.1| PREDICTED: RING finger protein 157 [Drosophila miranda]
>gi|1036714395|ref|XP_017141236.1| PREDICTED: DNA topoisomerase 2-binding protein 1-A [Drosop
>gi|1036739911|ref|XP_017155258.1| PREDICTED: AP-3 complex subunit delta [Drosophila miranda]
>gi|1036731592|ref|XP_017150790.1| PREDICTED: gamma-tubulin complex component 3 homolog [Drosop
>gi|1036731609|ref|XP_017150801.1| PREDICTED: probable methylthioribulose-1-phosphate dehydrata
>gi|1036731626|ref|XP_017150813.1| PREDICTED: peroxisomal multifunctional enzyme type 2 [Drosop
>gi|1036711983|ref|XP_017139800.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036711932|ref|XP_017139760.1| PREDICTED: programmed cell death protein 4 [Drosophila mira
>gi|1036712017|ref|XP_017139818.1| PREDICTED: gametocyte-specific factor 1 homolog isoform X2
>gi|1036712031|ref|XP_017139828.1| PREDICTED: gametocyte-specific factor 1 isoform X3 [Drosophi
>gi|1036712000|ref|XP_017139812.1| PREDICTED: gametocyte-specific factor 1 homolog isoform X1
>gi|1036711966|ref|XP_017139787.1| PREDICTED: probable RNA 3'-terminal phosphate cyclase-like p
>gi|1036711949|ref|XP_017139770.1| PREDICTED: vitellogenin-3 [Drosophila miranda]
>gi|1036711881|ref|XP_017139731.1| PREDICTED: protein retinal degeneration B isoform X3 [Drosop
>gi|1036711897|ref|XP_017139739.1| PREDICTED: protein retinal degeneration B isoform X4 [Drosop
>gi|1036711914|ref|XP_017139748.1| PREDICTED: protein retinal degeneration B isoform X5 [Drosop
>gi|1036711829|ref|XP_017139707.1| PREDICTED: protein retinal degeneration B isoform X1 [Drosop
>gi|1036711847|ref|XP_017139715.1| PREDICTED: protein retinal degeneration B isoform X1 [Drosop
>gi|1036711864|ref|XP_017139723.1| PREDICTED: protein retinal degeneration B isoform X2 [Drosop
>gi|1036746543|ref|XP_017156114.1| PREDICTED: cathepsin B [Drosophila miranda]
>gi|1036702340|ref|XP_017156305.1| PREDICTED: TM2 domain-containing protein CG11103 [Drosophila
>gi|1036721009|ref|XP_017144895.1| PREDICTED: probable tubulin polyglutamylase TTLL1 isoform X2
>gi|1036721046|ref|XP_017144916.1| PREDICTED: myb-like protein I [Drosophila miranda]
>gi|1036720990|ref|XP_017144886.1| PREDICTED: probable tubulin polyglutamylase TTLL1 isoform X
>gi|1036720953|ref|XP_017144868.1| PREDICTED: partitioning defective 3 homolog isoform X1 [Dros
>gi|1036720971|ref|XP_017144875.1| PREDICTED: partitioning defective 3 homolog isoform X2 [Dros
>gi|1036721028|ref|XP_017144902.1| PREDICTED: probable tubulin polyglutamylase TTLL1 isoform X3
>gi|1036706297|ref|XP_017136210.1| PREDICTED: keratin-associated protein 6-2 [Drosophila miranc
>gi|1036757481|ref|XP_017156729.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1036707913|ref|XP_017137303.1| PREDICTED: uncharacterized protein LOC108152451 [Drosophila
>gi|1036698686|ref|XP_017134677.1| PREDICTED: uncharacterized protein LOC108150846, partial [D
>gi|1036698703|ref|XP_017134789.1| PREDICTED: phospholipase A1 2-like [Drosophila miranda]
>gi|1036753104|ref|XP_017156478.1| PREDICTED: probable phospholipase A1 magnifin [Drosophila m
>gi|1036753087|ref|XP_017156477.1| PREDICTED: vitellogenin-3 [Drosophila miranda]
>gi|1036707490|ref|XP_017136991.1| PREDICTED: heterogeneous nuclear ribonucleoprotein A1 [Dros
>gi|1036706066|ref|XP_017136048.1| PREDICTED: glycine-rich cell wall structural protein 1.0 [D
>gi|1036705208|ref|XP_017135451.1| PREDICTED: glycine-rich cell wall structural protein 2 [Dros
>gi|1036707604|ref|XP_017137094.1| PREDICTED: ctenidin-1 [Drosophila miranda]
>gi|1036698719|ref|XP_017134907.1| PREDICTED: glycine-rich cell wall structural protein 1.0 [D
>gi|1036737299|ref|XP_017153840.1| PREDICTED: DExH-box ATP-dependent RNA helicase DExH6 [Drosop
>gi|1036737317|ref|XP_017153852.1| PREDICTED: uncharacterized protein C9orf85 homolog [Drosophi
>gi|1036737249|ref|XP_017153811.1| PREDICTED: uncharacterized protein LOC108163182 isoform X1
>gi|1036737267|ref|XP_017153820.1| PREDICTED: myb-like protein I isoform X2 [Drosophila miranda
>gi|1036737281|ref|XP_017153828.1| PREDICTED: espin isoform X3 [Drosophila miranda]
>gi|1036704126|ref|XP_017134701.1| PREDICTED: uncharacterized protein LOC108150899 [Drosophila
>gi|1036704439|ref|XP_017134894.1| PREDICTED: uncharacterized protein LOC108151036 [Drosophila
>gi|1036733525|ref|XP_017151683.1| PREDICTED: plastin-1 isoform X1 [Drosophila miranda]
>gi|1036733543|ref|XP_017151691.1| PREDICTED: plastin-1 isoform X2 [Drosophila miranda]
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>gi|1036746470|ref|XP_017156110.1| PREDICTED: uncharacterized protein C6orf106 homolog [Drosopl
>gi|1036754219|ref|XP_017156545.1| PREDICTED: homeobox protein B-H2 [Drosophila miranda]
>gi|1036755435|ref|XP_017156616.1| PREDICTED: homeobox protein B-H1 isoform X1 [Drosophila mira
>gi|1036755454|ref|XP_017156617.1| PREDICTED: homeobox protein B-H1 isoform X2 [Drosophila mira
>gi|1036704924|ref|XP_017135242.1| PREDICTED: E3 SUMO-protein ligase PIAS1-like [Drosophila mir
>gi|1036735263|ref|XP_017152676.1| PREDICTED: probable ATP-dependent RNA helicase CG8611 [Dros
>gi|1036707111|ref|XP_017136761.1| PREDICTED: uncharacterized protein LOC108152152 [Drosophila
>gi|1036741379|ref|XP_017155826.1| PREDICTED: cytosolic endo-beta-N-acetylglucosaminidase [Dros
>gi|1036741396|ref|XP_017155827.1| PREDICTED: cytosolic endo-beta-N-acetylglucosaminidase [Dros
>gi|1036741445|ref|XP_017155830.1| PREDICTED: uncharacterized protein LOC108164539 isoform X2
>gi|1036741426|ref|XP_017155829.1| PREDICTED: uncharacterized protein LOC108164539 isoform X1
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>gi|1036757284|ref|XP_017156716.1| PREDICTED: uncharacterized protein LOC108165160 [Drosophila
>gi|1036698734|ref|XP_017135031.1| PREDICTED: acidic leucine-rich nuclear phosphoprotein 32-re
>gi|1036747464|ref|XP_017156170.1| PREDICTED: ubiquitin-like modifier-activating enzyme 5 [Dros
>gi|1036747798|ref|XP_017156190.1| PREDICTED: signal peptidase complex subunit 2 [Drosophila m
>gi|1036747779|ref|XP_017156189.1| PREDICTED: probable tubulin beta chain CG32396 [Drosophila r
>gi|1036710810|ref|XP_017139125.1| PREDICTED: death-associated protein kinase related-like [Dro
>gi|1036710827|ref|XP_017139133.1| PREDICTED: death-associated protein kinase related-like [Dro
>gi|1036698751|ref|XP_017135152.1| PREDICTED: death-associated protein kinase related, partial
>gi|1036698770|ref|XP_017135281.1| PREDICTED: cytochrome P450 4g15-like, partial [Drosophila m
>gi|1036704155|ref|XP_017134724.1| PREDICTED: uncharacterized protein LOC108150906 isoform X2
>gi|1036704141|ref|XP_017134715.1| PREDICTED: uncharacterized protein LOC108150906 isoform X1
>gi|1036704173|ref|XP_017134732.1| PREDICTED: uncharacterized protein LOC108150906 isoform X3
>gi|1036698781|ref|XP_017135401.1| PREDICTED: maternal protein exuperantia-2-like [Drosophila r
>gi|1036751808|ref|XP_017156397.1| PREDICTED: cytochrome P450 4g15 isoform X1 [Drosophila mira
>gi|1036751825|ref|XP_017156398.1| PREDICTED: cytochrome P450 4g15 isoform X2 [Drosophila mira
>gi|1036750023|ref|XP_017156299.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750042|ref|XP_017156300.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750062|ref|XP_017156301.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750006|ref|XP_017156298.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750083|ref|XP_017156302.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750102|ref|XP_017156303.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036720713|ref|XP_017144781.1| PREDICTED: neurofilament heavy polypeptide [Drosophila miran
>gi|1036720694|ref|XP_017144769.1| PREDICTED: protein flightless-1 [Drosophila miranda]
>gi|1036747985|ref|XP_017156197.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748002|ref|XP_017156198.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748021|ref|XP_017156199.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748038|ref|XP_017156200.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748053|ref|XP_017156201.1| PREDICTED: protein tweety isoform X2 [Drosophila miranda]
>gi|1036748070|ref|XP_017156202.1| PREDICTED: protein tweety isoform X3 [Drosophila miranda]
>gi|1036742160|ref|XP_017155868.1| PREDICTED: pyridoxal-dependent decarboxylase domain-containi
>gi|1036757908|ref|XP_017156749.1| PREDICTED: OTU domain-containing protein 1-like [Drosophila
>gi|1036757927|ref|XP_017156750.1| PREDICTED: OTU domain-containing protein 1-like [Drosophila
>gi|1036726043|ref|XP_017147530.1| PREDICTED: acetylcholine receptor subunit alpha-like [Drosop
>gi|1036726076|ref|XP_017147552.1| PREDICTED: uncharacterized protein LOC108159091 isoform X2
>gi|1036726063|ref|XP_017147543.1| PREDICTED: uncharacterized protein LOC108159091 isoform X1
>gi|1036726093|ref|XP_017147565.1| PREDICTED: pyridoxine/pyridoxamine 5'-phosphate oxidase [Dro
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>gi|1036726024|ref|XP_017147517.1| PREDICTED: uncharacterized protein LOC108159070 isoform X2
>gi|1036726005|ref|XP_017147510.1| PREDICTED: uncharacterized protein LOC108159070 isoform X1
>gi|1036726112|ref|XP_017147578.1| PREDICTED: uncharacterized protein LOC108159109 [Drosophila
>gi|1036747817|ref|XP_017156191.1| PREDICTED: protein RFT1 homolog [Drosophila miranda]
>gi|1036726492|ref|XP_017147819.1| PREDICTED: uncharacterized protein LOC108159225 [Drosophila
>gi|1036726505|ref|XP_017147832.1| PREDICTED: UDP-N-acetylglucosamine transporter [Drosophila m
>gi|1036726524|ref|XP_017147846.1| PREDICTED: translation initiation factor eIF-2B subunit beta
>gi|1036726541|ref|XP_017147858.1| PREDICTED: phosphoglycolate phosphatase 2 [Drosophila mirano
>gi|1036726439|ref|XP_017147786.1| PREDICTED: uncharacterized protein LOC108159214 isoform X1
>gi|1036726458|ref|XP_017147797.1| PREDICTED: uncharacterized protein LOC108159214 isoform X2
>gi|1036726476|ref|XP_017147807.1| PREDICTED: uncharacterized protein LOC108159214 isoform X3
>gi|1036745594|ref|XP_017156057.1| PREDICTED: WW domain-binding protein 11 [Drosophila miranda]
>gi|1036732702|ref|XP_017151275.1| PREDICTED: activating signal cointegrator 1 complex subunit
>gi|1036732723|ref|XP_017151287.1| PREDICTED: mitosis initiation protein fs(1)Ya [Drosophila mi
>gi|1036732742|ref|XP_017151298.1| PREDICTED: RING finger protein vilya isoform X1 [Drosophila
>gi|1036732761|ref|XP_017151306.1| PREDICTED: RING finger protein vilya isoform X2 [Drosophila
>gi|1036718943|ref|XP_017143840.1| PREDICTED: zinc finger protein 391 [Drosophila miranda]
>gi|1036718964|ref|XP_017143851.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036698798|ref|XP_017135524.1| PREDICTED: zinc finger protein 569 [Drosophila miranda]
>gi|1036718923|ref|XP_017143828.1| PREDICTED: protein cramped [Drosophila miranda]
>gi|1036718983|ref|XP_017143862.1| PREDICTED: syntaxin-4 isoform X1 [Drosophila miranda]
>gi|1036719003|ref|XP_017143871.1| PREDICTED: syntaxin-4 isoform X2 [Drosophila miranda]
>gi|1036705326|ref|XP_017135528.1| PREDICTED: uncharacterized protein LOC108151430 isoform X2
>gi|1036705311|ref|XP_017135519.1| PREDICTED: uncharacterized protein LOC108151430 isoform X1
>gi|1036752884|ref|XP_017156465.1| PREDICTED: uncharacterized protein LOC108164964 [Drosophila
>gi|1036752901|ref|XP_017156466.1| PREDICTED: uncharacterized protein LOC108164964 [Drosophila
>gi|1036698813|ref|XP_017135648.1| PREDICTED: POC1 centriolar protein homolog A-like [Drosophil
>gi|1036752919|ref|XP_017156467.1| PREDICTED: ARL-6-interacting protein 1 homolog [Drosophila m
>gi|1036703713|ref|XP_017157108.1| PREDICTED: uncharacterized protein LOC108165541 [Drosophila
>gi|1036707146|ref|XP_017136781.1| PREDICTED: uncharacterized protein LOC108152167 [Drosophila
>gi|1036698828|ref|XP_017135766.1| PREDICTED: uncharacterized protein LOC108151568 [Drosophila
>gi|1036757172|ref|XP_017156709.1| PREDICTED: TAF5-like RNA polymerase II p300/CBP-associated f
>gi|1036705033|ref|XP_017135326.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1-like [Drosophil
>gi|1036735617|ref|XP_017152866.1| PREDICTED: transmembrane protein 120 homolog isoform X1 [Dro
>gi|1036735635|ref|XP_017152876.1| PREDICTED: transmembrane protein 120 homolog isoform X2 [Dro
>gi|1036735598|ref|XP_017152854.1| PREDICTED: protein white isoform X2 [Drosophila miranda]
>gi|1036735579|ref|XP_017152845.1| PREDICTED: protein white isoform X1 [Drosophila miranda]
>gi|1036735650|ref|XP_017152888.1| PREDICTED: protein angel-like [Drosophila miranda]
>gi|1036705224|ref|XP_017135465.1| PREDICTED: RNA polymerase-associated protein Rtf1 [Drosophil
>gi|1036704350|ref|XP_017134836.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036705156|ref|XP_017135417.1| PREDICTED: protein germ cell-less-like [Drosophila miranda]
>gi|1036705135|ref|XP_017135405.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036698842|ref|XP_017135892.1| PREDICTED: anaphase-promoting complex subunit cdh1 [Drosophi
>gi|1036698859|ref|XP_017136018.1| PREDICTED: uncharacterized protein LOC108151713 [Drosophila
>gi|1036698874|ref|XP_017136145.1| PREDICTED: putative cyclin-dependent serine/threonine-protei
>gi|1036706528|ref|XP_017136355.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036756826|ref|XP_017156692.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 [Drosophila mir
>gi|1036755996|ref|XP_017156644.1| PREDICTED: glycine, alanine and asparagine-rich protein [Dro
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>gi|1036708233|ref|XP_017137508.1| PREDICTED: uncharacterized protein LOC108152577 [Drosophila
>gi|1036698889|ref|XP_017136270.1| PREDICTED: eukaryotic translation initiation factor 4 gamma
>gi|1036706634|ref|XP_017136437.1| PREDICTED: uncharacterized protein LOC108151980 [Drosophila
>gi|1036698904|ref|XP_017136392.1| PREDICTED: PH domain-containing protein DDB_G0287875 [Drosop
>gi|1036698917|ref|XP_017136516.1| PREDICTED: putative uncharacterized protein DDB_G0274435 [Dr
>gi|1036698931|ref|XP_017136644.1| PREDICTED: protein PRY2 [Drosophila miranda]
>gi|1036698948|ref|XP_017136777.1| PREDICTED: uncharacterized protein LOC108152142 [Drosophila
>gi|1036698961|ref|XP_017136902.1| PREDICTED: uncharacterized protein LOC108152210 [Drosophila
>gi|1036754447|ref|XP_017156558.1| PREDICTED: estradiol 17-beta-dehydrogenase 8 [Drosophila mi
>gi|1036739106|ref|XP_017154782.1| PREDICTED: irregular chiasm C-roughest protein [Drosophila m
>gi|1036703531|ref|XP_017157001.1| PREDICTED: uncharacterized protein LOC108165462 [Drosophila
>gi|1036751645|ref|XP_017156389.1| PREDICTED: uncharacterized protein LOC108164914 [Drosophila
>gi|1036698975|ref|XP_017137021.1| PREDICTED: jacalin-related lectin 34 [Drosophila miranda]
>gi|1036755261|ref|XP_017156606.1| PREDICTED: hepatoma-derived growth factor-related protein 2-
>gi|1036755277|ref|XP_017156607.1| PREDICTED: uncharacterized protein LOC108165069 isoform X2
>gi|1036715997|ref|XP_017142187.1| PREDICTED: irregular chiasm C-roughest protein isoform X1 [
>gi|1036716017|ref|XP_017142194.1| PREDICTED: irregular chiasm C-roughest protein isoform X2 [
>gi|1036716037|ref|XP_017142203.1| PREDICTED: irregular chiasm C-roughest protein isoform X2 [
>gi|1036715979|ref|XP_017142172.1| PREDICTED: neurogenic locus Notch protein [Drosophila miran
>gi|1036752611|ref|XP_017156446.1| PREDICTED: UPF0729 protein GD16342 [Drosophila miranda]
>gi|1036742016|ref|XP_017155862.1| PREDICTED: follicle cell protein 3C-1 [Drosophila miranda]
>gi|1036742037|ref|XP_017155863.1| PREDICTED: uncharacterized protein LOC108164557 [Drosophila
>gi|1036742056|ref|XP_017155864.1| PREDICTED: thioredoxin domain-containing protein 17 [Drosop
>gi|1036719280|ref|XP_017144018.1| PREDICTED: uncharacterized protein LOC108156826 [Drosophila
>gi|1036719338|ref|XP_017144056.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036719319|ref|XP_017144043.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036719360|ref|XP_017144068.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036719299|ref|XP_017144031.1| PREDICTED: alpha-tubulin N-acetyltransferase 1-like [Drosop
>gi|1036719037|ref|XP_017143911.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719054|ref|XP_017143919.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719074|ref|XP_017143927.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719095|ref|XP_017143936.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719120|ref|XP_017143944.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719139|ref|XP_017143952.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719159|ref|XP_017143961.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719179|ref|XP_017143969.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719197|ref|XP_017143976.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase, is
>gi|1036719262|ref|XP_017144005.1| PREDICTED: katanin p60 ATPase-containing subunit A-like 2 [
>gi|1036719219|ref|XP_017143984.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719244|ref|XP_017143993.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036739750|ref|XP_017155172.1| PREDICTED: septin-5 isoform X1 [Drosophila miranda]
>gi|1036739769|ref|XP_017155178.1| PREDICTED: septin-5 isoform X2 [Drosophila miranda]
>gi|1036698998|ref|XP_017137147.1| PREDICTED: transmembrane protease serine 9-like [Drosophila
>gi|1036746332|ref|XP_017156100.1| PREDICTED: trypsin beta [Drosophila miranda]
>gi|1036746368|ref|XP_017156102.1| PREDICTED: serine protease SP24D [Drosophila miranda]
>gi|1036746351|ref|XP_017156101.1| PREDICTED: serine protease SP24D [Drosophila miranda]
>gi|1036746225|ref|XP_017156094.1| PREDICTED: carboxypeptidase D isoform X1 [Drosophila miranda
>gi|1036746244|ref|XP_017156095.1| PREDICTED: carboxypeptidase D isoform X1 [Drosophila miranda
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>gi|1036746263|ref|XP_017156096.1| PREDICTED: carboxypeptidase D isoform X2 [Drosophila miranda
>gi|1036746282|ref|XP_017156097.1| PREDICTED: carboxypeptidase D isoform X3 [Drosophila miranda
>gi|1036746297|ref|XP_017156098.1| PREDICTED: carboxypeptidase D isoform X4 [Drosophila miranda
>gi|1036746314|ref|XP_017156099.1| PREDICTED: carboxypeptidase D isoform X5 [Drosophila miranda
>gi|1036746209|ref|XP_017156093.1| PREDICTED: uncharacterized protein LOC108164719 [Drosophila
>gi|1036728992|ref|XP_017149297.1| PREDICTED: transforming growth factor beta-1-induced transcr
>gi|1036729007|ref|XP_017149305.1| PREDICTED: transforming growth factor beta-1-induced transcr
>gi|1036728909|ref|XP_017149239.1| PREDICTED: uncharacterized protein LOC108160007 isoform X2
>gi|1036728889|ref|XP_017149230.1| PREDICTED: uncharacterized protein LOC108160007 isoform X1
>gi|1036728873|ref|XP_017149217.1| PREDICTED: endothelin-converting enzyme 1-like isoform X2 [
>gi|1036728855|ref|XP_017149208.1| PREDICTED: endothelin-converting enzyme 1-like isoform X1 [
>gi|1036728940|ref|XP_017149263.1| PREDICTED: uncharacterized protein LOC108160031 [Drosophila
>gi|1036728959|ref|XP_017149274.1| PREDICTED: uncharacterized protein LOC108160038 [Drosophila
>gi|1036728976|ref|XP_017149286.1| PREDICTED: uncharacterized protein LOC108160049 [Drosophila
>gi|1036728925|ref|XP_017149251.1| PREDICTED: keratin, type I cytoskeletal 10-like [Drosophila
>gi|1036704855|ref|XP_017135194.1| PREDICTED: casein kinase II subunit beta'-like [Drosophila
>gi|1036753532|ref|XP_017156504.1| PREDICTED: TWiK family of potassium channels protein 7 [Dros
>gi|1036753553|ref|XP_017156505.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036752754|ref|XP_017156455.1| PREDICTED: transmembrane emp24 domain-containing protein 2
>gi|1036743155|ref|XP_017155927.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036743136|ref|XP_017155926.1| PREDICTED: uncharacterized protein LOC108164598 [Drosophila
>gi|1036730851|ref|XP_017150432.1| PREDICTED: uncharacterized protein LOC108160747 [Drosophila
>gi|1036730833|ref|XP_017150421.1| PREDICTED: structural maintenance of chromosomes protein 3
>gi|1036727974|ref|XP_017148695.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein pho
>gi|1036736569|ref|XP_017153421.1| PREDICTED: actin-related protein 2 isoform X1 [Drosophila mi
>gi|1036736587|ref|XP_017153430.1| PREDICTED: actin-related protein 2 isoform X1 [Drosophila mi
>gi|1036736607|ref|XP_017153438.1| PREDICTED: actin-related protein 2 isoform X2 [Drosophila mi
>gi|1036756469|ref|XP_017156672.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila
>gi|1036756487|ref|XP_017156673.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila
>gi|1036756505|ref|XP_017156674.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila
>gi|1036738091|ref|XP_017154268.1| PREDICTED: sodium channel protein para isoform X1 [Drosophil
>gi|1036738109|ref|XP_017154276.1| PREDICTED: sodium channel protein para isoform X2 [Drosophil
>gi|1036738359|ref|XP_017154388.1| PREDICTED: sodium channel protein para isoform X15 [Drosoph
>gi|1036738197|ref|XP_017154320.1| PREDICTED: sodium channel protein para isoform X7 [Drosophil
>gi|1036738461|ref|XP_017154434.1| PREDICTED: sodium channel protein para isoform X21 [Drosoph
>gi|1036738127|ref|XP_017154285.1| PREDICTED: sodium channel protein para isoform X3 [Drosophil
>gi|1036738145|ref|XP_017154294.1| PREDICTED: sodium channel protein para isoform X4 [Drosophil
>gi|1036738429|ref|XP_017154417.1| PREDICTED: sodium channel protein para isoform X19 [Drosoph
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>gi|1036738279|ref|XP_017154364.1| PREDICTED: sodium channel protein para isoform X12 [Drosoph
>gi|1036738325|ref|XP_017154373.1| PREDICTED: sodium channel protein para isoform X13 [Drosoph
>gi|1036738507|ref|XP_017154459.1| PREDICTED: sodium channel protein para isoform X24 [Drosoph
>gi|1036738375|ref|XP_017154396.1| PREDICTED: sodium channel protein para isoform X16 [Drosoph
>gi|1036738559|ref|XP_017154484.1| PREDICTED: sodium channel protein para isoform X27 [Drosoph
>gi|1036738343|ref|XP_017154380.1| PREDICTED: sodium channel protein para isoform X14 [Drosoph
>gi|1036738163|ref|XP_017154304.1| PREDICTED: sodium channel protein para isoform X5 [Drosophil
>gi|1036738230|ref|XP_017154337.1| PREDICTED: sodium channel protein para isoform X9 [Drosophil
>gi|1036738212|ref|XP_017154329.1| PREDICTED: sodium channel protein para isoform X8 [Drosophil
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>gi|1036738475|ref|XP_017154442.1| PREDICTED: sodium channel protein para isoform X22 [Drosophi
>gi|1036738525|ref|XP_017154468.1| PREDICTED: sodium channel protein para isoform X25 [Drosophi
>gi|1036738247|ref|XP_017154348.1| PREDICTED: sodium channel protein para isoform X10 [Drosophi
>gi|1036738491|ref|XP_017154451.1| PREDICTED: sodium channel protein para isoform X23 [Drosophi
>gi|1036738411|ref|XP_017154411.1| PREDICTED: sodium channel protein para isoform X18 [Drosophi
>gi|1036738393|ref|XP_017154404.1| PREDICTED: sodium channel protein para isoform X17 [Drosophi
>gi|1036738592|ref|XP_017154499.1| PREDICTED: sodium channel protein para isoform X29 [Drosophi
>gi|1036738576|ref|XP_017154491.1| PREDICTED: sodium channel protein para isoform X28 [Drosophi
>gi|1036738610|ref|XP_017154508.1| PREDICTED: sodium channel protein para isoform X30 [Drosophi
>gi|1036738179|ref|XP_017154312.1| PREDICTED: sodium channel protein para isoform X6 [Drosophil
>gi|1036738646|ref|XP_017154525.1| PREDICTED: sodium channel protein para isoform X32 [Drosophi
>gi|1036738628|ref|XP_017154515.1| PREDICTED: sodium channel protein para isoform X31 [Drosophi
>gi|1036738543|ref|XP_017154476.1| PREDICTED: sodium channel protein para isoform X26 [Drosophi
>gi|1036738447|ref|XP_017154426.1| PREDICTED: sodium channel protein para isoform X20 [Drosophi
>gi|1036738666|ref|XP_017154530.1| PREDICTED: sodium channel protein para isoform X33 [Drosophi
>gi|1036706974|ref|XP_017136670.1| PREDICTED: eukaryotic translation initiation factor 4H isofo
>gi|1036706934|ref|XP_017136649.1| PREDICTED: eukaryotic translation initiation factor 4H isofo
>gi|1036706955|ref|XP_017136659.1| PREDICTED: eukaryotic translation initiation factor 4H isofo
>gi|1036707024|ref|XP_017136699.1| PREDICTED: eukaryotic translation initiation factor 2 subuni
>gi|1036706875|ref|XP_017136615.1| PREDICTED: protein DDI1 homolog 2 [Drosophila miranda]
>gi|1036706991|ref|XP_017136682.1| PREDICTED: cell cycle control protein 50A [Drosophila mirand
>gi|1036706825|ref|XP_017136586.1| PREDICTED: zinc finger protein hangover isoform X2 [Drosophi
>gi|1036706808|ref|XP_017136578.1| PREDICTED: zinc finger protein hangover isoform X1 [Drosophi
>gi|1036706859|ref|XP_017136604.1| PREDICTED: zinc finger protein hangover isoform X4 [Drosophi
>gi|1036706842|ref|XP_017136596.1| PREDICTED: zinc finger protein hangover isoform X3 [Drosophi
>gi|1036706894|ref|XP_017136629.1| PREDICTED: regucalcin isoform X1 [Drosophila miranda]
>gi|1036706911|ref|XP_017136637.1| PREDICTED: regucalcin isoform X2 [Drosophila miranda]
>gi|1036707075|ref|XP_017136735.1| PREDICTED: chromatin accessibility complex protein 1 [Droso
>gi|1036707041|ref|XP_017136712.1| PREDICTED: protein LSM12 homolog A isoform X1 [Drosophila m
>gi|1036707058|ref|XP_017136721.1| PREDICTED: protein LSM12 homolog A isoform X2 [Drosophila m
>gi|1036710860|ref|XP_017139156.1| PREDICTED: pleiotropic regulator 1 [Drosophila miranda]
>gi|1036710844|ref|XP_017139146.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 7 [Drosophi
>gi|1036710877|ref|XP_017139168.1| PREDICTED: uncharacterized protein LOC108153594 [Drosophila
>gi|1036710928|ref|XP_017139206.1| PREDICTED: rhythmically expressed gene 2 protein [Drosophila
>gi|1036710894|ref|XP_017139181.1| PREDICTED: 26S protease regulatory subunit 10B [Drosophila
>gi|1036710911|ref|XP_017139193.1| PREDICTED: tRNA (guanine-N(7)-)-methyltransferase non-cataly
>gi|1036710232|ref|XP_017138730.1| PREDICTED: kinesin-like protein Klp10A [Drosophila miranda]
>gi|1036710253|ref|XP_017138737.1| PREDICTED: kinesin-like protein Klp10A [Drosophila miranda]
>gi|1036710287|ref|XP_017138765.1| PREDICTED: uncharacterized protein LOC108153337 isoform X1
>gi|1036710309|ref|XP_017138776.1| PREDICTED: homeotic protein ocelliless isoform X2 [Drosophil
>gi|1036710412|ref|XP_017138847.1| PREDICTED: venom toxin OcyC11 [Drosophila miranda]
>gi|1036710393|ref|XP_017138836.1| PREDICTED: uncharacterized protein LOC108153373 [Drosophila
>gi|1036710376|ref|XP_017138824.1| PREDICTED: uncharacterized protein LOC108153357 isoform X2
>gi|1036710359|ref|XP_017138816.1| PREDICTED: uncharacterized protein LOC108153357 isoform X1
>gi|1036710342|ref|XP_017138805.1| PREDICTED: uncharacterized protein LOC108153349 [Drosophila
>gi|1036710326|ref|XP_017138791.1| PREDICTED: GTP-binding nuclear protein Ran [Drosophila mira
>gi|1036710270|ref|XP_017138751.1| PREDICTED: probable G-protein coupled receptor Mth-like 1 [
>gi|1036741020|ref|XP_017155802.1| PREDICTED: DNA topoisomerase 3-beta [Drosophila miranda]
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>gi|1036719809|ref|XP_017144302.1| PREDICTED: ubiquitin [Drosophila miranda]
>gi|1036719789|ref|XP_017144290.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036719769|ref|XP_017144277.1| PREDICTED: zinc finger SWIM domain-containing protein 8 [Dro
>gi|1036699011|ref|XP_017137275.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miranc
>gi|1036723238|ref|XP_017145878.1| PREDICTED: CAD protein [Drosophila miranda]
>gi|1036723248|ref|XP_017145886.1| PREDICTED: uncharacterized protein LOC108158176 [Drosophila
>gi|1036699026|ref|XP_017137403.1| PREDICTED: LOW QUALITY PROTEIN: protein BUD31 homolog [Drosc
>gi|1036723373|ref|XP_017145935.1| PREDICTED: uncharacterized protein LOC108158217 [Drosophila
>gi|1036723354|ref|XP_017145922.1| PREDICTED: protein sisterless A [Drosophila miranda]
>gi|1036699039|ref|XP_017137528.1| PREDICTED: uncharacterized protein LOC108152566 [Drosophila
>gi|1036723390|ref|XP_017145948.1| PREDICTED: uncharacterized protein LOC108158224 [Drosophila
>gi|1036723337|ref|XP_017145911.1| PREDICTED: protein naked cuticle [Drosophila miranda]
>gi|1036707179|ref|XP_017136808.1| PREDICTED: uncharacterized protein LOC108152181 isoform X1
>gi|1036707200|ref|XP_017136817.1| PREDICTED: uncharacterized protein LOC108152181 isoform X2
>gi|1036704367|ref|XP_017134846.1| PREDICTED: uncharacterized protein LOC108151003 [Drosophila
>gi|1036704030|ref|XP_017134646.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036725562|ref|XP_017147235.1| PREDICTED: cytoplasmic dynein 1 light intermediate chain 1
>gi|1036725581|ref|XP_017147245.1| PREDICTED: cytoplasmic dynein 1 light intermediate chain 1
>gi|1036725683|ref|XP_017147298.1| PREDICTED: armadillo repeat-containing protein 2 [Drosophila
>gi|1036725661|ref|XP_017147286.1| PREDICTED: uncharacterized protein LOC108158962 [Drosophila
>gi|1036725600|ref|XP_017147257.1| PREDICTED: uncharacterized protein LOC108158945 isoform X1
>gi|1036725619|ref|XP_017147266.1| PREDICTED: uncharacterized protein LOC108158945 isoform X2
>gi|1036725638|ref|XP_017147273.1| PREDICTED: uncharacterized protein LOC108158945 isoform X3
>gi|1036725545|ref|XP_017147223.1| PREDICTED: uncharacterized protein LOC108158926 [Drosophila
>gi|1036735206|ref|XP_017152647.1| PREDICTED: bleomycin hydrolase [Drosophila miranda]
>gi|1036735225|ref|XP_017152659.1| PREDICTED: uncharacterized protein F58A4.6 [Drosophila miran
>gi|1036735244|ref|XP_017152667.1| PREDICTED: uncharacterized protein F58A4.6 [Drosophila miran
>gi|1036705294|ref|XP_017135507.1| PREDICTED: uncharacterized protein LOC108151424 [Drosophila
>gi|1036754095|ref|XP_017156537.1| PREDICTED: T-box transcription factor TBX1 [Drosophila miran
>gi|1036704783|ref|XP_017135141.1| PREDICTED: uncharacterized protein LOC108151194 [Drosophila
>gi|1036754656|ref|XP_017156570.1| PREDICTED: uncharacterized protein LOC108165043 [Drosophila
>gi|1036703385|ref|XP_017156917.1| PREDICTED: very low-density lipoprotein receptor-like [Drosc
>gi|1036703400|ref|XP_017156925.1| PREDICTED: very low-density lipoprotein receptor-like [Drosc
>gi|1036744849|ref|XP_017156011.1| PREDICTED: protein DGCR14 homolog [Drosophila miranda]
>gi|1036744868|ref|XP_017156012.1| PREDICTED: sepiapterin reductase [Drosophila miranda]
>gi|1036744784|ref|XP_017156009.1| PREDICTED: sepiapterin reductase [Drosophila miranda]
>gi|1036744748|ref|XP_017156007.1| PREDICTED: uncharacterized protein LOC108164665 isoform X1
>gi|1036744766|ref|XP_017156008.1| PREDICTED: uncharacterized protein LOC108164665 isoform X2
>gi|1036744731|ref|XP_017156006.1| PREDICTED: uncharacterized protein LOC108164663 isoform X2
>gi|1036744713|ref|XP_017156005.1| PREDICTED: uncharacterized protein LOC108164663 isoform X1
>gi|1036704836|ref|XP_017135182.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036704247|ref|XP_017134774.1| PREDICTED: chorion protein S36 [Drosophila miranda]
>gi|1036757301|ref|XP_017156717.1| PREDICTED: chorion protein S38 [Drosophila miranda]
>gi|1036726169|ref|XP_017147613.1| PREDICTED: protein ovarian tumor locus isoform X2 [Drosophil
>gi|1036726150|ref|XP_017147604.1| PREDICTED: protein ovarian tumor locus isoform X1 [Drosophil
>gi|1036726189|ref|XP_017147621.1| PREDICTED: protein ovarian tumor locus isoform X1 [Drosophil
>gi|1036726280|ref|XP_017147668.1| PREDICTED: solute carrier family 41 member 1 isoform X5 [Dro
>gi|1036726257|ref|XP_017147658.1| PREDICTED: solute carrier family 41 member 1 isoform X4 [Dro
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>gi|1036726241|ref|XP_017147648.1| PREDICTED: solute carrier family 41 member 2 isoform X3 [Dro
>gi|1036726204|ref|XP_017147632.1| PREDICTED: solute carrier family 41 member 1 isoform X1 [Dro
>gi|1036726222|ref|XP_017147640.1| PREDICTED: solute carrier family 41 member 1 isoform X2 [Dro
>gi|1036734410|ref|XP_017152186.1| PREDICTED: kinetochore protein NDC80 homolog [Drosophila mi
>gi|1036734394|ref|XP_017152174.1| PREDICTED: late secretory pathway protein AVL9 homolog [Dros
>gi|1036750613|ref|XP_017156329.1| PREDICTED: troponin T, skeletal muscle isoform X2 [Drosophil
>gi|1036750594|ref|XP_017156328.1| PREDICTED: troponin T, skeletal muscle isoform X1 [Drosophil
>gi|1036750665|ref|XP_017156332.1| PREDICTED: troponin T, skeletal muscle isoform X5 [Drosophil
>gi|1036750629|ref|XP_017156330.1| PREDICTED: troponin T, skeletal muscle isoform X3 [Drosophil
>gi|1036750832|ref|XP_017156343.1| PREDICTED: troponin T, skeletal muscle isoform X14 [Drosoph
>gi|1036750648|ref|XP_017156331.1| PREDICTED: troponin T, skeletal muscle isoform X4 [Drosophil
>gi|1036750812|ref|XP_017156342.1| PREDICTED: troponin T, skeletal muscle isoform X13 [Drosoph
>gi|1036750760|ref|XP_017156339.1| PREDICTED: troponin T, skeletal muscle isoform X10 [Drosoph
>gi|1036750741|ref|XP_017156338.1| PREDICTED: troponin T, skeletal muscle isoform X9 [Drosophi
>gi|1036750795|ref|XP_017156341.1| PREDICTED: troponin T, skeletal muscle isoform X12 [Drosoph
>gi|1036750777|ref|XP_017156340.1| PREDICTED: troponin T, skeletal muscle isoform X11 [Drosoph
>gi|1036750684|ref|XP_017156333.1| PREDICTED: troponin T, skeletal muscle isoform X6 [Drosophil
>gi|1036750703|ref|XP_017156334.1| PREDICTED: troponin T, skeletal muscle isoform X7 [Drosophil
>gi|1036750722|ref|XP_017156335.1| PREDICTED: troponin T, skeletal muscle isoform X8 [Drosophil
>gi|1036727090|ref|XP_017148180.1| PREDICTED: protein-tyrosine sulfotransferase isoform X1 [Dro
>gi|1036727109|ref|XP_017148188.1| PREDICTED: protein-tyrosine sulfotransferase isoform X1 [Dro
>gi|1036727126|ref|XP_017148197.1| PREDICTED: protein-tyrosine sulfotransferase isoform X2 [Dro
>gi|1036703433|ref|XP_017156947.1| PREDICTED: sperm flagellar protein 1-like [Drosophila mirand
>gi|1036703450|ref|XP_017156955.1| PREDICTED: sperm flagellar protein 1-like [Drosophila mirand
>gi|1036703463|ref|XP_017156961.1| PREDICTED: sperm flagellar protein 1-like [Drosophila mirand
>gi|1036707302|ref|XP_017136877.1| PREDICTED: uncharacterized protein LOC108152215 [Drosophila
>gi|1036706709|ref|XP_017136517.1| PREDICTED: uncharacterized protein LOC108152033 [Drosophila
>gi|1036705173|ref|XP_017135429.1| PREDICTED: uncharacterized protein LOC108151374 [Drosophila
>gi|1036705190|ref|XP_017135436.1| PREDICTED: uncharacterized protein LOC108151374 [Drosophila
>gi|1036706404|ref|XP_017136278.1| PREDICTED: larval cuticle protein LCP-14 [Drosophila miranda
>gi|1036705689|ref|XP_017135760.1| PREDICTED: uncharacterized protein LOC108151588 [Drosophila
>gi|1036707234|ref|XP_017136842.1| PREDICTED: uncharacterized protein LOC108152195 [Drosophila
>gi|1036737335|ref|XP_017153863.1| PREDICTED: uncharacterized protein LOC108163215 [Drosophila
>gi|1036737350|ref|XP_017153874.1| PREDICTED: mucin-2-like [Drosophila miranda]
>gi|1036757320|ref|XP_017156718.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1036757339|ref|XP_017156719.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1036757354|ref|XP_017156720.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1036752846|ref|XP_017156461.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1036752865|ref|XP_017156462.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1036699054|ref|XP_017137653.1| PREDICTED: proteasome subunit alpha type-2-like, partial [D
>gi|1036725774|ref|XP_017147365.1| PREDICTED: protein LST8 homolog [Drosophila miranda]
>gi|1036725755|ref|XP_017147351.1| PREDICTED: ADP-ribosylation factor-binding protein GGA2 [Dro
>gi|1036725793|ref|XP_017147377.1| PREDICTED: 40S ribosomal protein S5a [Drosophila miranda]
>gi|1036699069|ref|XP_017137778.1| PREDICTED: uncharacterized protein LOC108152715 [Drosophila
>gi|1036742141|ref|XP_017155867.1| PREDICTED: uncharacterized protein LOC108164561 isoform X2
>gi|1036742123|ref|XP_017155866.1| PREDICTED: uncharacterized protein LOC108164561 isoform X1
>gi|1036752972|ref|XP_017156470.1| PREDICTED: nuclear transcription factor Y subunit gamma-like
>gi|1036752955|ref|XP_017156469.1| PREDICTED: uncharacterized protein LOC108164968 [Drosophila
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>gi|1036731537|ref|XP_017150762.1| PREDICTED: uncharacterized protein LOC108160971 isoform X1
>gi|1036731556|ref|XP_017150770.1| PREDICTED: sterol 3-beta-glucosyltransferase isoform X2 [Dro
>gi|1036731573|ref|XP_017150778.1| PREDICTED: pleckstrin homology-like domain family B member
>gi|1036745632|ref|XP_017156059.1| PREDICTED: coatomer subunit delta [Drosophila miranda]
>gi|1036712871|ref|XP_017140293.1| PREDICTED: LOW QUALITY PROTEIN: acidic repeat-containing pro
>gi|1036712972|ref|XP_017140349.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036712888|ref|XP_017140304.1| PREDICTED: peroxisomal targeting signal 1 receptor [Drosoph
>gi|1036712853|ref|XP_017140283.1| PREDICTED: protein MMS22-like [Drosophila miranda]
>gi|1036712955|ref|XP_017140341.1| PREDICTED: serine/threonine-protein phosphatase Pgam5, mito
>gi|1036712938|ref|XP_017140333.1| PREDICTED: serine/threonine-protein phosphatase Pgam5, mito
>gi|1036712905|ref|XP_017140315.1| PREDICTED: calphotin isoform X1 [Drosophila miranda]
>gi|1036712922|ref|XP_017140323.1| PREDICTED: GATA type zinc finger protein asd-4 isoform X2 [
>gi|1036748739|ref|XP_017156228.1| PREDICTED: cytochrome P450 4d1 isoform X2 [Drosophila miran
>gi|1036748720|ref|XP_017156227.1| PREDICTED: cytochrome P450 4d1 isoform X1 [Drosophila miran
>gi|1036748758|ref|XP_017156229.1| PREDICTED: uncharacterized protein LOC108164819 [Drosophila
>gi|1036748815|ref|XP_017156232.1| PREDICTED: E3 ubiquitin-protein ligase RNF125-like [Drosoph
>gi|1036748777|ref|XP_017156230.1| PREDICTED: uncharacterized protein LOC108164820 isoform X1
>gi|1036748796|ref|XP_017156231.1| PREDICTED: uncharacterized protein LOC108164820 isoform X2
>gi|1036723794|ref|XP_017146152.1| PREDICTED: open rectifier potassium channel protein 1 isofor
>gi|1036723776|ref|XP_017146144.1| PREDICTED: open rectifier potassium channel protein 1 isofor
>gi|1036723813|ref|XP_017146163.1| PREDICTED: open rectifier potassium channel protein 1 isofor
>gi|1036723758|ref|XP_017146133.1| PREDICTED: putative ATP-dependent RNA helicase DHX57 [Drosop
>gi|1036723830|ref|XP_017146173.1| PREDICTED: uncharacterized protein LOC108158413 [Drosophila
>gi|1036705723|ref|XP_017135784.1| PREDICTED: uncharacterized protein LOC108151605 [Drosophila
>gi|1036706264|ref|XP_017136184.1| PREDICTED: E3 ubiquitin-protein ligase RNF125-like [Drosoph
>gi|1036707319|ref|XP_017136890.1| PREDICTED: cytochrome P450 4d1-like [Drosophila miranda]
>gi|1036745068|ref|XP_017156025.1| PREDICTED: uncharacterized protein LOC108164679 isoform X1
>gi|1036745087|ref|XP_017156026.1| PREDICTED: uncharacterized protein LOC108164679 isoform X1
>gi|1036745106|ref|XP_017156027.1| PREDICTED: uncharacterized protein LOC108164679 isoform X2
>gi|1036745125|ref|XP_017156028.1| PREDICTED: uncharacterized protein LOC108164679 isoform X3
>gi|1036745143|ref|XP_017156029.1| PREDICTED: uncharacterized protein LOC108164679 isoform X4
>gi|1036745809|ref|XP_017156068.1| PREDICTED: methylmalonic aciduria and homocystinuria type D
>gi|1036745827|ref|XP_017156069.1| PREDICTED: methylmalonic aciduria and homocystinuria type D
>gi|1036745846|ref|XP_017156070.1| PREDICTED: COMM domain-containing protein 3 [Drosophila mira
>gi|1036753185|ref|XP_017156483.1| PREDICTED: neuferricin homolog [Drosophila miranda]
>gi|1036726315|ref|XP_017147713.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726334|ref|XP_017147722.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726350|ref|XP_017147730.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726367|ref|XP_017147738.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726386|ref|XP_017147752.1| PREDICTED: nudC domain-containing protein 1 [Drosophila mira
>gi|1036726404|ref|XP_017147764.1| PREDICTED: transmembrane emp24 domain-containing protein 3
>gi|1036726422|ref|XP_017147774.1| PREDICTED: transmembrane emp24 domain-containing protein 3
>gi|1036699084|ref|XP_017137898.1| PREDICTED: transcription factor SPT20 homolog [Drosophila m
>gi|1036703584|ref|XP_017157032.1| PREDICTED: probable cytochrome P450 28c1 [Drosophila miranda
>gi|1036699098|ref|XP_017138026.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036709050|ref|XP_017138009.1| PREDICTED: breast carcinoma-amplified sequence 3 homolog iso
>gi|1036708999|ref|XP_017137984.1| PREDICTED: uncharacterized protein LOC108152864 isoform X1
>gi|1036709016|ref|XP_017137992.1| PREDICTED: uncharacterized protein LOC108152864 isoform X1
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>gi|1036709033|ref|XP_017138000.1| PREDICTED: uncharacterized protein LOC108152864 isoform X1
>gi|1036709065|ref|XP_017138021.1| PREDICTED: uncharacterized protein LOC108152890 [Drosophila
>gi|1036709079|ref|XP_017138031.1| PREDICTED: uncharacterized protein LOC108152890 [Drosophila
>gi|1036709100|ref|XP_017138042.1| PREDICTED: heat shock 70 kDa protein cognate 3 [Drosophila r
>gi|1036709119|ref|XP_017138051.1| PREDICTED: heat shock 70 kDa protein cognate 3 [Drosophila r
>gi|1036709136|ref|XP_017138061.1| PREDICTED: heat shock 70 kDa protein cognate 3 [Drosophila r
>gi|1036718613|ref|XP_017143633.1| PREDICTED: casein kinase II subunit beta isoform X1 [Drosopl
>gi|1036718631|ref|XP_017143639.1| PREDICTED: casein kinase II subunit beta isoform X1 [Drosopl
>gi|1036718647|ref|XP_017143648.1| PREDICTED: casein kinase II subunit beta isoform X2 [Drosopl
>gi|1036718576|ref|XP_017143609.1| PREDICTED: testis-expressed sequence 10 protein homolog [Dro
>gi|1036718594|ref|XP_017143621.1| PREDICTED: WD repeat-containing protein 46 [Drosophila mirar
>gi|1036755664|ref|XP_017156625.1| PREDICTED: uncharacterized protein LOC108165085 [Drosophila
>gi|1036710446|ref|XP_017138874.1| PREDICTED: transcriptional repressor CTCF [Drosophila mirand
>gi|1036710429|ref|XP_017138860.1| PREDICTED: dynein regulatory complex protein 1 homolog [Dros
>gi|1036710523|ref|XP_017138934.1| PREDICTED: zinc finger protein 311 isoform X1 [Drosophila mi
>gi|1036710536|ref|XP_017138943.1| PREDICTED: zinc finger protein 311 isoform X2 [Drosophila mi
>gi|1036710480|ref|XP_017138900.1| PREDICTED: mothers against decapentaplegic homolog 3 [Drosop
>gi|1036710553|ref|XP_017138955.1| PREDICTED: uncharacterized protein LOC108153446 [Drosophila
>gi|1036710463|ref|XP_017138887.1| PREDICTED: phenylalanine--tRNA ligase alpha subunit [Drosopl
>gi|1036710496|ref|XP_017138912.1| PREDICTED: zinc finger protein 311 [Drosophila miranda]
>gi|1036710570|ref|XP_017138968.1| PREDICTED: zinc finger protein 32 [Drosophila miranda]
>gi|1036710510|ref|XP_017138923.1| PREDICTED: zinc finger protein 813 [Drosophila miranda]
>gi|1036754365|ref|XP_017156553.1| PREDICTED: guanine nucleotide-binding protein subunit beta-5
>gi|1036751364|ref|XP_017156373.1| PREDICTED: zinc finger protein 32 [Drosophila miranda]
>gi|1036745562|ref|XP_017156055.1| PREDICTED: hepatic triacylglycerol lipase-like [Drosophila r
>gi|1036745544|ref|XP_017156054.1| PREDICTED: TNF receptor-associated factor 6 [Drosophila mira
>gi|1036748332|ref|XP_017156215.1| PREDICTED: acidic phospholipase A2 PA4 [Drosophila miranda]
>gi|1036748314|ref|XP_017156214.1| PREDICTED: uncharacterized protein LOC108164803 [Drosophila
>gi|1036699113|ref|XP_017138156.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 [Drosophila mi
>gi|1036699128|ref|XP_017138281.1| PREDICTED: LOW QUALITY PROTEIN: odorant receptor 7a [Drosopl
>gi|1036751974|ref|XP_017156407.1| PREDICTED: uncharacterized protein LOC108164930 [Drosophila
>gi|1036735669|ref|XP_017152900.1| PREDICTED: mitogen-activated protein kinase kinase kinase [I
>gi|1036727799|ref|XP_017148607.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036727834|ref|XP_017148623.1| PREDICTED: uncharacterized protein LOC108159648 isoform X3
>gi|1036727899|ref|XP_017148657.1| PREDICTED: uncharacterized protein LOC108159648 isoform X5
>gi|1036727848|ref|XP_017148632.1| PREDICTED: uncharacterized protein LOC108159648 isoform X4
>gi|1036727864|ref|XP_017148640.1| PREDICTED: uncharacterized protein LOC108159648 isoform X4
>gi|1036727880|ref|XP_017148648.1| PREDICTED: uncharacterized protein LOC108159648 isoform X4
>gi|1036727917|ref|XP_017148666.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X6 [Drosophi
>gi|1036727815|ref|XP_017148616.1| PREDICTED: uncharacterized protein LOC108159648 isoform X2
>gi|1036727936|ref|XP_017148676.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X7 [Drosophi
>gi|1036727955|ref|XP_017148684.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X7 [Drosophi
>gi|1036756326|ref|XP_017156663.1| PREDICTED: rRNA 2'-O-methyltransferase fibrillarin [Drosophi
>gi|1036699142|ref|XP_017138396.1| PREDICTED: glycine-rich protein DOT1 [Drosophila miranda]
>gi|1036736056|ref|XP_017153110.1| PREDICTED: DNA-binding protein Ewg isoform X1 [Drosophila mi
>gi|1036736097|ref|XP_017153121.1| PREDICTED: DNA-binding protein Ewg isoform X3 [Drosophila mi
>gi|1036736079|ref|XP_017153113.1| PREDICTED: DNA-binding protein Ewg isoform X2 [Drosophila mi
>gi|1036754185|ref|XP_017156543.1| PREDICTED: ras-related protein Rap-2b [Drosophila miranda]
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>gi|1036754202|ref|XP_017156544.1| PREDICTED: uncharacterized protein LOC108165017 [Drosophila
>gi|1036706676|ref|XP_017136483.1| PREDICTED: uncharacterized protein LOC108152014 [Drosophila
>gi|1036757081|ref|XP_017156704.1| PREDICTED: uncharacterized protein LOC108165148 [Drosophila
>gi|1036699157|ref|XP_017138516.1| PREDICTED: odorant receptor 1a [Drosophila miranda]
>gi|1036755921|ref|XP_017156640.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036755940|ref|XP_017156641.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036755959|ref|XP_017156642.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036755978|ref|XP_017156643.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036753933|ref|XP_017156528.1| PREDICTED: nuclear pore complex protein DDB_G0274915 [Drosop
>gi|1036699171|ref|XP_017138641.1| PREDICTED: uncharacterized protein LOC108153252 [Drosophila
>gi|1036699186|ref|XP_017138770.1| PREDICTED: uncharacterized protein LOC108153321 [Drosophila
>gi|1036757116|ref|XP_017156706.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1036702374|ref|XP_017156337.1| PREDICTED: achaete-scute complex protein T5 [Drosophila mira
>gi|1036702357|ref|XP_017156322.1| PREDICTED: achaete-scute complex protein T4 [Drosophila mira
>gi|1036702271|ref|XP_017156257.1| PREDICTED: achaete-scute complex protein T3 [Drosophila mira
>gi|1036757246|ref|XP_017156714.1| PREDICTED: pepsin-2B [Drosophila miranda]
>gi|1036754816|ref|XP_017156580.1| PREDICTED: achaete-scute complex protein T8 [Drosophila mira
>gi|1036727688|ref|XP_017148548.1| PREDICTED: cytochrome P450 4g1 [Drosophila miranda]
>gi|1036727670|ref|XP_017148537.1| PREDICTED: exportin-6-A [Drosophila miranda]
>gi|1036743242|ref|XP_017155933.1| PREDICTED: apoptosis regulatory protein Siva isoform X2 [Dro
>gi|1036743224|ref|XP_017155932.1| PREDICTED: apoptosis regulatory protein Siva isoform X1 [Dro
>gi|1036743277|ref|XP_017155934.1| PREDICTED: uncharacterized protein LOC108164606 [Drosophila
>gi|1036738775|ref|XP_017154589.1| PREDICTED: ATP-binding cassette sub-family B member 10, mito
>gi|1036738787|ref|XP_017154598.1| PREDICTED: probable methylmalonate-semialdehyde dehydrogenas
>gi|1036704384|ref|XP_017134859.1| PREDICTED: uncharacterized protein LOC108151009 [Drosophila
>gi|1036726558|ref|XP_017147870.1| PREDICTED: carboxypeptidase D isoform X1 [Drosophila miranda
>gi|1036726577|ref|XP_017147878.1| PREDICTED: carboxypeptidase D isoform X2 [Drosophila miranda
>gi|1036722294|ref|XP_017145462.1| PREDICTED: arginase-1 [Drosophila miranda]
>gi|1036722311|ref|XP_017145474.1| PREDICTED: protein elav [Drosophila miranda]
>gi|1036722277|ref|XP_017145450.1| PREDICTED: endoplasmic reticulum-Golgi intermediate compartm
>gi|1036740402|ref|XP_017155476.1| PREDICTED: chromodomain-helicase-DNA-binding protein 1 isofo
>gi|1036740420|ref|XP_017155480.1| PREDICTED: chromodomain-helicase-DNA-binding protein 1 isofo
>gi|1036740438|ref|XP_017155495.1| PREDICTED: NADH dehydrogenase [ubiquinone] flavoprotein 2, m
>gi|1036754256|ref|XP_017156547.1| PREDICTED: uncharacterized protein LOC108165021 [Drosophila
>gi|1036749413|ref|XP_017156264.1| PREDICTED: uncharacterized protein LOC108164846 [Drosophila
>gi|1036742717|ref|XP_017155901.1| PREDICTED: transmembrane protein 41 homolog [Drosophila mira
>gi|1036740314|ref|XP_017155458.1| PREDICTED: sn1-specific diacylglycerol lipase alpha isoform
>gi|1036740298|ref|XP_017155448.1| PREDICTED: sn1-specific diacylglycerol lipase alpha isoform
>gi|1036740332|ref|XP_017155469.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036732353|ref|XP_017151106.1| PREDICTED: regulator of nonsense transcripts 1 homolog [Dros
>gi|1036744658|ref|XP_017156002.1| PREDICTED: poly(A) RNA polymerase gld-2 homolog B [Drosophil
>gi|1036699201|ref|XP_017138899.1| PREDICTED: NADH dehydrogenase (ubiquinone) complex I, assemb
>gi|1036732299|ref|XP_017151071.1| PREDICTED: glycine-rich selenoprotein [Drosophila miranda]
>gi|1036732220|ref|XP_017151039.1| PREDICTED: anoctamin-6 isoform X1 [Drosophila miranda]
>gi|1036732243|ref|XP_017151048.1| PREDICTED: anoctamin-6 isoform X1 [Drosophila miranda]
>gi|1036732281|ref|XP_017151065.1| PREDICTED: anoctamin-6 isoform X3 [Drosophila miranda]
>gi|1036732264|ref|XP_017151057.1| PREDICTED: anoctamin-6 isoform X2 [Drosophila miranda]
>gi|1036752810|ref|XP_017156458.1| PREDICTED: 4-nitrophenylphosphatase [Drosophila miranda]
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>gi|1036752791|ref|XP_017156457.1| PREDICTED: pyridoxal phosphate phosphatase [Drosophila mira
>gi|1036752482|ref|XP_017156439.1| PREDICTED: loss of heterozygosity 12 chromosomal region 1 p
>gi|1036751699|ref|XP_017156390.1| PREDICTED: AH receptor-interacting protein [Drosophila mira
>gi|1036747387|ref|XP_017156165.1| PREDICTED: SH3 domain-binding protein 5-like isoform X2 [Dro
>gi|1036747368|ref|XP_017156164.1| PREDICTED: keratin, type II cytoskeletal 1 isoform X1 [Droso
>gi|1036732370|ref|XP_017151115.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732388|ref|XP_017151121.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732407|ref|XP_017151128.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732447|ref|XP_017151142.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732483|ref|XP_017151158.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732426|ref|XP_017151134.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732464|ref|XP_017151149.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732500|ref|XP_017151166.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036705277|ref|XP_017135496.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036703079|ref|XP_017156747.1| PREDICTED: venom allergen 5-like [Drosophila miranda]
>gi|1036703214|ref|XP_017156818.1| PREDICTED: venom allergen 5-like [Drosophila miranda]
>gi|1036751106|ref|XP_017156359.1| PREDICTED: solute carrier family 46 member 3 [Drosophila mi
>gi|1036735975|ref|XP_017153058.1| PREDICTED: netrin-B [Drosophila miranda]
>gi|1036703251|ref|XP_017156840.1| PREDICTED: gamma-butyrobetaine dioxygenase [Drosophila mira
>gi|1036736208|ref|XP_017153196.1| PREDICTED: netrin-A isoform X1 [Drosophila miranda]
>gi|1036736226|ref|XP_017153204.1| PREDICTED: netrin-A isoform X2 [Drosophila miranda]
>gi|1036741231|ref|XP_017155816.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X2
>gi|1036741282|ref|XP_017155819.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X5
>gi|1036741213|ref|XP_017155815.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X1
>gi|1036741248|ref|XP_017155817.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X3
>gi|1036741265|ref|XP_017155818.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X4
>gi|1036741343|ref|XP_017155823.1| PREDICTED: nucleoside diphosphate kinase 6 [Drosophila mira
>gi|1036741326|ref|XP_017155822.1| PREDICTED: probable 28S ribosomal protein S25, mitochondrial
>gi|1036741299|ref|XP_017155820.1| PREDICTED: probable RNA-binding protein 18 isoform X1 [Droso
>gi|1036741313|ref|XP_017155821.1| PREDICTED: probable RNA-binding protein 18 isoform X2 [Droso
>gi|1036706228|ref|XP_017136158.1| PREDICTED: uncharacterized protein LOC108151811 [Drosophila
>gi|1036699216|ref|XP_017139141.1| PREDICTED: uncharacterized protein LOC108153545 [Drosophila
>gi|1036758052|ref|XP_017156753.1| PREDICTED: atrophin-1 [Drosophila miranda]
>gi|1036746085|ref|XP_017156085.1| PREDICTED: titin homolog isoform X2 [Drosophila miranda]
>gi|1036746028|ref|XP_017156081.1| PREDICTED: titin homolog isoform X1 [Drosophila miranda]
>gi|1036746047|ref|XP_017156082.1| PREDICTED: titin homolog isoform X1 [Drosophila miranda]
>gi|1036746066|ref|XP_017156083.1| PREDICTED: titin homolog isoform X1 [Drosophila miranda]
>gi|1036746103|ref|XP_017156086.1| PREDICTED: protein TsetseEP isoform X1 [Drosophila miranda]
>gi|1036746139|ref|XP_017156088.1| PREDICTED: transcriptional regulatory protein AlgP isoform
>gi|1036746120|ref|XP_017156087.1| PREDICTED: protein TsetseEP isoform X2 [Drosophila miranda]
>gi|1036745996|ref|XP_017156079.1| PREDICTED: uncharacterized protein LOC108164713 [Drosophila
>gi|1036746011|ref|XP_017156080.1| PREDICTED: uncharacterized protein LOC108164713 [Drosophila
>gi|1036703967|ref|XP_017134609.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036699231|ref|XP_017139260.1| PREDICTED: uncharacterized protein LOC108153628 [Drosophila
>gi|1036732853|ref|XP_017151354.1| PREDICTED: transcription factor TBF1-like [Drosophila miran
>gi|1036732871|ref|XP_017151361.1| PREDICTED: transcription factor TBF1-like [Drosophila miran
>gi|1036712798|ref|XP_017140241.1| PREDICTED: sodium leak channel non-selective protein isofor
>gi|1036712764|ref|XP_017140226.1| PREDICTED: sodium leak channel non-selective protein isofor
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>gi|1036712781|ref|XP_017140234.1| PREDICTED: sodium leak channel non-selective protein isofor
>gi|1036712832|ref|XP_017140273.1| PREDICTED: plasminogen receptor (KT) [Drosophila miranda]
>gi|1036712815|ref|XP_017140258.1| PREDICTED: tyrosine aminotransferase [Drosophila miranda]
>gi|1036712747|ref|XP_017140216.1| PREDICTED: golgin subfamily B member 1 [Drosophila miranda]
>gi|1036751473|ref|XP_017156379.1| PREDICTED: uncharacterized protein LOC108164910 isoform X1
>gi|1036751492|ref|XP_017156380.1| PREDICTED: uncharacterized protein LOC108164910 isoform X2
>gi|1036751511|ref|XP_017156382.1| PREDICTED: uncharacterized protein LOC108164910 isoform X2
>gi|1036714882|ref|XP_017141522.1| PREDICTED: mRNA-capping enzyme [Drosophila miranda]
>gi|1036714937|ref|XP_017141563.1| PREDICTED: tRNA-dihydrouridine(20) synthase [NAD(P)+]-like
>gi|1036714950|ref|XP_017141577.1| PREDICTED: vacuolar protein sorting-associated protein 4B [
>gi|1036714965|ref|XP_017141587.1| PREDICTED: vacuolar protein sorting-associated protein 4B [
>gi|1036714981|ref|XP_017141595.1| PREDICTED: vacuolar protein sorting-associated protein 4B [
>gi|1036714920|ref|XP_017141550.1| PREDICTED: uncharacterized protein LOC108155323 [Drosophila
>gi|1036714903|ref|XP_017141535.1| PREDICTED: serine/threonine-protein kinase ATR [Drosophila m
>gi|1036734889|ref|XP_017152458.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734907|ref|XP_017152466.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734761|ref|XP_017152404.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734923|ref|XP_017152473.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734871|ref|XP_017152450.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734852|ref|XP_017152441.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734816|ref|XP_017152426.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734833|ref|XP_017152434.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734960|ref|XP_017152490.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734780|ref|XP_017152408.1| PREDICTED: uncharacterized protein LOC108162264 isoform X2
>gi|1036734944|ref|XP_017152482.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734797|ref|XP_017152419.1| PREDICTED: dual specificity protein kinase splA isoform X3
>gi|1036734979|ref|XP_017152503.1| PREDICTED: uncharacterized protein LOC108162326 isoform X1
>gi|1036734998|ref|XP_017152510.1| PREDICTED: uncharacterized protein LOC108162326 isoform X2
>gi|1036699246|ref|XP_017139376.1| PREDICTED: axonemal 84 kDa protein [Drosophila miranda]
>gi|1036718432|ref|XP_017143530.1| PREDICTED: dynein heavy chain 6, axonemal [Drosophila mirand
>gi|1036718451|ref|XP_017143542.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036718474|ref|XP_017143551.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036718518|ref|XP_017143576.1| PREDICTED: transcriptional adapter 3 [Drosophila miranda]
>gi|1036718556|ref|XP_017143600.1| PREDICTED: 3-hydroxyacyl-CoA dehydrogenase type-2 [Drosophil
>gi|1036718495|ref|XP_017143563.1| PREDICTED: zinc finger protein 132-like [Drosophila miranda]
>gi|1036718536|ref|XP_017143590.1| PREDICTED: rho GTPase-activating protein 15 [Drosophila mira
>gi|1036735017|ref|XP_017152522.1| PREDICTED: multivesicular body subunit 12A isoform X1 [Droso
>gi|1036735033|ref|XP_017152531.1| PREDICTED: multivesicular body subunit 12A isoform X2 [Droso
>gi|1036735051|ref|XP_017152545.1| PREDICTED: transcription initiation factor TFIID subunit 8
>gi|1036736366|ref|XP_017153295.1| PREDICTED: amyloid protein-binding protein 2 isoform X1 [Dro
>gi|1036736383|ref|XP_017153303.1| PREDICTED: amyloid protein-binding protein 2 isoform X2 [Dro
>gi|1036748202|ref|XP_017156210.1| PREDICTED: uncharacterized protein LOC108164797 isoform X1
>gi|1036748221|ref|XP_017156211.1| PREDICTED: uncharacterized protein LOC108164797 isoform X2
>gi|1036748240|ref|XP_017156212.1| PREDICTED: type-1 angiotensin II receptor-associated protei
>gi|1036740280|ref|XP_017155437.1| PREDICTED: uncharacterized protein LOC108164297 [Drosophila
>gi|1036740262|ref|XP_017155426.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036724452|ref|XP_017146586.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1036724469|ref|XP_017146594.1| PREDICTED: neprilysin-2 [Drosophila miranda]
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>gi|1036724486|ref|XP_017146603.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1036724505|ref|XP_017146615.1| PREDICTED: multiple inositol polyphosphate phosphatase 1 [D:
>gi|1036724436|ref|XP_017146576.1| PREDICTED: regulatory-associated protein of mTOR [Drosophila
>gi|1036756986|ref|XP_017156699.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036757005|ref|XP_017156700.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036757024|ref|XP_017156701.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036754997|ref|XP_017156591.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036754867|ref|XP_017156583.1| PREDICTED: tetraspanin-9 [Drosophila miranda]
>gi|1036754886|ref|XP_017156584.1| PREDICTED: uncharacterized protein LOC108165054 [Drosophila
>gi|1036704420|ref|XP_017134880.1| PREDICTED: uncharacterized protein LOC108151028 [Drosophila
>gi|1036751718|ref|XP_017156391.1| PREDICTED: hyphally regulated cell wall protein 3 [Drosophil
>gi|1036743445|ref|XP_017155944.1| PREDICTED: brefeldin A-inhibited guanine nucleotide-exchange
>gi|1036752773|ref|XP_017156456.1| PREDICTED: malignant T-cell-amplified sequence 1 homolog [D:
>gi|1036752230|ref|XP_017156423.1| PREDICTED: cell cycle negative regulator roughex [Drosophila
>gi|1036751899|ref|XP_017156403.1| PREDICTED: protein mab-21 [Drosophila miranda]
>gi|1036754710|ref|XP_017156574.1| PREDICTED: protein mab-21-like [Drosophila miranda]
>gi|1036754725|ref|XP_017156575.1| PREDICTED: protein mab-21-like [Drosophila miranda]
>gi|1036750957|ref|XP_017156349.1| PREDICTED: pancreatic triacylglycerol lipase [Drosophila mi:
>gi|1036741159|ref|XP_017155811.1| PREDICTED: glutamate receptor-interacting protein 1 isoform
>gi|1036741141|ref|XP_017155810.1| PREDICTED: glutamate receptor-interacting protein 1 isoform
>gi|1036740711|ref|XP_017155683.1| PREDICTED: uncharacterized protein LOC108164460 [Drosophila
>gi|1036740729|ref|XP_017155697.1| PREDICTED: uncharacterized protein LOC108164466 [Drosophila
>gi|1036706194|ref|XP_017136132.1| PREDICTED: trypsin eta-like [Drosophila miranda]
>gi|1036704001|ref|XP_017134631.1| PREDICTED: trypsin-1-like [Drosophila miranda]
>gi|1036745287|ref|XP_017156037.1| PREDICTED: trypsin I-P1 isoform X1 [Drosophila miranda]
>gi|1036745304|ref|XP_017156039.1| PREDICTED: trypsin I-P1 isoform X2 [Drosophila miranda]
>gi|1036745321|ref|XP_017156040.1| PREDICTED: trypsin I-P1 isoform X3 [Drosophila miranda]
>gi|1036745338|ref|XP_017156041.1| PREDICTED: uncharacterized protein LOC108164685 isoform X4
>gi|1036745354|ref|XP_017156042.1| PREDICTED: urokinase-type plasminogen activator isoform X5
>gi|1036745370|ref|XP_017156043.1| PREDICTED: trypsin eta isoform X1 [Drosophila miranda]
>gi|1036745387|ref|XP_017156044.1| PREDICTED: trypsin eta isoform X2 [Drosophila miranda]
>gi|1036745404|ref|XP_017156045.1| PREDICTED: trypsin eta isoform X3 [Drosophila miranda]
>gi|1036745421|ref|XP_017156046.1| PREDICTED: uncharacterized protein LOC108164687 [Drosophila
>gi|1036745233|ref|XP_017156034.1| PREDICTED: protein bric-a-brac 1 isoform X1 [Drosophila mira
>gi|1036745269|ref|XP_017156036.1| PREDICTED: protein bric-a-brac 1 isoform X3 [Drosophila mira
>gi|1036745252|ref|XP_017156035.1| PREDICTED: protein bric-a-brac 1 isoform X2 [Drosophila mira
>gi|1036712511|ref|XP_017140078.1| PREDICTED: serine hydroxymethyltransferase, cytosolic [Dros
>gi|1036712594|ref|XP_017140128.1| PREDICTED: transmembrane protein 256 homolog [Drosophila mi:
>gi|1036712528|ref|XP_017140088.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036712545|ref|XP_017140096.1| PREDICTED: modifier of mdg4 isoform X2 [Drosophila miranda]
>gi|1036712561|ref|XP_017140108.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosop]
>gi|1036712577|ref|XP_017140118.1| PREDICTED: actin-5C [Drosophila miranda]
>gi|1036742251|ref|XP_017155874.1| PREDICTED: serine/threonine-protein phosphatase beta isoform
>gi|1036742233|ref|XP_017155873.1| PREDICTED: serine/threonine-protein phosphatase beta isoform
>gi|1036755592|ref|XP_017156621.1| PREDICTED: uncharacterized protein LOC108165082 isoform X1
>gi|1036755609|ref|XP_017156622.1| PREDICTED: uncharacterized protein LOC108165082 isoform X2
>gi|1036703787|ref|XP_017157160.1| PREDICTED: sphingomyelin synthase-related 1-like [Drosophila
>gi|1036699261|ref|XP_017139456.1| PREDICTED: odorant receptor 9a [Drosophila miranda]
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>gi|1036743789|ref|XP_017155959.1| PREDICTED: uncharacterized protein LOC108164624 [Drosophila
>gi|1036743624|ref|XP_017155952.1| PREDICTED: uncharacterized protein LOC108164621 isoform X4
>gi|1036743549|ref|XP_017155948.1| PREDICTED: uncharacterized protein LOC108164621 isoform X1
>gi|1036743568|ref|XP_017155949.1| PREDICTED: uncharacterized protein LOC108164621 isoform X1
>gi|1036743587|ref|XP_017155950.1| PREDICTED: uncharacterized protein LOC108164621 isoform X2
>gi|1036743606|ref|XP_017155951.1| PREDICTED: uncharacterized protein LOC108164621 isoform X3
>gi|1036743716|ref|XP_017155957.1| PREDICTED: uncharacterized protein LOC108164621 isoform X8
>gi|1036743643|ref|XP_017155953.1| PREDICTED: uncharacterized protein LOC108164621 isoform X5
>gi|1036743661|ref|XP_017155954.1| PREDICTED: uncharacterized protein LOC108164621 isoform X6
>gi|1036743680|ref|XP_017155955.1| PREDICTED: uncharacterized protein LOC108164621 isoform X6
>gi|1036743699|ref|XP_017155956.1| PREDICTED: uncharacterized protein LOC108164621 isoform X7
>gi|1036748257|ref|XP_017156213.1| PREDICTED: up-regulated during skeletal muscle growth prote
>gi|1036719849|ref|XP_017144338.1| PREDICTED: neurogenic protein mastermind [Drosophila miranda
>gi|1036719827|ref|XP_017144316.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036733434|ref|XP_017151633.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1036733452|ref|XP_017151641.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1036733470|ref|XP_017151650.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1036733489|ref|XP_017151662.1| PREDICTED: leucine-rich repeat-containing protein 58 [Droso
>gi|1036722475|ref|XP_017145554.1| PREDICTED: protein sprint isoform X3 [Drosophila miranda]
>gi|1036722494|ref|XP_017145555.1| PREDICTED: protein sprint isoform X3 [Drosophila miranda]
>gi|1036722512|ref|XP_017145556.1| PREDICTED: protein sprint isoform X4 [Drosophila miranda]
>gi|1036722437|ref|XP_017145540.1| PREDICTED: protein sprint isoform X1 [Drosophila miranda]
>gi|1036722456|ref|XP_017145548.1| PREDICTED: protein sprint isoform X2 [Drosophila miranda]
>gi|1036722529|ref|XP_017145557.1| PREDICTED: protein sprint isoform X5 [Drosophila miranda]
>gi|1036722550|ref|XP_017145558.1| PREDICTED: protein sprint isoform X5 [Drosophila miranda]
>gi|1036722569|ref|XP_017145559.1| PREDICTED: protein sprint isoform X6 [Drosophila miranda]
>gi|1036722646|ref|XP_017145587.1| PREDICTED: uncharacterized protein LOC108157950 [Drosophila
>gi|1036722667|ref|XP_017145595.1| PREDICTED: uncharacterized protein LOC108157950 [Drosophila
>gi|1036722592|ref|XP_017145560.1| PREDICTED: microspherule protein 1-like [Drosophila miranda]
>gi|1036722609|ref|XP_017145567.1| PREDICTED: microspherule protein 1-like [Drosophila miranda]
>gi|1036722682|ref|XP_017145606.1| PREDICTED: uncharacterized protein LOC108157968 [Drosophila
>gi|1036722628|ref|XP_017145576.1| PREDICTED: uncharacterized protein F09G8.5-like [Drosophila
>gi|1036706161|ref|XP_017136113.1| PREDICTED: GTP-binding protein ypt1-like [Drosophila miranda
>gi|1036731735|ref|XP_017150828.1| PREDICTED: uncharacterized protein LOC108161083 [Drosophila
>gi|1036738937|ref|XP_017154693.1| PREDICTED: FAST kinase domain-containing protein 5, mitochor
>gi|1036738953|ref|XP_017154706.1| PREDICTED: LOW QUALITY PROTEIN: mitochondrial ornithine tra
>gi|1036751088|ref|XP_017156357.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036741801|ref|XP_017155852.1| PREDICTED: uncharacterized protein LOC108164549 [Drosophila
>gi|1036741693|ref|XP_017155845.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741710|ref|XP_017155846.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741731|ref|XP_017155847.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741747|ref|XP_017155848.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741768|ref|XP_017155849.1| PREDICTED: uncharacterized protein LOC108164547 isoform X2
>gi|1036741784|ref|XP_017155850.1| PREDICTED: uncharacterized protein LOC108164548 [Drosophila
>gi|1036705425|ref|XP_017135596.1| PREDICTED: rRNA biogenesis protein rrp36-like [Drosophila m
>gi|1036725961|ref|XP_017147489.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase a
>gi|1036725984|ref|XP_017147497.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase a
>gi|1036725923|ref|XP_017147468.1| PREDICTED: BTB/POZ domain-containing protein 9 isoform X1 [
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>gi|1036725942|ref|XP_017147476.1| PREDICTED: BTB/POZ domain-containing protein 9 isoform X2 [
>gi|1036717963|ref|XP_017143290.1| PREDICTED: uncharacterized protein LOC108156382 [Drosophila
>gi|1036717885|ref|XP_017143246.1| PREDICTED: ADP,ATP carrier protein [Drosophila miranda]
>gi|1036717904|ref|XP_017143257.1| PREDICTED: ADP,ATP carrier protein [Drosophila miranda]
>gi|1036717926|ref|XP_017143266.1| PREDICTED: ADP,ATP carrier protein [Drosophila miranda]
>gi|1036717866|ref|XP_017143234.1| PREDICTED: protein penguin [Drosophila miranda]
>gi|1036717944|ref|XP_017143278.1| PREDICTED: putative peptidyl-prolyl cis-trans isomerase dod
>gi|1036728836|ref|XP_017149195.1| PREDICTED: sphingosine kinase 2 [Drosophila miranda]
>gi|1036728748|ref|XP_017149147.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728767|ref|XP_017149154.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728784|ref|XP_017149163.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728801|ref|XP_017149173.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728819|ref|XP_017149182.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036746435|ref|XP_017156107.1| PREDICTED: glutamine synthetase 2 cytoplasmic isoform X3 [D
>gi|1036746403|ref|XP_017156105.1| PREDICTED: glutamine synthetase 2 cytoplasmic isoform X1 [D
>gi|1036746420|ref|XP_017156106.1| PREDICTED: glutamine synthetase 2 cytoplasmic isoform X2 [D
>gi|1036746453|ref|XP_017156108.1| PREDICTED: odorant receptor 10a [Drosophila miranda]
>gi|1036746385|ref|XP_017156103.1| PREDICTED: gustatory receptor 10a [Drosophila miranda]
>gi|1036707473|ref|XP_017136981.1| PREDICTED: putative gustatory receptor 10b [Drosophila mira
>gi|1036715328|ref|XP_017141794.1| PREDICTED: uncharacterized protein LOC108155479 [Drosophila
>gi|1036715568|ref|XP_017141910.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715549|ref|XP_017141902.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715511|ref|XP_017141885.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715475|ref|XP_017141866.1| PREDICTED: disks large 1 tumor suppressor protein isoform X8
>gi|1036715606|ref|XP_017141927.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715458|ref|XP_017141857.1| PREDICTED: disks large 1 tumor suppressor protein isoform X7
>gi|1036715437|ref|XP_017141848.1| PREDICTED: disks large 1 tumor suppressor protein isoform X6
>gi|1036715419|ref|XP_017141841.1| PREDICTED: disks large 1 tumor suppressor protein isoform X5
>gi|1036715402|ref|XP_017141833.1| PREDICTED: disks large 1 tumor suppressor protein isoform X4
>gi|1036715383|ref|XP_017141823.1| PREDICTED: disks large 1 tumor suppressor protein isoform X3
>gi|1036715366|ref|XP_017141814.1| PREDICTED: disks large 1 tumor suppressor protein isoform X2
>gi|1036715347|ref|XP_017141805.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715587|ref|XP_017141918.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715530|ref|XP_017141893.1| PREDICTED: disks large 1 tumor suppressor protein isoform X
>gi|1036715494|ref|XP_017141876.1| PREDICTED: disks large 1 tumor suppressor protein isoform X9
>gi|1036715624|ref|XP_017141939.1| PREDICTED: uncharacterized protein LOC108155555 [Drosophila
>gi|1036752008|ref|XP_017156410.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036732597|ref|XP_017151220.1| PREDICTED: tyrosine-protein kinase hopscotch [Drosophila mi
>gi|1036751627|ref|XP_017156388.1| PREDICTED: uncharacterized protein LOC108164913 [Drosophila
>gi|1036743173|ref|XP_017155928.1| PREDICTED: segment polarity protein dishevelled [Drosophila
>gi|1036724085|ref|XP_017146341.1| PREDICTED: uncharacterized protein LOC108158504 [Drosophila
>gi|1036724143|ref|XP_017146384.1| PREDICTED: uncharacterized protein LOC108158514 [Drosophila
>gi|1036724124|ref|XP_017146371.1| PREDICTED: uncharacterized protein C1orf131 [Drosophila mira
>gi|1036724106|ref|XP_017146356.1| PREDICTED: alanine--glyoxylate aminotransferase 2-like [Dros
>gi|1036699276|ref|XP_017139662.1| PREDICTED: E3 ubiquitin-protein ligase RFWD3-like, partial
>gi|1036733820|ref|XP_017151829.1| PREDICTED: putative vitellogenin receptor isoform X2 [Drosop
>gi|1036733801|ref|XP_017151821.1| PREDICTED: putative vitellogenin receptor isoform X1 [Drosop
>gi|1036733890|ref|XP_017151883.1| PREDICTED: uncharacterized protein LOC108161948 [Drosophila
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>gi|1036733947|ref|XP_017151921.1| PREDICTED: prisilkin-39 [Drosophila miranda]
>gi|1036733873|ref|XP_017151869.1| PREDICTED: probable ATP-dependent RNA helicase ddx17 [Droso
>gi|1036733928|ref|XP_017151910.1| PREDICTED: uncharacterized protein LOC108161959 [Drosophila
>gi|1036733855|ref|XP_017151856.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
>gi|1036699291|ref|XP_017139786.1| PREDICTED: uncharacterized protein LOC108154124 [Drosophila
>gi|1036699305|ref|XP_017139917.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036699320|ref|XP_017140034.1| PREDICTED: proline-rich protein HaeIII subfamily 1 [Drosophi
>gi|1036733909|ref|XP_017151898.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
>gi|1036733837|ref|XP_017151842.1| PREDICTED: hyphal wall protein 1 [Drosophila miranda]
>gi|1036707092|ref|XP_017136749.1| PREDICTED: MICOS complex subunit Mic10 [Drosophila miranda]
>gi|1036704766|ref|XP_017135126.1| PREDICTED: probable mitochondrial import inner membrane tra
>gi|1036705546|ref|XP_017135668.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036705072|ref|XP_017135358.1| PREDICTED: uncharacterized protein LOC108151319 [Drosophila
>gi|1036706211|ref|XP_017136146.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036705102|ref|XP_017135382.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036704941|ref|XP_017135256.1| PREDICTED: uncharacterized protein LOC108151248 [Drosophila
>gi|1036704956|ref|XP_017135264.1| PREDICTED: uncharacterized protein LOC108151248 [Drosophila
>gi|1036699335|ref|XP_017140145.1| PREDICTED: uncharacterized protein LOC108154357 [Drosophila
>gi|1036744345|ref|XP_017155986.1| PREDICTED: ubiquitin-conjugating enzyme E2 N [Drosophila mi
>gi|1036744363|ref|XP_017155987.1| PREDICTED: ubiquitin-conjugating enzyme E2 N [Drosophila mi
>gi|1036700935|ref|XP_017152298.1| PREDICTED: uncharacterized protein LOC108162178 [Drosophila
>gi|1036701120|ref|XP_017153709.1| PREDICTED: uncharacterized protein LOC108163093 [Drosophila
>gi|1036700721|ref|XP_017150989.1| PREDICTED: uncharacterized protein LOC108161245 [Drosophila
>gi|1036700662|ref|XP_017150645.1| PREDICTED: AMP deaminase 2 isoform X2 [Drosophila miranda]
>gi|1036700645|ref|XP_017150564.1| PREDICTED: AMP deaminase 2 isoform X1 [Drosophila miranda]
>gi|1036700677|ref|XP_017150723.1| PREDICTED: AMP deaminase 2 isoform X3 [Drosophila miranda]
>gi|1036700707|ref|XP_017150886.1| PREDICTED: LIM domain-containing protein jub [Drosophila mi
>gi|1036700614|ref|XP_017150318.1| PREDICTED: nuclear pore complex protein Nup93 [Drosophila m
>gi|1036700983|ref|XP_017152620.1| PREDICTED: chloride intracellular channel exc-4 isoform X2
>gi|1036700969|ref|XP_017152540.1| PREDICTED: chloride intracellular channel exc-4 isoform X1
>gi|1036700954|ref|XP_017152418.1| PREDICTED: nucleoside diphosphate-linked moiety X motif 8 [
>gi|1036700600|ref|XP_017150194.1| PREDICTED: AP-1 complex subunit beta-1 [Drosophila miranda]
>gi|1036700868|ref|XP_017151767.1| PREDICTED: transmembrane protein 198 [Drosophila miranda]
>gi|1036701105|ref|XP_017153584.1| PREDICTED: cytochrome b-c1 complex subunit 10 [Drosophila m
>gi|1036701090|ref|XP_017153460.1| PREDICTED: cytochrome c oxidase subunit 6B1 [Drosophila mira
>gi|1036700882|ref|XP_017151897.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1036700895|ref|XP_017151980.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1036700907|ref|XP_017152065.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1036701045|ref|XP_017153097.1| PREDICTED: ribonuclease P protein subunit p20 [Drosophila m
>gi|1036701075|ref|XP_017153340.1| PREDICTED: transmembrane protein 203 [Drosophila miranda]
>gi|1036700922|ref|XP_017152189.1| PREDICTED: beta-parvin [Drosophila miranda]
>gi|1036700998|ref|XP_017152732.1| PREDICTED: FAD-linked sulfhydryl oxidase ALR [Drosophila mi
>gi|1036701060|ref|XP_017153217.1| PREDICTED: uncharacterized protein LOC108162787 [Drosophila
>gi|1036700693|ref|XP_017150824.1| PREDICTED: uncharacterized protein LOC108161006 [Drosophila
>gi|1036700395|ref|XP_017148872.1| PREDICTED: nuclear pore complex protein Nup153 isoform X1 [
>gi|1036700409|ref|XP_017148950.1| PREDICTED: nuclear pore complex protein Nup153 isoform X1 [
>gi|1036700424|ref|XP_017149035.1| PREDICTED: nuclear pore complex protein Nup153 isoform X2 [
>gi|1036700453|ref|XP_017149204.1| PREDICTED: nuclear pore complex protein Nup153 isoform X4 [
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>gi|1036700438|ref|XP_017149115.1| PREDICTED: nuclear pore complex protein Nup153 isoform X3 [1
>gi|1036700736|ref|XP_017151097.1| PREDICTED: serine/threonine-protein kinase PAK mbt [Drosophi
>gi|1036700336|ref|XP_017148507.1| PREDICTED: homeotic protein female sterile isoform X5 [Drosc
>gi|1036700307|ref|XP_017148328.1| PREDICTED: homeotic protein female sterile isoform X3 [Drosc
>gi|1036700292|ref|XP_017148241.1| PREDICTED: homeotic protein female sterile isoform X2 [Drosc
>gi|1036700321|ref|XP_017148412.1| PREDICTED: homeotic protein female sterile isoform X4 [Drosc
>gi|1036700202|ref|XP_017147727.1| PREDICTED: homeotic protein female sterile isoform X1 [Drosc
>gi|1036700216|ref|XP_017147815.1| PREDICTED: homeotic protein female sterile isoform X1 [Drosc
>gi|1036700229|ref|XP_017147901.1| PREDICTED: homeotic protein female sterile isoform X1 [Drosc
>gi|1036700242|ref|XP_017147986.1| PREDICTED: homeotic protein female sterile isoform X1 [Drosc
>gi|1036700256|ref|XP_017148072.1| PREDICTED: homeotic protein female sterile isoform X1 [Drosc
>gi|1036700273|ref|XP_017148151.1| PREDICTED: homeotic protein female sterile isoform X1 [Drosc
>gi|1036700353|ref|XP_017148586.1| PREDICTED: homeotic protein female sterile isoform X6 [Drosc
>gi|1036700381|ref|XP_017148753.1| PREDICTED: homeotic protein female sterile isoform X8 [Drosc
>gi|1036700367|ref|XP_017148671.1| PREDICTED: homeotic protein female sterile isoform X7 [Drosc
>gi|1036700826|ref|XP_017151493.1| PREDICTED: multifunctional protein ADE2 [Drosophila miranda]
>gi|1036700630|ref|XP_017150441.1| PREDICTED: TBC domain-containing protein kinase-like protein
>gi|1036700811|ref|XP_017151391.1| PREDICTED: heparan-alpha-glucosaminide N-acetyltransferase
>gi|1036700750|ref|XP_017151204.1| PREDICTED: uncharacterized protein LOC108161422 [Drosophila
>gi|1036700787|ref|XP_017151279.1| PREDICTED: uncharacterized protein LOC108161422 [Drosophila
>gi|1036700844|ref|XP_017151607.1| PREDICTED: zinc finger BED domain-containing protein 4-like
>gi|1036700526|ref|XP_017149660.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X3 [Dros
>gi|1036700468|ref|XP_017149328.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036700483|ref|XP_017149411.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036700497|ref|XP_017149493.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036700511|ref|XP_017149577.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X2 [Dros
>gi|1036701028|ref|XP_017152973.1| PREDICTED: uncharacterized protein LOC108162621 [Drosophila
>gi|1036701013|ref|XP_017152853.1| PREDICTED: CTD nuclear envelope phosphatase 1A-like [Drosoph
>gi|1036700541|ref|XP_017149790.1| PREDICTED: protein ovo isoform X1 [Drosophila miranda]
>gi|1036700555|ref|XP_017149875.1| PREDICTED: protein ovo isoform X2 [Drosophila miranda]
>gi|1036700570|ref|XP_017149993.1| PREDICTED: zinc finger protein 236 [Drosophila miranda]
>gi|1036700585|ref|XP_017150075.1| PREDICTED: zinc finger protein 236 [Drosophila miranda]
>gi|1036739071|ref|XP_017154762.1| PREDICTED: innexin inx1 [Drosophila miranda]
>gi|1036739089|ref|XP_017154770.1| PREDICTED: innexin inx1 [Drosophila miranda]
>gi|1036753592|ref|XP_017156507.1| PREDICTED: uncharacterized protein LOC108164993 [Drosophila
>gi|1036755344|ref|XP_017156611.1| PREDICTED: innexin inx7 [Drosophila miranda]
>gi|1036732044|ref|XP_017150958.1| PREDICTED: innexin inx2 [Drosophila miranda]
>gi|1036736115|ref|XP_017153134.1| PREDICTED: mitochondrial import receptor subunit TOM7 homolo
>gi|1036736151|ref|XP_017153160.1| PREDICTED: AP-3 complex subunit mu-2 [Drosophila miranda]
>gi|1036736174|ref|XP_017153171.1| PREDICTED: small integral membrane protein 4 [Drosophila mir
>gi|1036736190|ref|XP_017153183.1| PREDICTED: uncharacterized protein C12orf73 homolog [Drosoph
>gi|1036736133|ref|XP_017153146.1| PREDICTED: zinc finger protein 782 [Drosophila miranda]
>gi|1036748513|ref|XP_017156216.1| PREDICTED: uncharacterized protein LOC108164807 [Drosophila
>gi|1036748551|ref|XP_017156218.1| PREDICTED: protein nullo [Drosophila miranda]
>gi|1036748532|ref|XP_017156217.1| PREDICTED: uncharacterized protein LOC108164808 [Drosophila
>gi|1036748570|ref|XP_017156219.1| PREDICTED: uncharacterized protein LOC108164810 [Drosophila
>gi|1036731237|ref|XP_017150618.1| PREDICTED: uncharacterized protein LOC108160864 isoform X1
>gi|1036731292|ref|XP_017150646.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
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>gi|1036731307|ref|XP_017150653.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036731273|ref|XP_017150635.1| PREDICTED: uncharacterized protein LOC108160864 isoform X3
>gi|1036731254|ref|XP_017150626.1| PREDICTED: uncharacterized protein LOC108160864 isoform X2
>gi|1036722404|ref|XP_017145521.1| PREDICTED: peptide transporter family 1 [Drosophila miranda]
>gi|1036699352|ref|XP_017140257.1| PREDICTED: uncharacterized protein LOC108154456 [Drosophila
>gi|1036722387|ref|XP_017145509.1| PREDICTED: peptide transporter family 1 isoform X4 [Drosophi
>gi|1036722349|ref|XP_017145494.1| PREDICTED: peptide transporter family 1 isoform X2 [Drosophi
>gi|1036722368|ref|XP_017145502.1| PREDICTED: peptide transporter family 1 isoform X3 [Drosophi
>gi|1036722328|ref|XP_017145485.1| PREDICTED: peptide transporter family 1 isoform X1 [Drosophi
>gi|1036741634|ref|XP_017155841.1| PREDICTED: zinc finger protein ZPR1 [Drosophila miranda]
>gi|1036741672|ref|XP_017155844.1| PREDICTED: uncharacterized protein LOC108164546 [Drosophila
>gi|1036741653|ref|XP_017155843.1| PREDICTED: uncharacterized protein LOC108164545 [Drosophila
>gi|1036722164|ref|XP_017145379.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1036722185|ref|XP_017145391.1| PREDICTED: histone H3.3 [Drosophila miranda]
>gi|1036722145|ref|XP_017145369.1| PREDICTED: basic proline-rich protein [Drosophila miranda]
>gi|1036729638|ref|XP_017149692.1| PREDICTED: frataxin homolog, mitochondrial [Drosophila mira
>gi|1036729619|ref|XP_017149678.1| PREDICTED: uncharacterized protein CG7065 [Drosophila miran
>gi|1036754237|ref|XP_017156546.1| PREDICTED: probable sodium/metabolite cotransporter BASS5,
>gi|1036739338|ref|XP_017154934.1| PREDICTED: transmembrane protein fend [Drosophila miranda]
>gi|1036722204|ref|XP_017145403.1| PREDICTED: LOW QUALITY PROTEIN: AP-1 complex subunit gamma-
>gi|1036722241|ref|XP_017145426.1| PREDICTED: ADP-ribosylation factor-related protein 1 [Drosop
>gi|1036722260|ref|XP_017145438.1| PREDICTED: protein CutA homolog [Drosophila miranda]
>gi|1036722222|ref|XP_017145415.1| PREDICTED: puff-specific protein Bx42 [Drosophila miranda]
>gi|1036731362|ref|XP_017150681.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731447|ref|XP_017150719.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731483|ref|XP_017150735.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731326|ref|XP_017150665.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731344|ref|XP_017150673.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731394|ref|XP_017150696.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731431|ref|XP_017150711.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731377|ref|XP_017150687.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731412|ref|XP_017150704.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731466|ref|XP_017150727.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731518|ref|XP_017150751.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731500|ref|XP_017150743.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036743858|ref|XP_017155961.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036743839|ref|XP_017155960.1| PREDICTED: T-complex protein 1 subunit beta [Drosophila mira
>gi|1036757646|ref|XP_017156735.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036753388|ref|XP_017156495.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036753426|ref|XP_017156498.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036753443|ref|XP_017156499.1| PREDICTED: sporulation-specific protein 15-like isoform X4
>gi|1036753459|ref|XP_017156500.1| PREDICTED: uncharacterized protein LOC108164985 isoform X5
>gi|1036753407|ref|XP_017156496.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036753479|ref|XP_017156501.1| PREDICTED: golgin subfamily A member 6-like protein 10 [Dros
>gi|1036704479|ref|XP_017134919.1| PREDICTED: uncharacterized protein LOC108151052 [Drosophila
>gi|1036705342|ref|XP_017135541.1| PREDICTED: uncharacterized protein LOC108151439 isoform X1
>gi|1036705360|ref|XP_017135550.1| PREDICTED: uncharacterized protein LOC108151439 isoform X2
>gi|1036705060|ref|XP_017135345.1| PREDICTED: uncharacterized protein LOC108151308 isoform X2
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>gi|1036705046|ref|XP_017135338.1| PREDICTED: uncharacterized protein LOC108151308 isoform X1
>gi|1036707523|ref|XP_017137022.1| PREDICTED: chondroadherin [Drosophila miranda]
>gi|1036756414|ref|XP_017156669.1| PREDICTED: innexin inx6 [Drosophila miranda]
>gi|1036756432|ref|XP_017156670.1| PREDICTED: innexin inx6 [Drosophila miranda]
>gi|1036756450|ref|XP_017156671.1| PREDICTED: uncharacterized protein LOC108165120 [Drosophila
>gi|1036740839|ref|XP_017155773.1| PREDICTED: N-acetylglucosamine-6-phosphate deacetylase [Dros
>gi|1036740856|ref|XP_017155781.1| PREDICTED: ras-related protein Rab-10 [Drosophila miranda]
>gi|1036741107|ref|XP_017155808.1| PREDICTED: uncharacterized protein LOC108164527 [Drosophila
>gi|1036741123|ref|XP_017155809.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036755363|ref|XP_017156612.1| PREDICTED: peritrophin-1 [Drosophila miranda]
>gi|1036713844|ref|XP_017140937.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713861|ref|XP_017140946.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713876|ref|XP_017140954.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713891|ref|XP_017140964.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713913|ref|XP_017140971.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713964|ref|XP_017140998.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713979|ref|XP_017141008.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713930|ref|XP_017140979.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713947|ref|XP_017140987.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714030|ref|XP_017141030.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714071|ref|XP_017141046.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714088|ref|XP_017141052.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713996|ref|XP_017141014.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714013|ref|XP_017141022.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714044|ref|XP_017141037.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714117|ref|XP_017141076.1| PREDICTED: annexin B10 isoform X1 [Drosophila miranda]
>gi|1036714131|ref|XP_017141086.1| PREDICTED: annexin B10 isoform X2 [Drosophila miranda]
>gi|1036714180|ref|XP_017141120.1| PREDICTED: TIP41-like protein isoform X2 [Drosophila miranda
>gi|1036714162|ref|XP_017141111.1| PREDICTED: TIP41-like protein isoform X1 [Drosophila miranda
>gi|1036714103|ref|XP_017141064.1| PREDICTED: probable Xaa-Pro aminopeptidase 3 [Drosophila mir
>gi|1036714146|ref|XP_017141099.1| PREDICTED: delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mito
>gi|1036714216|ref|XP_017141140.1| PREDICTED: PHD finger protein 7 [Drosophila miranda]
>gi|1036742216|ref|XP_017155872.1| PREDICTED: ras-related protein Rab-35 [Drosophila miranda]
>gi|1036754076|ref|XP_017156536.1| PREDICTED: neurogenic locus notch homolog protein 3 isoform
>gi|1036754057|ref|XP_017156535.1| PREDICTED: neurogenic locus notch homolog protein 3 isoform
>gi|1036754039|ref|XP_017156534.1| PREDICTED: neurogenic locus notch homolog protein 3 isoform
>gi|1036699367|ref|XP_017140365.1| PREDICTED: putative odorant receptor 19b [Drosophila miranda
>gi|1036699381|ref|XP_017140474.1| PREDICTED: serine/threonine-protein kinase Warts, partial [D
>gi|1036704989|ref|XP_017135290.1| PREDICTED: uncharacterized protein LOC108151272 [Drosophila
>gi|1036699395|ref|XP_017140596.1| PREDICTED: uncharacterized protein LOC108154714 [Drosophila
>gi|1036707438|ref|XP_017136960.1| PREDICTED: uncharacterized protein LOC108152259 [Drosophila
>gi|1036699410|ref|XP_017140835.1| PREDICTED: uncharacterized protein LOC108154869 [Drosophila
>gi|1036752500|ref|XP_017156440.1| PREDICTED: dopamine D2-like receptor isoform X1 [Drosophila
>gi|1036752593|ref|XP_017156445.1| PREDICTED: dopamine D2-like receptor isoform X6 [Drosophila
>gi|1036752538|ref|XP_017156442.1| PREDICTED: dopamine D2-like receptor isoform X3 [Drosophila
>gi|1036752558|ref|XP_017156443.1| PREDICTED: dopamine D2-like receptor isoform X4 [Drosophila
>gi|1036752519|ref|XP_017156441.1| PREDICTED: dopamine D2-like receptor isoform X2 [Drosophila
>gi|1036752577|ref|XP_017156444.1| PREDICTED: dopamine D2-like receptor isoform X5 [Drosophila
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>gi|1036749517|ref|XP_017156270.1| PREDICTED: endothelin-converting enzyme 1 isoform X2 [Droso
>gi|1036749500|ref|XP_017156269.1| PREDICTED: endothelin-converting enzyme 1 isoform X1 [Droso
>gi|1036703827|ref|XP_017134536.1| PREDICTED: glycine-rich cell wall structural protein [Droso
>gi|1036723465|ref|XP_017145989.1| PREDICTED: uncharacterized protein LOC108158260 [Drosophila
>gi|1036744574|ref|XP_017155997.1| PREDICTED: uncharacterized protein LOC108164656 [Drosophila
>gi|1036711778|ref|XP_017139678.1| PREDICTED: nuclear pore complex protein Nup205 isoform X1 [
>gi|1036711795|ref|XP_017139686.1| PREDICTED: nuclear pore complex protein Nup205 isoform X2 [
>gi|1036711812|ref|XP_017139694.1| PREDICTED: ras-associated and pleckstrin homology domains-co
>gi|1036749550|ref|XP_017156272.1| PREDICTED: uncharacterized protein LOC108164854 isoform X1
>gi|1036749569|ref|XP_017156273.1| PREDICTED: uncharacterized protein LOC108164854 isoform X2
>gi|1036731861|ref|XP_017150873.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036726653|ref|XP_017147924.1| PREDICTED: uncharacterized protein LOC108159272 [Drosophila
>gi|1036726671|ref|XP_017147935.1| PREDICTED: ATP-binding cassette sub-family D member 3 [Dros
>gi|1036726689|ref|XP_017147944.1| PREDICTED: ATP-binding cassette sub-family D member 3 [Dros
>gi|1036742734|ref|XP_017155902.1| PREDICTED: bifunctional lysine-specific demethylase and hist
>gi|1036742752|ref|XP_017155903.1| PREDICTED: 39S ribosomal protein L33, mitochondrial [Drosop
>gi|1036721355|ref|XP_017145072.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721297|ref|XP_017145045.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721333|ref|XP_017145062.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721278|ref|XP_017145036.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721314|ref|XP_017145053.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036716143|ref|XP_017142245.1| PREDICTED: protein transport protein sec31 [Drosophila mira
>gi|1036716158|ref|XP_017142259.1| PREDICTED: 60S ribosomal protein L22 [Drosophila miranda]
>gi|1036716232|ref|XP_017142306.1| PREDICTED: eukaryotic translation initiation factor 3 subun
>gi|1036716251|ref|XP_017142315.1| PREDICTED: eukaryotic translation initiation factor 3 subun
>gi|1036716212|ref|XP_017142293.1| PREDICTED: histone-lysine N-methyltransferase SMYD3 [Drosop
>gi|1036716194|ref|XP_017142280.1| PREDICTED: centromere/kinetochore protein zw10 [Drosophila
>gi|1036735957|ref|XP_017153045.1| PREDICTED: chromosome-associated kinesin KIF4 [Drosophila m
>gi|1036734580|ref|XP_017152284.1| PREDICTED: beta-1,4-mannosyltransferase egh [Drosophila mir
>gi|1036734598|ref|XP_017152293.1| PREDICTED: beta-1,4-mannosyltransferase egh [Drosophila mir
>gi|1036729655|ref|XP_017149704.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036729674|ref|XP_017149717.1| PREDICTED: dynein light chain 1, cytoplasmic [Drosophila mi
>gi|1036729693|ref|XP_017149725.1| PREDICTED: dynein light chain 1, cytoplasmic [Drosophila mi
>gi|1036729713|ref|XP_017149733.1| PREDICTED: dynein light chain 1, cytoplasmic [Drosophila mi
>gi|1036729187|ref|XP_017149419.1| PREDICTED: uncharacterized protein LOC108160132 [Drosophila
>gi|1036734331|ref|XP_017152134.1| PREDICTED: uncharacterized protein LOC108162040 isoform X10
>gi|1036734278|ref|XP_017152117.1| PREDICTED: uncharacterized protein LOC108162040 isoform X8
>gi|1036734255|ref|XP_017152109.1| PREDICTED: uncharacterized protein LOC108162040 isoform X7
>gi|1036734222|ref|XP_017152092.1| PREDICTED: uncharacterized protein LOC108162040 isoform X5
>gi|1036734168|ref|XP_017152066.1| PREDICTED: uncharacterized protein LOC108162040 isoform X2
>gi|1036734241|ref|XP_017152100.1| PREDICTED: uncharacterized protein LOC108162040 isoform X6
>gi|1036734185|ref|XP_017152075.1| PREDICTED: uncharacterized protein LOC108162040 isoform X3
>gi|1036734296|ref|XP_017152125.1| PREDICTED: uncharacterized protein LOC108162040 isoform X9
>gi|1036734204|ref|XP_017152084.1| PREDICTED: uncharacterized protein LOC108162040 isoform X4
>gi|1036734150|ref|XP_017152057.1| PREDICTED: uncharacterized protein LOC108162040 isoform X1
>gi|1036743208|ref|XP_017155931.1| PREDICTED: uncharacterized protein DDB_G0271670 [Drosophila
>gi|1036743192|ref|XP_017155929.1| PREDICTED: uncharacterized protein LOC108164602 [Drosophila
>gi|1036740539|ref|XP_017155570.1| PREDICTED: anoctamin-10 [Drosophila miranda]
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>gi|1036740556|ref|XP_017155584.1| PREDICTED: uncharacterized protein LOC108164415 [Drosophila
>gi|1036740575|ref|XP_017155594.1| PREDICTED: endochitinase A-like [Drosophila miranda]
>gi|1036740592|ref|XP_017155607.1| PREDICTED: uncharacterized protein LOC108164430 [Drosophila
>gi|1036740608|ref|XP_017155617.1| PREDICTED: uncharacterized protein LOC108164430 [Drosophila
>gi|1036718667|ref|XP_017143662.1| PREDICTED: DENN domain-containing protein 4C isoform X1 [Dro
>gi|1036718684|ref|XP_017143669.1| PREDICTED: DENN domain-containing protein 4C isoform X2 [Dro
>gi|1036718700|ref|XP_017143684.1| PREDICTED: chromatin assembly factor 1 subunit A-B [Drosophi
>gi|1036706368|ref|XP_017136253.1| PREDICTED: rab-like protein 3 [Drosophila miranda]
>gi|1036746562|ref|XP_017156115.1| PREDICTED: homeotic protein ocelliless [Drosophila miranda]
>gi|1036745865|ref|XP_017156071.1| PREDICTED: uncharacterized protein LOC108164709 isoform X1
>gi|1036745884|ref|XP_017156072.1| PREDICTED: uncharacterized protein LOC108164709 isoform X2
>gi|1036736644|ref|XP_017153461.1| PREDICTED: carbonic anhydrase 2 [Drosophila miranda]
>gi|1036736626|ref|XP_017153450.1| PREDICTED: palmitoyl-protein thioesterase 1 [Drosophila mira
>gi|1036705706|ref|XP_017135771.1| PREDICTED: paired mesoderm homeobox protein 2 [Drosophila mi
>gi|1036740507|ref|XP_017155546.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036740523|ref|XP_017155558.1| PREDICTED: uncharacterized protein LOC108164404 [Drosophila
>gi|1036731792|ref|XP_017150837.1| PREDICTED: uncharacterized protein CG1785 [Drosophila miran
>gi|1036731754|ref|XP_017150829.1| PREDICTED: RNA cytidine acetyltransferase isoform X1 [Drosop
>gi|1036731773|ref|XP_017150830.1| PREDICTED: RNA cytidine acetyltransferase isoform X2 [Drosop
>gi|1036702255|ref|XP_017156245.1| PREDICTED: probable U3 small nucleolar RNA-associated protei
>gi|1036705880|ref|XP_017135899.1| PREDICTED: uncharacterized protein LOC108151668 [Drosophila
>gi|1036747447|ref|XP_017156169.1| PREDICTED: LIM/homeobox protein Lhx1 [Drosophila miranda]
>gi|1036706758|ref|XP_017136542.1| PREDICTED: uncharacterized protein LOC108152047 [Drosophila
>gi|1036706774|ref|XP_017136550.1| PREDICTED: uncharacterized protein LOC108152047 [Drosophila
>gi|1036749202|ref|XP_017156253.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D
>gi|1036749220|ref|XP_017156254.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D
>gi|1036729431|ref|XP_017149561.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036729452|ref|XP_017149569.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036729471|ref|XP_017149578.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036729491|ref|XP_017149586.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036753967|ref|XP_017156530.1| PREDICTED: uroporphyrinogen-III synthase [Drosophila miranda
>gi|1036753984|ref|XP_017156531.1| PREDICTED: uncharacterized protein LOC108165008 [Drosophila
>gi|1036756564|ref|XP_017156677.1| PREDICTED: calumenin-like [Drosophila miranda]
>gi|1036745439|ref|XP_017156047.1| PREDICTED: uncharacterized protein LOC108164688 [Drosophila
>gi|1036745457|ref|XP_017156048.1| PREDICTED: uncharacterized protein LOC108164688 [Drosophila
>gi|1036735406|ref|XP_017152756.1| PREDICTED: protein MAK16 homolog [Drosophila miranda]
>gi|1036735424|ref|XP_017152769.1| PREDICTED: protein enhancer of rudimentary [Drosophila mira
>gi|1036712278|ref|XP_017139951.1| PREDICTED: eye-specific diacylglycerol kinase isoform X14 [
>gi|1036712430|ref|XP_017140039.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036712445|ref|XP_017140051.1| PREDICTED: uncharacterized protein LOC108154304 isoform X1
>gi|1036712462|ref|XP_017140059.1| PREDICTED: uncharacterized protein LOC108154304 isoform X2
>gi|1036712115|ref|XP_017139878.1| PREDICTED: eye-specific diacylglycerol kinase isoform X5 [Di
>gi|1036712165|ref|XP_017139904.1| PREDICTED: eye-specific diacylglycerol kinase isoform X8 [Di
>gi|1036712049|ref|XP_017139841.1| PREDICTED: eye-specific diacylglycerol kinase isoform X1 [Di
>gi|1036712132|ref|XP_017139887.1| PREDICTED: eye-specific diacylglycerol kinase isoform X6 [Di
>gi|1036712082|ref|XP_017139856.1| PREDICTED: eye-specific diacylglycerol kinase isoform X3 [Di
>gi|1036712099|ref|XP_017139867.1| PREDICTED: eye-specific diacylglycerol kinase isoform X4 [Di
>gi|1036712067|ref|XP_017139847.1| PREDICTED: eye-specific diacylglycerol kinase isoform X2 [Di
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>gi|1036712148|ref|XP_017139897.1| PREDICTED: eye-specific diacylglycerol kinase isoform X7 [D
>gi|1036712182|ref|XP_017139913.1| PREDICTED: eye-specific diacylglycerol kinase isoform X9 [D
>gi|1036712247|ref|XP_017139937.1| PREDICTED: eye-specific diacylglycerol kinase isoform X12 [
>gi|1036712199|ref|XP_017139922.1| PREDICTED: eye-specific diacylglycerol kinase isoform X10 [
>gi|1036712216|ref|XP_017139930.1| PREDICTED: eye-specific diacylglycerol kinase isoform X11 [
>gi|1036712396|ref|XP_017140016.1| PREDICTED: nucleosome assembly protein 1-like 1 [Drosophila
>gi|1036712412|ref|XP_017140027.1| PREDICTED: nucleosome assembly protein 1-like 1 [Drosophila
>gi|1036712261|ref|XP_017139944.1| PREDICTED: eye-specific diacylglycerol kinase isoform X13 [
>gi|1036712330|ref|XP_017139978.1| PREDICTED: nostrin isoform X3 [Drosophila miranda]
>gi|1036712364|ref|XP_017139993.1| PREDICTED: nostrin isoform X5 [Drosophila miranda]
>gi|1036712347|ref|XP_017139985.1| PREDICTED: nostrin isoform X4 [Drosophila miranda]
>gi|1036712381|ref|XP_017140002.1| PREDICTED: nostrin isoform X6 [Drosophila miranda]
>gi|1036712295|ref|XP_017139965.1| PREDICTED: uncharacterized protein LOC108154240 isoform X1
>gi|1036712313|ref|XP_017139973.1| PREDICTED: uncharacterized protein LOC108154240 isoform X2
>gi|1036704189|ref|XP_017134744.1| PREDICTED: uncharacterized protein DDB_G0271670-like [Droso
>gi|1036753292|ref|XP_017156490.1| PREDICTED: transcription factor SPT20 homolog isoform X2 [D
>gi|1036753274|ref|XP_017156489.1| PREDICTED: transcription factor SPT20 homolog isoform X1 [D
>gi|1036749623|ref|XP_017156275.1| PREDICTED: acylphosphatase-1 [Drosophila miranda]
>gi|1036749640|ref|XP_017156276.1| PREDICTED: acylphosphatase-1 [Drosophila miranda]
>gi|1036749604|ref|XP_017156274.1| PREDICTED: uncharacterized protein LOC108164856 [Drosophila
>gi|1036749657|ref|XP_017156277.1| PREDICTED: uncharacterized protein LOC108164859 [Drosophila
>gi|1036724197|ref|XP_017146421.1| PREDICTED: U3 small nucleolar RNA-associated protein 6 homol
>gi|1036724233|ref|XP_017146446.1| PREDICTED: chromobox protein homolog 3-like [Drosophila mira
>gi|1036724160|ref|XP_017146396.1| PREDICTED: anaphase-promoting complex subunit 4 isoform X1
>gi|1036724179|ref|XP_017146406.1| PREDICTED: anaphase-promoting complex subunit 4 isoform X2
>gi|1036724212|ref|XP_017146435.1| PREDICTED: uncharacterized protein F54F2.9 [Drosophila mira
>gi|1036746844|ref|XP_017156133.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036749076|ref|XP_017156246.1| PREDICTED: band 7 protein AGAP004871 isoform X1 [Drosophila
>gi|1036749095|ref|XP_017156247.1| PREDICTED: band 7 protein AGAP004871 isoform X2 [Drosophila
>gi|1036749059|ref|XP_017156244.1| PREDICTED: band 7 protein AGAP004871 [Drosophila miranda]
>gi|1036749114|ref|XP_017156248.1| PREDICTED: NADH-cytochrome b5 reductase-like [Drosophila mi
>gi|1036732084|ref|XP_017150975.1| PREDICTED: transmembrane protein 185B [Drosophila miranda]
>gi|1036732121|ref|XP_017150995.1| PREDICTED: barrier-to-autointegration factor-like [Drosophil
>gi|1036732103|ref|XP_017150985.1| PREDICTED: LOW QUALITY PROTEIN: post-GPI attachment to prote
>gi|1036732161|ref|XP_017151012.1| PREDICTED: barrier-to-autointegration factor-like isoform X2
>gi|1036732182|ref|XP_017151019.1| PREDICTED: barrier-to-autointegration factor-like isoform X2
>gi|1036732142|ref|XP_017151005.1| PREDICTED: barrier-to-autointegration factor-like isoform X1
>gi|1036745683|ref|XP_017156062.1| PREDICTED: post-GPI attachment to proteins factor 2-like iso
>gi|1036745702|ref|XP_017156064.1| PREDICTED: post-GPI attachment to proteins factor 2-like iso
>gi|1036728207|ref|XP_017148840.1| PREDICTED: uncharacterized protein LOC108159778 isoform X2
>gi|1036728190|ref|XP_017148831.1| PREDICTED: uncharacterized protein LOC108159778 isoform X1
>gi|1036756844|ref|XP_017156693.1| PREDICTED: uncharacterized protein LOC108165139 [Drosophila
>gi|1036755160|ref|XP_017156600.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036754693|ref|XP_017156572.1| PREDICTED: dentin sialophosphoprotein [Drosophila miranda]
>gi|1036744419|ref|XP_017155991.1| PREDICTED: uncharacterized protein LOC108164650 [Drosophila
>gi|1036744162|ref|XP_017155976.1| PREDICTED: uncharacterized protein LOC108164641 isoform X1
>gi|1036744181|ref|XP_017155977.1| PREDICTED: uncharacterized protein LOC108164641 isoform X2
>gi|1036744200|ref|XP_017155978.1| PREDICTED: uncharacterized protein LOC108164641 isoform X3
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>gi|1036744219|ref|XP_017155979.1| PREDICTED: uncharacterized protein LOC108164641 isoform X3
>gi|1036744236|ref|XP_017155980.1| PREDICTED: uncharacterized protein LOC108164641 isoform X4
>gi|1036744137|ref|XP_017155975.1| PREDICTED: uncharacterized protein LOC108164640 [Drosophila
>gi|1036751381|ref|XP_017156374.1| PREDICTED: uncharacterized protein LOC108164907 [Drosophila
>gi|1036751398|ref|XP_017156375.1| PREDICTED: uncharacterized protein LOC108164907 [Drosophila
>gi|1036751417|ref|XP_017156376.1| PREDICTED: uncharacterized protein LOC108164907 [Drosophila
>gi|1036732516|ref|XP_017151179.1| PREDICTED: uncharacterized protein LOC108161426 [Drosophila
>gi|1036755749|ref|XP_017156631.1| PREDICTED: monocarboxylate transporter 7 isoform X1 [Drosoph
>gi|1036755768|ref|XP_017156632.1| PREDICTED: monocarboxylate transporter 7 isoform X2 [Drosoph
>gi|1036746524|ref|XP_017156113.1| PREDICTED: receptor-binding cancer antigen expressed on SiSo
>gi|1036746505|ref|XP_017156112.1| PREDICTED: uncharacterized protein LOC108164730 [Drosophila
>gi|1036736451|ref|XP_017153348.1| PREDICTED: protein vav isoform X2 [Drosophila miranda]
>gi|1036736433|ref|XP_017153341.1| PREDICTED: protein vav isoform X1 [Drosophila miranda]
>gi|1036736470|ref|XP_017153360.1| PREDICTED: uncharacterized protein LOC108162898 [Drosophila
>gi|1036731700|ref|XP_017150826.1| PREDICTED: rapamycin-insensitive companion of mTOR [Drosophi
>gi|1036731718|ref|XP_017150827.1| PREDICTED: rapamycin-insensitive companion of mTOR [Drosophi
>gi|1036754832|ref|XP_017156581.1| PREDICTED: heparan sulfate glucosamine 3-O-sulfotransferase
>gi|1036720563|ref|XP_017144695.1| PREDICTED: uncharacterized protein LOC108157237 [Drosophila
>gi|1036720585|ref|XP_017144701.1| PREDICTED: uncharacterized protein LOC108157237 [Drosophila
>gi|1036720621|ref|XP_017144726.1| PREDICTED: protein-lysine N-methyltransferase EEF2KMT [Droso
>gi|1036720545|ref|XP_017144680.1| PREDICTED: pre-mRNA-splicing factor ATP-dependent RNA helica
>gi|1036720639|ref|XP_017144739.1| PREDICTED: uncharacterized protein LOC108157265 isoform X1
>gi|1036720658|ref|XP_017144748.1| PREDICTED: uncharacterized protein LOC108157265 isoform X2
>gi|1036720676|ref|XP_017144757.1| PREDICTED: uncharacterized protein LOC108157265 isoform X2
>gi|1036720605|ref|XP_017144714.1| PREDICTED: tenascin [Drosophila miranda]
>gi|1036730325|ref|XP_017150131.1| PREDICTED: 39S ribosomal protein L38, mitochondrial [Drosoph
>gi|1036730268|ref|XP_017150097.1| PREDICTED: type I inositol 3,4-bisphosphate 4-phosphatase [
>gi|1036730287|ref|XP_017150106.1| PREDICTED: type I inositol 3,4-bisphosphate 4-phosphatase [
>gi|1036730306|ref|XP_017150119.1| PREDICTED: F-box/LRR-repeat protein 4 [Drosophila miranda]
>gi|1036713343|ref|XP_017140558.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036713360|ref|XP_017140568.1| PREDICTED: neurogenic protein mastermind isoform X3 [Drosoph
>gi|1036713327|ref|XP_017140549.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036713376|ref|XP_017140576.1| PREDICTED: neurogenic protein mastermind isoform X4 [Drosoph
>gi|1036713393|ref|XP_017140588.1| PREDICTED: UMP-CMP kinase 2-like [Drosophila miranda]
>gi|1036713410|ref|XP_017140600.1| PREDICTED: protein Turandot Z [Drosophila miranda]
>gi|1036738073|ref|XP_017154255.1| PREDICTED: ras-related protein Ral-a isoform X2 [Drosophila
>gi|1036738037|ref|XP_017154241.1| PREDICTED: ras-related protein Ral-a isoform X1 [Drosophila
>gi|1036738055|ref|XP_017154249.1| PREDICTED: ras-related protein Ral-a isoform X1 [Drosophila
>gi|1036742862|ref|XP_017155910.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X2
>gi|1036742843|ref|XP_017155909.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036743462|ref|XP_017155945.1| PREDICTED: salivary glue protein Sgs-4-like [Drosophila mira
>gi|1036742339|ref|XP_017155880.1| PREDICTED: poly(ADP-ribose) glycohydrolase [Drosophila miran
>gi|1036742355|ref|XP_017155881.1| PREDICTED: probable insulin-like peptide 7 [Drosophila miran
>gi|1036706078|ref|XP_017136061.1| PREDICTED: uncharacterized protein LOC108151746 [Drosophila
>gi|1036754294|ref|XP_017156549.1| PREDICTED: allatostatin-A receptor [Drosophila miranda]
>gi|1036744015|ref|XP_017155971.1| PREDICTED: nucleolar complex protein 4 homolog B [Drosophila
>gi|1036743998|ref|XP_017155970.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1036750244|ref|XP_017156310.1| PREDICTED: limbic system-associated membrane protein [Drosop
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>gi|1036750279|ref|XP_017156312.1| PREDICTED: uncharacterized protein LOC108164877 isoform X2
>gi|1036750295|ref|XP_017156313.1| PREDICTED: uncharacterized protein LOC108164877 isoform X2
>gi|1036750314|ref|XP_017156314.1| PREDICTED: uncharacterized protein LOC108164877 isoform X3
>gi|1036750263|ref|XP_017156311.1| PREDICTED: uncharacterized protein LOC108164877 isoform X1
>gi|1036739123|ref|XP_017154793.1| PREDICTED: growth arrest-specific protein 8 homolog [Drosop]
>gi|1036739158|ref|XP_017154806.1| PREDICTED: uncharacterized protein LOC108163820 isoform X2
>gi|1036739141|ref|XP_017154799.1| PREDICTED: uncharacterized protein LOC108163820 isoform X1
>gi|1036746952|ref|XP_017156140.1| PREDICTED: neurogenic protein mastermind [Drosophila miranda
>gi|1036746916|ref|XP_017156138.1| PREDICTED: uncharacterized protein LOC108164750 isoform X2
>gi|1036746899|ref|XP_017156136.1| PREDICTED: uncharacterized protein LOC108164750 isoform X1
>gi|1036746933|ref|XP_017156139.1| PREDICTED: voltage-gated potassium channel subunit beta-2-l
>gi|1036756342|ref|XP_017156664.1| PREDICTED: uncharacterized protein LOC108165117 [Drosophila
>gi|1036719606|ref|XP_017144204.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719625|ref|XP_017144213.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719645|ref|XP_017144222.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719586|ref|XP_017144197.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719665|ref|XP_017144230.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719704|ref|XP_017144256.1| PREDICTED: activating signal cointegrator 1 [Drosophila mir
>gi|1036719749|ref|XP_017144265.1| PREDICTED: migration and invasion enhancer 1 [Drosophila mi
>gi|1036719685|ref|XP_017144244.1| PREDICTED: cactin [Drosophila miranda]
>gi|1036742500|ref|XP_017155889.1| PREDICTED: molybdenum cofactor sulfurase isoform X2 [Drosop]
>gi|1036742482|ref|XP_017155888.1| PREDICTED: molybdenum cofactor sulfurase isoform X1 [Drosop]
>gi|1036754003|ref|XP_017156532.1| PREDICTED: small G protein signaling modulator 1 isoform X1
>gi|1036754019|ref|XP_017156533.1| PREDICTED: small G protein signaling modulator 1 isoform X2
>gi|1036752077|ref|XP_017156414.1| PREDICTED: LOW QUALITY PROTEIN: small G protein signaling mo
>gi|1036752096|ref|XP_017156415.1| PREDICTED: glutathione S-transferase theta-1 [Drosophila mi
>gi|1036720782|ref|XP_017144813.1| PREDICTED: uncharacterized protein LOC108157307 isoform X2
>gi|1036720800|ref|XP_017144822.1| PREDICTED: uncharacterized protein LOC108157307 isoform X3
>gi|1036720766|ref|XP_017144804.1| PREDICTED: uncharacterized protein LOC108157307 isoform X1
>gi|1036720817|ref|XP_017144830.1| PREDICTED: uncharacterized protein LOC108157307 isoform X4
>gi|1036725719|ref|XP_017147323.1| PREDICTED: kelch-like protein 22 [Drosophila miranda]
>gi|1036725702|ref|XP_017147311.1| PREDICTED: mitogen-activated protein kinase kinase kinase 7
>gi|1036725736|ref|XP_017147337.1| PREDICTED: uncharacterized protein LOC108158994 [Drosophila
>gi|1036752249|ref|XP_017156424.1| PREDICTED: uncharacterized protein LOC108164946 [Drosophila
>gi|1036757833|ref|XP_017156744.1| PREDICTED: uncharacterized protein LOC108165185 [Drosophila
>gi|1036757852|ref|XP_017156745.1| PREDICTED: uncharacterized protein LOC108165186 [Drosophila
>gi|1036757774|ref|XP_017156741.1| PREDICTED: uncharacterized protein LOC108165182 [Drosophila
>gi|1036704333|ref|XP_017134824.1| PREDICTED: runt-related transcription factor 3 isoform X2 [
>gi|1036704316|ref|XP_017134815.1| PREDICTED: protein lozenge isoform X1 [Drosophila miranda]
>gi|1036745942|ref|XP_017156076.1| PREDICTED: probable cytochrome P450 6v1 isoform X1 [Drosoph
>gi|1036745960|ref|XP_017156077.1| PREDICTED: probable cytochrome P450 6v1 isoform X2 [Drosoph
>gi|1036750556|ref|XP_017156326.1| PREDICTED: segmentation protein Runt isoform X1 [Drosophila
>gi|1036750575|ref|XP_017156327.1| PREDICTED: segmentation protein Runt isoform X2 [Drosophila
>gi|1036756015|ref|XP_017156645.1| PREDICTED: uncharacterized protein LOC108165101 [Drosophila
>gi|1036756034|ref|XP_017156646.1| PREDICTED: uncharacterized protein LOC108165101 [Drosophila
>gi|1036703603|ref|XP_017157044.1| PREDICTED: uncharacterized protein LOC108165494 [Drosophila
>gi|1036739559|ref|XP_017155056.1| PREDICTED: innexin shaking-B isoform X1 [Drosophila miranda]
>gi|1036739595|ref|XP_017155073.1| PREDICTED: innexin shaking-B isoform X3 [Drosophila miranda]
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>gi|1036739577|ref|XP_017155065.1| PREDICTED: innexin shaking-B isoform X2 [Drosophila miranda]
>gi|1036739528|ref|XP_017155040.1| PREDICTED: glycerol-3-phosphate acyltransferase 3 [Drosophil
>gi|1036739543|ref|XP_017155048.1| PREDICTED: glycerol-3-phosphate acyltransferase 3 [Drosophil
>gi|1036739508|ref|XP_017155028.1| PREDICTED: uncharacterized protein LOC108163978 [Drosophila
>gi|1036739612|ref|XP_017155084.1| PREDICTED: protein phosphatase 1 regulatory subunit 11 [Dros
>gi|1036699425|ref|XP_017141085.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036755626|ref|XP_017156623.1| PREDICTED: Niemann-Pick C1 protein [Drosophila miranda]
>gi|1036703095|ref|XP_017156755.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036757682|ref|XP_017156737.1| PREDICTED: uncharacterized protein LOC108165177 [Drosophila
>gi|1036748966|ref|XP_017156239.1| PREDICTED: uncharacterized protein LOC108164827 [Drosophila
>gi|1036748983|ref|XP_017156240.1| PREDICTED: glyoxalase 1 [Drosophila miranda]
>gi|1036730613|ref|XP_017150288.1| PREDICTED: protein Cep78 homolog isoform X3 [Drosophila mira
>gi|1036730594|ref|XP_017150281.1| PREDICTED: protein Cep78 homolog isoform X2 [Drosophila mira
>gi|1036730575|ref|XP_017150272.1| PREDICTED: protein Cep78 homolog isoform X1 [Drosophila mira
>gi|1036730743|ref|XP_017150372.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036730651|ref|XP_017150312.1| PREDICTED: uncharacterized protein LOC108160674 [Drosophila
>gi|1036730632|ref|XP_017150300.1| PREDICTED: uncharacterized protein LOC108160665 [Drosophila
>gi|1036699442|ref|XP_017141213.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036730669|ref|XP_017150326.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036730707|ref|XP_017150348.1| PREDICTED: protein NipSnap [Drosophila miranda]
>gi|1036730726|ref|XP_017150359.1| PREDICTED: protein NipSnap [Drosophila miranda]
>gi|1036730688|ref|XP_017150336.1| PREDICTED: zinc finger matrin-type protein 3 [Drosophila mi
>gi|1036752936|ref|XP_017156468.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1036702288|ref|XP_017156267.1| PREDICTED: UPF0488 protein CG14286 [Drosophila miranda]
>gi|1036721260|ref|XP_017145024.1| PREDICTED: protein Tob1 isoform X2 [Drosophila miranda]
>gi|1036721133|ref|XP_017144968.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721153|ref|XP_017144976.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
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>gi|1036721203|ref|XP_017144998.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721222|ref|XP_017145007.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721241|ref|XP_017145016.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721078|ref|XP_017144940.1| PREDICTED: uncharacterized protein LOC108157404 [Drosophila
>gi|1036721095|ref|XP_017144952.1| PREDICTED: uncharacterized protein LOC108157410 [Drosophila
>gi|1036721114|ref|XP_017144959.1| PREDICTED: uncharacterized protein LOC108157410 [Drosophila
>gi|1036721063|ref|XP_017144928.1| PREDICTED: putative uncharacterized protein DDB_G0291608 [D
>gi|1036757373|ref|XP_017156722.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop
>gi|1036704496|ref|XP_017134931.1| PREDICTED: fork head domain transcription factor slp1, part
>gi|1036747428|ref|XP_017156168.1| PREDICTED: uncharacterized protein LOC108164767 [Drosophila
>gi|1036749002|ref|XP_017156241.1| PREDICTED: protein disconnected [Drosophila miranda]
>gi|1036705986|ref|XP_017135981.1| PREDICTED: uncharacterized protein LOC108151716 isoform X1
>gi|1036706007|ref|XP_017135989.1| PREDICTED: uncharacterized protein LOC108151716 isoform X2
>gi|1036699457|ref|XP_017141336.1| PREDICTED: uncharacterized protein LOC108155149 [Drosophila
>gi|1036732797|ref|XP_017151323.1| PREDICTED: katanin p80 WD40 repeat-containing subunit B1 is
>gi|1036732778|ref|XP_017151317.1| PREDICTED: katanin p80 WD40 repeat-containing subunit B1 is
>gi|1036732815|ref|XP_017151335.1| PREDICTED: peroxisomal multifunctional enzyme type 2 [Droso
>gi|1036732834|ref|XP_017151343.1| PREDICTED: peroxisomal multifunctional enzyme type 2 [Droso
>gi|1036746489|ref|XP_017156111.1| PREDICTED: calpain-C [Drosophila miranda]
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>gi|1036744677|ref|XP_017156003.1| PREDICTED: uncharacterized protein LOC108164661 [Drosophila
>gi|1036744695|ref|XP_017156004.1| PREDICTED: HIG1 domain family member 2A, mitochondrial [Dros
>gi|1036741037|ref|XP_017155803.1| PREDICTED: high mobility group protein DSP1 isoform X1 [Dros
>gi|1036741055|ref|XP_017155804.1| PREDICTED: high mobility group protein DSP1 isoform X2 [Dros
>gi|1036741072|ref|XP_017155805.1| PREDICTED: high mobility group protein DSP1 isoform X3 [Dros
>gi|1036741089|ref|XP_017155807.1| PREDICTED: high mobility group protein DSP1 isoform X4 [Dros
>gi|1036717210|ref|XP_017142880.1| PREDICTED: acyl-CoA Delta(11) desaturase-like [Drosophila mi
>gi|1036717122|ref|XP_017142829.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036717194|ref|XP_017142868.1| PREDICTED: splicing factor U2AF 50 kDa subunit [Drosophila m
>gi|1036717138|ref|XP_017142840.1| PREDICTED: uncharacterized protein LOC108156065 [Drosophila
>gi|1036717154|ref|XP_017142848.1| PREDICTED: uncharacterized protein LOC108156065 [Drosophila
>gi|1036717171|ref|XP_017142856.1| PREDICTED: uncharacterized protein LOC108156065 [Drosophila
>gi|1036729166|ref|XP_017149407.1| PREDICTED: uncharacterized protein LOC108160122 [Drosophila
>gi|1036729148|ref|XP_017149396.1| PREDICTED: protein no-on-transient A [Drosophila miranda]
>gi|1036723090|ref|XP_017145799.1| PREDICTED: furin-like protease 2 isoform X1 [Drosophila mira
>gi|1036723112|ref|XP_017145808.1| PREDICTED: furin-like protease 2 isoform X1 [Drosophila mira
>gi|1036723130|ref|XP_017145820.1| PREDICTED: furin-like protease 2 isoform X2 [Drosophila mira
>gi|1036723148|ref|XP_017145831.1| PREDICTED: furin-like protease 2 isoform X3 [Drosophila mira
>gi|1036723167|ref|XP_017145840.1| PREDICTED: furin-like protease 2 isoform X4 [Drosophila mira
>gi|1036723203|ref|XP_017145861.1| PREDICTED: uncharacterized protein LOC108158142 isoform X2
>gi|1036723186|ref|XP_017145853.1| PREDICTED: uncharacterized protein LOC108158142 isoform X1
>gi|1036723214|ref|XP_017145869.1| PREDICTED: uncharacterized protein LOC108158142 isoform X3
>gi|1036736680|ref|XP_017153482.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736698|ref|XP_017153491.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736714|ref|XP_017153499.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736733|ref|XP_017153508.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736663|ref|XP_017153471.1| PREDICTED: negative elongation factor D [Drosophila miranda]
>gi|1036756892|ref|XP_017156698.1| PREDICTED: probable phospholipid-transporting ATPase IF [Dr
>gi|1036751992|ref|XP_017156409.1| PREDICTED: probable phospholipid-transporting ATPase IF [Dr
>gi|1036747113|ref|XP_017156149.1| PREDICTED: TNF receptor-associated factor 3 isoform X1 [Dros
>gi|1036747131|ref|XP_017156151.1| PREDICTED: TNF receptor-associated factor 3 isoform X2 [Dros
>gi|1036747097|ref|XP_017156148.1| PREDICTED: annexin B11 isoform X2 [Drosophila miranda]
>gi|1036747062|ref|XP_017156146.1| PREDICTED: annexin B11 isoform X1 [Drosophila miranda]
>gi|1036747078|ref|XP_017156147.1| PREDICTED: annexin B11 isoform X1 [Drosophila miranda]
>gi|1036733599|ref|XP_017151723.1| PREDICTED: gamma-glutamyltranspeptidase 1 [Drosophila mirand
>gi|1036733650|ref|XP_017151741.1| PREDICTED: gamma-glutamyltranspeptidase 1 [Drosophila mirand
>gi|1036733690|ref|XP_017151759.1| PREDICTED: uncharacterized protein LOC108161868 isoform X2
>gi|1036733669|ref|XP_017151752.1| PREDICTED: uncharacterized protein LOC108161868 isoform X1
>gi|1036733709|ref|XP_017151771.1| PREDICTED: uncharacterized protein LOC108161884 [Drosophila
>gi|1036733561|ref|XP_017151703.1| PREDICTED: sushi, von Willebrand factor type A, EGF and pent
>gi|1036733581|ref|XP_017151710.1| PREDICTED: sushi, von Willebrand factor type A, EGF and pent
>gi|1036756308|ref|XP_017156662.1| PREDICTED: fibrous sheath CABYR-binding protein [Drosophila
>gi|1036746809|ref|XP_017156131.1| PREDICTED: serine protease gd [Drosophila miranda]
>gi|1036746826|ref|XP_017156132.1| PREDICTED: protein twisted gastrulation [Drosophila miranda]
>gi|1036755307|ref|XP_017156608.1| PREDICTED: uncharacterized protein LOC108165071 [Drosophila
>gi|1036755325|ref|XP_017156610.1| PREDICTED: uncharacterized protein LOC108165071 [Drosophila
>gi|1036717982|ref|XP_017143303.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718002|ref|XP_017143313.1| PREDICTED: voltage-dependent calcium channel type A subunit
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>gi|1036718021|ref|XP_017143323.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718059|ref|XP_017143341.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718040|ref|XP_017143331.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718076|ref|XP_017143349.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718113|ref|XP_017143364.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718095|ref|XP_017143356.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718128|ref|XP_017143372.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718147|ref|XP_017143378.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718166|ref|XP_017143387.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718188|ref|XP_017143393.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718207|ref|XP_017143400.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036754527|ref|XP_017156563.1| PREDICTED: PDZ domain-containing protein 8 [Drosophila miran
>gi|1036756053|ref|XP_017156647.1| PREDICTED: leucine-rich repeat-containing protein 20 isofor
>gi|1036756072|ref|XP_017156648.1| PREDICTED: leucine-rich repeat-containing protein 20 isofor
>gi|1036703879|ref|XP_017134567.1| PREDICTED: leucine-rich repeat-containing protein 69-like i
>gi|1036703930|ref|XP_017134588.1| PREDICTED: protein lap1-like isoform X4 [Drosophila miranda]
>gi|1036703913|ref|XP_017134582.1| PREDICTED: leucine-rich repeat-containing protein 69-like i
>gi|1036703896|ref|XP_017134574.1| PREDICTED: leucine-rich repeat-containing protein 69-like i
>gi|1036748107|ref|XP_017156204.1| PREDICTED: uncharacterized protein LOC108164792 [Drosophila
>gi|1036748088|ref|XP_017156203.1| PREDICTED: uncharacterized protein LOC108164791 [Drosophila
>gi|1036754275|ref|XP_017156548.1| PREDICTED: probable cytochrome P450 28a5 [Drosophila mirand
>gi|1036732315|ref|XP_017151082.1| PREDICTED: protein xmas-2 [Drosophila miranda]
>gi|1036732332|ref|XP_017151094.1| PREDICTED: uncharacterized protein LOC108161359 [Drosophila
>gi|1036699471|ref|XP_017141454.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop
>gi|1036729976|ref|XP_017149886.1| PREDICTED: uncharacterized protein LOC108160402 [Drosophila
>gi|1036699488|ref|XP_017141582.1| PREDICTED: uncharacterized protein LOC108155322 [Drosophila
>gi|1036740746|ref|XP_017155712.1| PREDICTED: histone H1.3-like [Drosophila miranda]
>gi|1036740775|ref|XP_017155735.1| PREDICTED: histone H2B-like [Drosophila miranda]
>gi|1036740760|ref|XP_017155723.1| PREDICTED: coiled-coil-helix-coiled-coil-helix domain-conta
>gi|1036743427|ref|XP_017155943.1| PREDICTED: hexokinase type 2 [Drosophila miranda]
>gi|1036699502|ref|XP_017141705.1| PREDICTED: LOW QUALITY PROTEIN: protein vreteno-like [Drosop
>gi|1036727344|ref|XP_017148347.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036727361|ref|XP_017148355.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036727295|ref|XP_017148325.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036727325|ref|XP_017148337.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036723885|ref|XP_017146206.1| PREDICTED: probable cytochrome P450 318a1 [Drosophila miran
>gi|1036723848|ref|XP_017146187.1| PREDICTED: protein strawberry notch isoform X1 [Drosophila
>gi|1036723867|ref|XP_017146196.1| PREDICTED: protein strawberry notch isoform X2 [Drosophila
>gi|1036723919|ref|XP_017146228.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036723902|ref|XP_017146217.1| PREDICTED: 2-methoxy-6-polyprenyl-1,4-benzoquinol methylase
>gi|1036699514|ref|XP_017141828.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036723936|ref|XP_017146241.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036727144|ref|XP_017148209.1| PREDICTED: relaxin receptor 2 [Drosophila miranda]
>gi|1036727215|ref|XP_017148264.1| PREDICTED: uncharacterized protein LOC108159462 [Drosophila
>gi|1036727198|ref|XP_017148250.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 5 [
>gi|1036727181|ref|XP_017148235.1| PREDICTED: brahma-associated protein of 60 kDa [Drosophila
>gi|1036727162|ref|XP_017148223.1| PREDICTED: cleft lip and palate transmembrane protein 1-like
>gi|1036756184|ref|XP_017156655.1| PREDICTED: uncharacterized protein LOC108165108 [Drosophila
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>gi|1036745049|ref|XP_017156024.1| PREDICTED: voltage-dependent T-type calcium channel subunit
>gi|1036745032|ref|XP_017156022.1| PREDICTED: voltage-dependent T-type calcium channel subunit
>gi|1036754602|ref|XP_017156567.1| PREDICTED: vanin-like protein 3 [Drosophila miranda]
>gi|1036754584|ref|XP_017156566.1| PREDICTED: vanin-like protein 1 [Drosophila miranda]
>gi|1036757265|ref|XP_017156715.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036756797|ref|XP_017156691.1| PREDICTED: vanin-like protein 2 [Drosophila miranda]
>gi|1036740626|ref|XP_017155631.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740645|ref|XP_017155638.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740661|ref|XP_017155648.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740677|ref|XP_017155656.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740693|ref|XP_017155669.1| PREDICTED: myosin regulatory light chain sqh [Drosophila mi
>gi|1036727653|ref|XP_017148524.1| PREDICTED: transmembrane protein 132B [Drosophila miranda]
>gi|1036703295|ref|XP_017156876.1| PREDICTED: uncharacterized protein LOC108165351 isoform X1
>gi|1036703334|ref|XP_017156891.1| PREDICTED: uncharacterized protein LOC108165351 isoform X3
>gi|1036703369|ref|XP_017156906.1| PREDICTED: uncharacterized protein LOC108165351 isoform X5
>gi|1036703315|ref|XP_017156883.1| PREDICTED: uncharacterized protein LOC108165351 isoform X2
>gi|1036703354|ref|XP_017156899.1| PREDICTED: uncharacterized protein LOC108165351 isoform X4
>gi|1036729305|ref|XP_017149482.1| PREDICTED: dmX-like protein 2 [Drosophila miranda]
>gi|1036751755|ref|XP_017156394.1| PREDICTED: splicing factor 3B subunit 4 isoform X2 [Drosophi
>gi|1036751736|ref|XP_017156393.1| PREDICTED: splicing factor 3B subunit 4 isoform X1 [Drosophi
>gi|1036746971|ref|XP_017156141.1| PREDICTED: ceramide synthase 6 [Drosophila miranda]
>gi|1036746990|ref|XP_017156142.1| PREDICTED: josephin-like protein [Drosophila miranda]
>gi|1036724726|ref|XP_017146751.1| PREDICTED: transcription elongation factor SPT6 [Drosophila
>gi|1036747479|ref|XP_017156171.1| PREDICTED: uncharacterized protein LOC108164770 [Drosophila
>gi|1036747496|ref|XP_017156172.1| PREDICTED: uncharacterized protein LOC108164770 [Drosophila
>gi|1036747534|ref|XP_017156174.1| PREDICTED: cytochrome b5 isoform X2 [Drosophila miranda]
>gi|1036747515|ref|XP_017156173.1| PREDICTED: cytochrome b5 isoform X1 [Drosophila miranda]
>gi|1036747555|ref|XP_017156175.1| PREDICTED: probable cytochrome b5 isoform X3 [Drosophila mi
>gi|1036710068|ref|XP_017138615.1| PREDICTED: uncharacterized protein LOC108153261 [Drosophila
>gi|1036710166|ref|XP_017138682.1| PREDICTED: uncharacterized protein LOC108153299 [Drosophila
>gi|1036710183|ref|XP_017138692.1| PREDICTED: uncharacterized protein LOC108153299 [Drosophila
>gi|1036710113|ref|XP_017138650.1| PREDICTED: U4/U6.U5 tri-snRNP-associated protein 2 isoform
>gi|1036710130|ref|XP_017138660.1| PREDICTED: U4/U6.U5 tri-snRNP-associated protein 2 isoform
>gi|1036710098|ref|XP_017138642.1| PREDICTED: U4/U6.U5 tri-snRNP-associated protein 2 isoform
>gi|1036710215|ref|XP_017138717.1| PREDICTED: L-xylulose reductase [Drosophila miranda]
>gi|1036710083|ref|XP_017138628.1| PREDICTED: mucin-12 [Drosophila miranda]
>gi|1036710147|ref|XP_017138672.1| PREDICTED: probable sulfite oxidase, mitochondrial [Drosophi
>gi|1036710200|ref|XP_017138706.1| PREDICTED: proteasome subunit alpha type-7-1 [Drosophila mi
>gi|1036749844|ref|XP_017156288.1| PREDICTED: XK-related protein 6 isoform X2 [Drosophila mira
>gi|1036749826|ref|XP_017156287.1| PREDICTED: XK-related protein 6 isoform X1 [Drosophila mira
>gi|1036749375|ref|XP_017156263.1| PREDICTED: 27 kDa hemolymph protein [Drosophila miranda]
>gi|1036726872|ref|XP_017148064.1| PREDICTED: uncharacterized protein LOC108159354 [Drosophila
>gi|1036726909|ref|XP_017148080.1| PREDICTED: lambda-crystallin homolog [Drosophila miranda]
>gi|1036726852|ref|XP_017148050.1| PREDICTED: protein IWS1 homolog [Drosophila miranda]
>gi|1036749770|ref|XP_017156284.1| PREDICTED: WASH complex subunit 7 homolog isoform X1 [Drosop
>gi|1036749789|ref|XP_017156285.1| PREDICTED: WASH complex subunit 7 homolog isoform X2 [Drosop
>gi|1036745580|ref|XP_017156056.1| PREDICTED: peptidyl-prolyl cis-trans isomerase [Drosophila
>gi|1036738801|ref|XP_017154611.1| PREDICTED: myotubularin-related protein 4 isoform X1 [Drosop
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>gi|1036738817|ref|XP_017154619.1| PREDICTED: myotubularin-related protein 4 isoform X2 [Droso
>gi|1036738832|ref|XP_017154628.1| PREDICTED: myotubularin-related protein 4 isoform X2 [Droso
>gi|1036738849|ref|XP_017154637.1| PREDICTED: myotubularin-related protein 4 isoform X2 [Droso
>gi|1036738865|ref|XP_017154644.1| PREDICTED: myotubularin-related protein 4 isoform X3 [Droso
>gi|1036747408|ref|XP_017156166.1| PREDICTED: endoplasmic reticulum resident protein 44 [Droso
>gi|1036744291|ref|XP_017155983.1| PREDICTED: RNA-binding protein cabeza isoform X1 [Drosophila
>gi|1036744308|ref|XP_017155984.1| PREDICTED: RNA-binding protein cabeza isoform X2 [Drosophila
>gi|1036749023|ref|XP_017156242.1| PREDICTED: vesicle transport protein GOT1B [Drosophila mirar
>gi|1036749041|ref|XP_017156243.1| PREDICTED: cytochrome b-c1 complex subunit 7 [Drosophila mi
>gi|1036747007|ref|XP_017156143.1| PREDICTED: ethanolamine kinase [Drosophila miranda]
>gi|1036747025|ref|XP_017156144.1| PREDICTED: ethanolamine kinase [Drosophila miranda]
>gi|1036747044|ref|XP_017156145.1| PREDICTED: ethanolamine kinase [Drosophila miranda]
>gi|1036699531|ref|XP_017141956.1| PREDICTED: uncharacterized protein LOC108155546 [Drosophila
>gi|1036729267|ref|XP_017149466.1| PREDICTED: vascular endothelial growth factor C [Drosophila
>gi|1036729240|ref|XP_017149454.1| PREDICTED: zinc finger protein 596 [Drosophila miranda]
>gi|1036729203|ref|XP_017149433.1| PREDICTED: protein FAM188A homolog isoform X1 [Drosophila m
>gi|1036729222|ref|XP_017149441.1| PREDICTED: protein FAM188A homolog isoform X2 [Drosophila m
>gi|1036740872|ref|XP_017155796.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D
>gi|1036740890|ref|XP_017155797.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D
>gi|1036756671|ref|XP_017156684.1| PREDICTED: succinate dehydrogenase [ubiquinone] iron-sulfur
>gi|1036753001|ref|XP_017156472.1| PREDICTED: dual specificity protein phosphatase 3 isoform X
>gi|1036753018|ref|XP_017156473.1| PREDICTED: dual specificity protein phosphatase 3 isoform X
>gi|1036753037|ref|XP_017156474.1| PREDICTED: dual specificity protein phosphatase 3 isoform X
>gi|1036753054|ref|XP_017156475.1| PREDICTED: dual specificity protein phosphatase 3 isoform X
>gi|1036720526|ref|XP_017144670.1| PREDICTED: putative uncharacterized protein DDB_G0277255 [D
>gi|1036706051|ref|XP_017136033.1| PREDICTED: uncharacterized protein LOC108151737 [Drosophila
>gi|1036755195|ref|XP_017156602.1| PREDICTED: uncharacterized protein LOC108165065 [Drosophila
>gi|1036705477|ref|XP_017135628.1| PREDICTED: uncharacterized protein LOC108151496 [Drosophila
>gi|1036705493|ref|XP_017135637.1| PREDICTED: uncharacterized protein LOC108151496 [Drosophila
>gi|1036699547|ref|XP_017142209.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036754112|ref|XP_017156538.1| PREDICTED: uncharacterized protein LOC108165013 [Drosophila
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>gi|1036754150|ref|XP_017156541.1| PREDICTED: uncharacterized protein LOC108165014 [Drosophila
>gi|1036703695|ref|XP_017157096.1| PREDICTED: myotrophin [Drosophila miranda]
>gi|1036703654|ref|XP_017157074.1| PREDICTED: uncharacterized protein LOC108165523 isoform X1
>gi|1036703678|ref|XP_017157084.1| PREDICTED: uncharacterized protein LOC108165523 isoform X2
>gi|1036706280|ref|XP_017136196.1| PREDICTED: cAMP-regulated phosphoprotein 19-A-like [Drosoph
>gi|1036704972|ref|XP_017135276.1| PREDICTED: nuclear pore complex protein Nup50-like [Drosoph
>gi|1036703843|ref|XP_017134547.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036703862|ref|XP_017134555.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036732909|ref|XP_017151377.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
>gi|1036732928|ref|XP_017151386.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
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>gi|1036733006|ref|XP_017151403.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
>gi|1036732890|ref|XP_017151370.1| PREDICTED: uncharacterized protein LOC108161591 isoform X1
>gi|1036733044|ref|XP_017151421.1| PREDICTED: vacuolar ATPase assembly integral membrane prote
>gi|1036733025|ref|XP_017151411.1| PREDICTED: innexin inx5 [Drosophila miranda]
>gi|1036704047|ref|XP_017134656.1| PREDICTED: uncharacterized protein LOC108150861 [Drosophila
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>gi|1036704109|ref|XP_017134691.1| PREDICTED: uncharacterized protein LOC108150889 [Drosophila
>gi|1036735457|ref|XP_017152788.1| PREDICTED: uncharacterized protein LOC108162511 isoform X2
>gi|1036735440|ref|XP_017152781.1| PREDICTED: uncharacterized protein LOC108162511 isoform X1
>gi|1036725417|ref|XP_017147153.1| PREDICTED: uncharacterized protein LOC108158886 isoform X1
>gi|1036725435|ref|XP_017147164.1| PREDICTED: uncharacterized protein LOC108158886 isoform X1
>gi|1036725472|ref|XP_017147180.1| PREDICTED: uncharacterized protein LOC108158886 isoform X3
>gi|1036725491|ref|XP_017147189.1| PREDICTED: uncharacterized protein LOC108158886 isoform X4
>gi|1036725510|ref|XP_017147197.1| PREDICTED: uncharacterized protein LOC108158886 isoform X5
>gi|1036725452|ref|XP_017147172.1| PREDICTED: uncharacterized protein LOC108158886 isoform X2
>gi|1036725528|ref|XP_017147210.1| PREDICTED: lectin subunit alpha [Drosophila miranda]
>gi|1036739293|ref|XP_017154907.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1036739309|ref|XP_017154916.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1036739323|ref|XP_017154924.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1036739278|ref|XP_017154896.1| PREDICTED: DNA replication licensing factor Mcm3 [Drosophila
>gi|1036743982|ref|XP_017155969.1| PREDICTED: protein DENND6B [Drosophila miranda]
>gi|1036741869|ref|XP_017155853.1| PREDICTED: DNA repair protein XRCC1 [Drosophila miranda]
>gi|1036741887|ref|XP_017155854.1| PREDICTED: calcineurin subunit B type 1 [Drosophila miranda]
>gi|1036741905|ref|XP_017155855.1| PREDICTED: calcineurin subunit B type 1 [Drosophila miranda]
>gi|1036741924|ref|XP_017155856.1| PREDICTED: calcineurin subunit B type 1 [Drosophila miranda]
>gi|1036742951|ref|XP_017155915.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036742969|ref|XP_017155917.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036742988|ref|XP_017155918.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743007|ref|XP_017155919.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743026|ref|XP_017155920.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743045|ref|XP_017155921.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743064|ref|XP_017155922.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743079|ref|XP_017155923.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743098|ref|XP_017155924.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743117|ref|XP_017155925.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036754489|ref|XP_017156560.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036723016|ref|XP_017145754.1| PREDICTED: A-kinase anchor protein 1, mitochondrial [Drosoph
>gi|1036723059|ref|XP_017145780.1| PREDICTED: 60S ribosomal protein L36 [Drosophila miranda]
>gi|1036723075|ref|XP_017145788.1| PREDICTED: 60S ribosomal protein L36 [Drosophila miranda]
>gi|1036723040|ref|XP_017145767.1| PREDICTED: histone H4 transcription factor [Drosophila miran
>gi|1036733112|ref|XP_017151453.1| PREDICTED: uncharacterized protein LOC108161665 [Drosophila
>gi|1036733131|ref|XP_017151467.1| PREDICTED: uncharacterized protein LOC108161671 [Drosophila
>gi|1036733095|ref|XP_017151447.1| PREDICTED: frizzled-4 [Drosophila miranda]
>gi|1036741962|ref|XP_017155858.1| PREDICTED: uncharacterized protein LOC108164555 isoform X1
>gi|1036741998|ref|XP_017155860.1| PREDICTED: uncharacterized protein LOC108164555 isoform X3
>gi|1036741981|ref|XP_017155859.1| PREDICTED: uncharacterized protein LOC108164555 isoform X2
>gi|1036757097|ref|XP_017156705.1| PREDICTED: uncharacterized protein LOC108165149 [Drosophila
>gi|1036703496|ref|XP_017156979.1| PREDICTED: thioredoxin domain-containing protein 9-like [Dro
>gi|1036731843|ref|XP_017150862.1| PREDICTED: zinc finger CCCH domain-containing protein 18 iso
>gi|1036731825|ref|XP_017150856.1| PREDICTED: zinc finger CCCH domain-containing protein 18 iso
>gi|1036731807|ref|XP_017150848.1| PREDICTED: zinc finger CCCH domain-containing protein 18 iso
>gi|1036753574|ref|XP_017156506.1| PREDICTED: protein ABHD11 [Drosophila miranda]
>gi|1036747291|ref|XP_017156161.1| PREDICTED: protein unc-119 homolog [Drosophila miranda]
>gi|1036743512|ref|XP_017155946.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
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>gi|1036744830|ref|XP_017156010.1| PREDICTED: autophagy protein 5 [Drosophila miranda]
>gi|1036736920|ref|XP_017153624.1| PREDICTED: rho GTPase-activating protein gacZ [Drosophila m
>gi|1036752137|ref|XP_017156417.1| PREDICTED: acyl-CoA synthetase family member 3, mitochondria
>gi|1036752155|ref|XP_017156418.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036742807|ref|XP_017155906.1| PREDICTED: sarcoplasmic calcium-binding protein [Drosophila
>gi|1036742826|ref|XP_017155908.1| PREDICTED: uncharacterized protein LOC108164593 [Drosophila
>gi|1036705562|ref|XP_017135680.1| PREDICTED: uncharacterized protein LOC108151534 isoform X1
>gi|1036705592|ref|XP_017135689.1| PREDICTED: uncharacterized protein LOC108151534 isoform X2
>gi|1036705241|ref|XP_017135477.1| PREDICTED: uncharacterized protein LOC108151397 [Drosophila
>gi|1036705258|ref|XP_017135483.1| PREDICTED: uncharacterized protein LOC108151397 [Drosophila
>gi|1036703620|ref|XP_017157054.1| PREDICTED: uncharacterized protein LOC108165505 [Drosophila
>gi|1036703637|ref|XP_017157061.1| PREDICTED: uncharacterized protein LOC108165505 [Drosophila
>gi|1036753884|ref|XP_017156525.1| PREDICTED: condensin complex subunit 2-like isoform X1 [Dros
>gi|1036753900|ref|XP_017156526.1| PREDICTED: condensin complex subunit 2-like isoform X2 [Dros
>gi|1036753916|ref|XP_017156527.1| PREDICTED: uncharacterized protein LOC108165003 isoform X3
>gi|1036705646|ref|XP_017135726.1| PREDICTED: uncharacterized protein LOC108151562 [Drosophila
>gi|1036706387|ref|XP_017136266.1| PREDICTED: uncharacterized protein LOC108151877 [Drosophila
>gi|1036705526|ref|XP_017135657.1| PREDICTED: uncharacterized protein LOC108151511 isoform X2
>gi|1036705509|ref|XP_017135649.1| PREDICTED: uncharacterized protein LOC108151511 isoform X1
>gi|1036757737|ref|XP_017156739.1| PREDICTED: uncharacterized protein LOC108165180 [Drosophila
>gi|1036705392|ref|XP_017135575.1| PREDICTED: uncharacterized protein LOC108151461 [Drosophila
>gi|1036706547|ref|XP_017136368.1| PREDICTED: uncharacterized protein LOC108151944 [Drosophila
>gi|1036707574|ref|XP_017137057.1| PREDICTED: uncharacterized protein LOC108152322 [Drosophila
>gi|1036705003|ref|XP_017135302.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1036699562|ref|XP_017142462.1| PREDICTED: F-box/WD repeat-containing protein A [Drosophila
>gi|1036707369|ref|XP_017136915.1| PREDICTED: uncharacterized protein LOC108152233 [Drosophila
>gi|1036707163|ref|XP_017136794.1| PREDICTED: small nuclear ribonucleoprotein-associated prote
>gi|1036707420|ref|XP_017136949.1| PREDICTED: uncharacterized protein LOC108152250 [Drosophila
>gi|1036699577|ref|XP_017142588.1| PREDICTED: uncharacterized protein LOC108155912 [Drosophila
>gi|1036753311|ref|XP_017156491.1| PREDICTED: transcription factor GTE3, chloroplastic-like [D
>gi|1036753369|ref|XP_017156494.1| PREDICTED: uncharacterized protein LOC108164984 [Drosophila
>gi|1036753351|ref|XP_017156493.1| PREDICTED: uncharacterized protein LOC108164983 [Drosophila
>gi|1036753332|ref|XP_017156492.1| PREDICTED: uncharacterized protein LOC108164981 [Drosophila
>gi|1036699591|ref|XP_017142714.1| PREDICTED: uncharacterized protein LOC108155989 [Drosophila
>gi|1036756128|ref|XP_017156651.1| PREDICTED: uncharacterized protein LOC108165105 [Drosophila
>gi|1036705800|ref|XP_017135843.1| PREDICTED: uncharacterized protein LOC108151644 [Drosophila
>gi|1036705815|ref|XP_017135854.1| PREDICTED: uncharacterized protein LOC108151644 [Drosophila
>gi|1036735122|ref|XP_017152591.1| PREDICTED: homeobox protein cut isoform X1 [Drosophila miran
>gi|1036735140|ref|XP_017152600.1| PREDICTED: homeobox protein cut isoform X2 [Drosophila miran
>gi|1036735155|ref|XP_017152612.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein pho
>gi|1036757227|ref|XP_017156712.1| PREDICTED: uncharacterized protein LOC108165157 [Drosophila
>gi|1036730870|ref|XP_017150446.1| PREDICTED: uncharacterized protein LOC108160754 [Drosophila
>gi|1036730889|ref|XP_017150453.1| PREDICTED: uncharacterized protein LOC108160754 [Drosophila
>gi|1036706590|ref|XP_017136397.1| PREDICTED: uncharacterized protein LOC108151962 [Drosophila
>gi|1036704207|ref|XP_017134751.1| PREDICTED: LOW QUALITY PROTEIN: AF4/FMR2 family member 4 [D
>gi|1036715798|ref|XP_017142050.1| PREDICTED: protein HIRA homolog [Drosophila miranda]
>gi|1036715815|ref|XP_017142063.1| PREDICTED: negative elongation factor B [Drosophila miranda]
>gi|1036715831|ref|XP_017142077.1| PREDICTED: uncharacterized protein LOC108155639 [Drosophila
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>gi|1036715851|ref|XP_017142089.1| PREDICTED: pyruvate dehydrogenase [acetyl-transferring]-phos
>gi|1036715871|ref|XP_017142102.1| PREDICTED: ras-related protein Rab-39B [Drosophila miranda]
>gi|1036742374|ref|XP_017155882.1| PREDICTED: mitochondrial import receptor subunit TOM40 homol
>gi|1036758069|ref|XP_017156754.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036699606|ref|XP_017142847.1| PREDICTED: uncharacterized protein LOC108156047 [Drosophila
>gi|1036706649|ref|XP_017136459.1| PREDICTED: uncharacterized protein LOC108151998 [Drosophila
>gi|1036699621|ref|XP_017142968.1| PREDICTED: uncharacterized protein LOC108156128 [Drosophila
>gi|1036753796|ref|XP_017156519.1| PREDICTED: venom carboxylesterase-6 [Drosophila miranda]
>gi|1036749861|ref|XP_017156289.1| PREDICTED: L-asparaginase [Drosophila miranda]
>gi|1036737180|ref|XP_017153765.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036737128|ref|XP_017153742.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036737146|ref|XP_017153748.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036737164|ref|XP_017153757.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036725887|ref|XP_017147442.1| PREDICTED: uncharacterized protein LOC108159029 [Drosophila
>gi|1036725905|ref|XP_017147454.1| PREDICTED: CAS1 domain-containing protein 1 [Drosophila mira
>gi|1036699638|ref|XP_017143087.1| PREDICTED: uncharacterized protein LOC108156219 [Drosophila
>gi|1036757946|ref|XP_017156751.1| PREDICTED: uncharacterized protein ZK546.14 [Drosophila mira
>gi|1036757963|ref|XP_017156752.1| PREDICTED: uncharacterized protein ZK546.14 [Drosophila mira
>gi|1036745490|ref|XP_017156050.1| PREDICTED: uncharacterized protein LOC108164691 [Drosophila
>gi|1036745474|ref|XP_017156049.1| PREDICTED: V-type proton ATPase subunit d 1 [Drosophila mira
>gi|1036734365|ref|XP_017152153.1| PREDICTED: fizzy-related protein homolog [Drosophila miranda
>gi|1036734383|ref|XP_017152162.1| PREDICTED: fizzy-related protein homolog [Drosophila miranda
>gi|1036751528|ref|XP_017156383.1| PREDICTED: uncharacterized protein LOC108164911 isoform X1
>gi|1036751546|ref|XP_017156384.1| PREDICTED: acidic phospholipase A2 PA4 isoform X2 [Drosophil
>gi|1036751567|ref|XP_017156385.1| PREDICTED: uncharacterized protein LOC108164911 isoform X3
>gi|1036751588|ref|XP_017156386.1| PREDICTED: uncharacterized protein LOC108164911 isoform X3
>gi|1036734545|ref|XP_017152252.1| PREDICTED: nuclear cap-binding protein subunit 1 [Drosophila
>gi|1036734562|ref|XP_017152263.1| PREDICTED: mastermind-like domain-containing protein 1 [Dros
>gi|1036746577|ref|XP_017156116.1| PREDICTED: torsin-like protein [Drosophila miranda]
>gi|1036746594|ref|XP_017156118.1| PREDICTED: 39S ribosomal protein L30, mitochondrial [Drosoph
>gi|1036706790|ref|XP_017136564.1| PREDICTED: helix-loop-helix protein 1 [Drosophila miranda]
>gi|1036705672|ref|XP_017135748.1| PREDICTED: protein Flattop homolog [Drosophila miranda]
>gi|1036708216|ref|XP_017137496.1| PREDICTED: uncharacterized protein LOC108152568 [Drosophila
>gi|1036747833|ref|XP_017156192.1| PREDICTED: ras-interacting protein RIP3 [Drosophila miranda]
>gi|1036739472|ref|XP_017155007.1| PREDICTED: COPII coat assembly protein SEC16 isoform X1 [Dro
>gi|1036739490|ref|XP_017155016.1| PREDICTED: uncharacterized protein LOC108163962 isoform X2
>gi|1036721467|ref|XP_017145136.1| PREDICTED: uncharacterized abhydrolase domain-containing pro
>gi|1036721430|ref|XP_017145114.1| PREDICTED: serine/threonine-protein kinase minibrain isoform
>gi|1036721411|ref|XP_017145106.1| PREDICTED: serine/threonine-protein kinase minibrain isoform
>gi|1036721448|ref|XP_017145124.1| PREDICTED: zinc finger protein 622 [Drosophila miranda]
>gi|1036742270|ref|XP_017155875.1| PREDICTED: actin-related protein 8 [Drosophila miranda]
>gi|1036742285|ref|XP_017155876.1| PREDICTED: putative sulfiredoxin isoform X1 [Drosophila mira
>gi|1036742303|ref|XP_017155877.1| PREDICTED: putative sulfiredoxin isoform X2 [Drosophila mira
>gi|1036713586|ref|XP_017140747.1| PREDICTED: laccase-3 [Drosophila miranda]
>gi|1036713719|ref|XP_017140839.1| PREDICTED: uncharacterized protein LOC108154897 [Drosophila
>gi|1036713619|ref|XP_017140768.1| PREDICTED: uncharacterized protein LOC108154855 [Drosophila
>gi|1036713671|ref|XP_017140803.1| PREDICTED: RING finger protein 121 [Drosophila miranda]
>gi|1036713636|ref|XP_017140781.1| PREDICTED: mitochondrial uncoupling protein 4 [Drosophila m
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>gi|1036713655|ref|XP_017140791.1| PREDICTED: mitochondrial uncoupling protein 4 [Drosophila m
>gi|1036713603|ref|XP_017140755.1| PREDICTED: replication factor C subunit 4 [Drosophila miran
>gi|1036713704|ref|XP_017140827.1| PREDICTED: transcription initiation factor TFIID subunit 9
>gi|1036713688|ref|XP_017140815.1| PREDICTED: uncharacterized protein LOC108154880 [Drosophila
>gi|1036739732|ref|XP_017155160.1| PREDICTED: uncharacterized protein LOC108164082 [Drosophila
>gi|1036705782|ref|XP_017135831.1| PREDICTED: uncharacterized protein LOC108151634 [Drosophila
>gi|1036754620|ref|XP_017156568.1| PREDICTED: homeobox protein unc-4 isoform X1 [Drosophila mi
>gi|1036754637|ref|XP_017156569.1| PREDICTED: homeobox protein unc-4 isoform X2 [Drosophila mi
>gi|1036704299|ref|XP_017134806.1| PREDICTED: homeobox protein unc-4 [Drosophila miranda]
>gi|1036706421|ref|XP_017136290.1| PREDICTED: uncharacterized protein LOC108151894 [Drosophila
>gi|1036709987|ref|XP_017138564.1| PREDICTED: probable Ras GTPase-activating protein isoform X4
>gi|1036709970|ref|XP_017138558.1| PREDICTED: probable Ras GTPase-activating protein isoform X3
>gi|1036709954|ref|XP_017138549.1| PREDICTED: probable Ras GTPase-activating protein isoform X2
>gi|1036709938|ref|XP_017138541.1| PREDICTED: probable Ras GTPase-activating protein isoform X1
>gi|1036710004|ref|XP_017138577.1| PREDICTED: DNA topoisomerase 1 isoform X1 [Drosophila mirano
>gi|1036710021|ref|XP_017138586.1| PREDICTED: DNA topoisomerase 1 isoform X2 [Drosophila mirano
>gi|1036710036|ref|XP_017138596.1| PREDICTED: DNA topoisomerase 1 isoform X3 [Drosophila mirano
>gi|1036710053|ref|XP_017138606.1| PREDICTED: DNA topoisomerase 1 isoform X4 [Drosophila mirano
>gi|1036739696|ref|XP_017155141.1| PREDICTED: dystrotelin [Drosophila miranda]
>gi|1036739714|ref|XP_017155149.1| PREDICTED: uncharacterized protein LOC108164074 [Drosophila
>gi|1036724829|ref|XP_017146830.1| PREDICTED: histone deacetylase 6 isoform X1 [Drosophila mira
>gi|1036724896|ref|XP_017146861.1| PREDICTED: histone deacetylase 6 isoform X4 [Drosophila mira
>gi|1036724914|ref|XP_017146869.1| PREDICTED: histone deacetylase 6 isoform X5 [Drosophila mira
>gi|1036724877|ref|XP_017146852.1| PREDICTED: histone deacetylase 6 isoform X3 [Drosophila mira
>gi|1036724845|ref|XP_017146839.1| PREDICTED: histone deacetylase 6 isoform X2 [Drosophila mira
>gi|1036724859|ref|XP_017146845.1| PREDICTED: histone deacetylase 6 isoform X2 [Drosophila mira
>gi|1036724932|ref|XP_017146882.1| PREDICTED: LOW QUALITY PROTEIN: p21-activated protein kinase
>gi|1036735729|ref|XP_017152924.1| PREDICTED: probable ATP-dependent RNA helicase DDX46 [Drosop
>gi|1036752409|ref|XP_017156434.1| PREDICTED: inhibitory POU protein isoform X4 [Drosophila mi
>gi|1036752390|ref|XP_017156433.1| PREDICTED: inhibitory POU protein isoform X3 [Drosophila mi
>gi|1036752372|ref|XP_017156432.1| PREDICTED: inhibitory POU protein isoform X2 [Drosophila mi
>gi|1036752356|ref|XP_017156431.1| PREDICTED: inhibitory POU protein isoform X1 [Drosophila mi
>gi|1036752427|ref|XP_017156435.1| PREDICTED: serine/threonine-protein phosphatase alpha-3 iso
>gi|1036752444|ref|XP_017156437.1| PREDICTED: uncharacterized protein LOC108164952 [Drosophila
>gi|1036752463|ref|XP_017156438.1| PREDICTED: uncharacterized protein LOC108164952 [Drosophila
>gi|1036757464|ref|XP_017156727.1| PREDICTED: WSCD family member GA21586 isoform X2 [Drosophila
>gi|1036757447|ref|XP_017156726.1| PREDICTED: WSCD family member GA21586 isoform X1 [Drosophila
>gi|1036748644|ref|XP_017156223.1| PREDICTED: uncharacterized protein LOC108164814 isoform X2
>gi|1036748625|ref|XP_017156221.1| PREDICTED: uncharacterized protein LOC108164814 isoform X1
>gi|1036748663|ref|XP_017156224.1| PREDICTED: uncharacterized protein LOC108164814 isoform X3
>gi|1036699658|ref|XP_017143210.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036748682|ref|XP_017156225.1| PREDICTED: uncharacterized protein LOC108164815 [Drosophila
>gi|1036736261|ref|XP_017153228.1| PREDICTED: secretory carrier-associated membrane protein 1
>gi|1036736246|ref|XP_017153218.1| PREDICTED: adenosylhomocysteinase [Drosophila miranda]
>gi|1036716747|ref|XP_017142604.1| PREDICTED: P3 protein [Drosophila miranda]
>gi|1036716708|ref|XP_017142580.1| PREDICTED: anaphase-promoting complex subunit 1 [Drosophila
>gi|1036716785|ref|XP_017142630.1| PREDICTED: uncharacterized protein LOC108155963 [Drosophila
>gi|1036716804|ref|XP_017142637.1| PREDICTED: uncharacterized protein LOC108155963 [Drosophila
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>gi|1036716824|ref|XP_017142647.1| PREDICTED: uncharacterized protein LOC108155963 [Drosophila
>gi|1036716847|ref|XP_017142659.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036716726|ref|XP_017142593.1| PREDICTED: serine/threonine-protein kinase Nek11 [Drosophila
>gi|1036716766|ref|XP_017142616.1| PREDICTED: acetyl-CoA acetyltransferase, mitochondrial [Dros
>gi|1036757191|ref|XP_017156710.1| PREDICTED: uncharacterized protein LOC108165155 [Drosophila
>gi|1036704800|ref|XP_017135157.1| PREDICTED: selenide, water dikinase-like [Drosophila miranda
>gi|1036707217|ref|XP_017136828.1| PREDICTED: uncharacterized protein LOC108152189 [Drosophila
>gi|1036699672|ref|XP_017143336.1| PREDICTED: uncharacterized protein LOC108156386 [Drosophila
>gi|1036699685|ref|XP_017143453.1| PREDICTED: uncharacterized protein LOC108156460 [Drosophila
>gi|1036751014|ref|XP_017156353.1| PREDICTED: uncharacterized protein LOC108164897 [Drosophila
>gi|1036751032|ref|XP_017156354.1| PREDICTED: defective chorion-1 protein, FC106 isoform [Dros
>gi|1036757627|ref|XP_017156734.1| PREDICTED: golgin subfamily A member 4-like [Drosophila mira
>gi|1036699701|ref|XP_017143575.1| PREDICTED: uncharacterized protein LOC108156538 [Drosophila
>gi|1036699714|ref|XP_017143700.1| PREDICTED: mitochondrial carrier homolog 2-like [Drosophila
>gi|1036699731|ref|XP_017143948.1| PREDICTED: mitochondrial carrier homolog 2-like [Drosophila
>gi|1036699746|ref|XP_017144072.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036757755|ref|XP_017156740.1| PREDICTED: carbonic anhydrase-related protein 10 [Drosophila
>gi|1036755734|ref|XP_017156630.1| PREDICTED: AT-rich interactive domain-containing protein 1B
>gi|1036753867|ref|XP_017156524.1| PREDICTED: uncharacterized protein LOC108165002 [Drosophila
>gi|1036743371|ref|XP_017155940.1| PREDICTED: cap-specific mRNA (nucleoside-2'-O-)-methyltransi
>gi|1036747312|ref|XP_017156162.1| PREDICTED: uncharacterized protein LOC108164762 [Drosophila
>gi|1036747331|ref|XP_017156163.1| PREDICTED: uncharacterized protein LOC108164763 [Drosophila
>gi|1036757135|ref|XP_017156707.1| PREDICTED: LOW QUALITY PROTEIN: polycystic kidney disease 2-
>gi|1036756376|ref|XP_017156667.1| PREDICTED: polycystic kidney disease 2-like 1 protein isofor
>gi|1036756359|ref|XP_017156666.1| PREDICTED: polycystic kidney disease 2-like 2 protein isofor
>gi|1036756395|ref|XP_017156668.1| PREDICTED: uncharacterized protein LOC108165118 isoform X3
>gi|1036755228|ref|XP_017156604.1| PREDICTED: collagen alpha-1(II) chain-like isoform X1 [Dros
>gi|1036755244|ref|XP_017156605.1| PREDICTED: collagen alpha-1(III) chain-like isoform X2 [Dros
>gi|1036737545|ref|XP_017153972.1| PREDICTED: uncharacterized protein LOC108163279 [Drosophila
>gi|1036723410|ref|XP_017145959.1| PREDICTED: serine/threonine-protein kinase TAO3 isoform X1
>gi|1036723429|ref|XP_017145967.1| PREDICTED: serine/threonine-protein kinase TAO3 isoform X2
>gi|1036723446|ref|XP_017145979.1| PREDICTED: vacuolar protein sorting-associated protein 33A
>gi|1036735745|ref|XP_017152937.1| PREDICTED: gamma-tubulin complex component 2 homolog isoform
>gi|1036735764|ref|XP_017152944.1| PREDICTED: gamma-tubulin complex component 2 homolog isoform
>gi|1036699760|ref|XP_017144189.1| PREDICTED: homeobox protein ARX [Drosophila miranda]
>gi|1036730153|ref|XP_017150023.1| PREDICTED: REST corepressor isoform X2 [Drosophila miranda]
>gi|1036730134|ref|XP_017150015.1| PREDICTED: REST corepressor isoform X1 [Drosophila miranda]
>gi|1036730172|ref|XP_017150031.1| PREDICTED: REST corepressor isoform X3 [Drosophila miranda]
>gi|1036745649|ref|XP_017156060.1| PREDICTED: probable isocitrate dehydrogenase [NAD] subunit a
>gi|1036745667|ref|XP_017156061.1| PREDICTED: probable isocitrate dehydrogenase [NAD] subunit a
>gi|1036718720|ref|XP_017143701.1| PREDICTED: uncharacterized protein LOC108156629 [Drosophila
>gi|1036718740|ref|XP_017143712.1| PREDICTED: uncharacterized protein LOC108156629 [Drosophila
>gi|1036718769|ref|XP_017143733.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036718754|ref|XP_017143724.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036718788|ref|XP_017143741.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036718807|ref|XP_017143751.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036702925|ref|XP_017156678.1| PREDICTED: protein kish [Drosophila miranda]
>gi|1036702323|ref|XP_017156292.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prote
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>gi|1036721393|ref|XP_017145094.1| PREDICTED: LOW QUALITY PROTEIN: 5'-3' exoribonuclease 1 [Dro
>gi|1036721374|ref|XP_017145083.1| PREDICTED: probable ATP-dependent RNA helicase DHX34 [Droso
>gi|1036734653|ref|XP_017152326.1| PREDICTED: 28S ribosomal protein S14, mitochondrial [Drosop
>gi|1036734615|ref|XP_017152306.1| PREDICTED: N-alpha-acetyltransferase 16, NatA auxiliary subu
>gi|1036734634|ref|XP_017152314.1| PREDICTED: N-alpha-acetyltransferase 16, NatA auxiliary subu
>gi|1036699775|ref|XP_017144311.1| PREDICTED: uncharacterized protein LOC108156988 [Drosophila
>gi|1036722703|ref|XP_017145617.1| PREDICTED: RING-box protein 1-like [Drosophila miranda]
>gi|1036757889|ref|XP_017156748.1| PREDICTED: uncharacterized protein LOC108165188 [Drosophila
>gi|1036704819|ref|XP_017135169.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036757392|ref|XP_017156723.1| PREDICTED: nose resistant to fluoxetine protein 6-like isof
>gi|1036757410|ref|XP_017156724.1| PREDICTED: nose resistant to fluoxetine protein 6-like isof
>gi|1036757429|ref|XP_017156725.1| PREDICTED: nose resistant to fluoxetine protein 6-like isof
>gi|1036757588|ref|XP_017156732.1| PREDICTED: nardilysin-like isoform X1 [Drosophila miranda]
>gi|1036757607|ref|XP_017156733.1| PREDICTED: nardilysin-like isoform X2 [Drosophila miranda]
>gi|1036756780|ref|XP_017156690.1| PREDICTED: nose resistant to fluoxetine protein 6-like [Dros
>gi|1036708492|ref|XP_017137675.1| PREDICTED: 40S ribosomal protein S10b [Drosophila miranda]
>gi|1036708380|ref|XP_017137603.1| PREDICTED: sin3 histone deacetylase corepressor complex comp
>gi|1036708431|ref|XP_017137642.1| PREDICTED: heat shock protein beta-6 isoform X1 [Drosophila
>gi|1036708447|ref|XP_017137648.1| PREDICTED: heat shock protein beta-1 isoform X2 [Drosophila
>gi|1036708318|ref|XP_017137562.1| PREDICTED: solute carrier family 25 member 40 isoform X1 [D
>gi|1036708331|ref|XP_017137571.1| PREDICTED: solute carrier family 25 member 40 isoform X2 [D
>gi|1036708284|ref|XP_017137542.1| PREDICTED: solute carrier family 25 member 40 isoform X1 [D
>gi|1036708301|ref|XP_017137550.1| PREDICTED: solute carrier family 25 member 40 isoform X2 [D
>gi|1036708509|ref|XP_017137687.1| PREDICTED: coiled-coil domain-containing protein 86 [Drosop
>gi|1036708397|ref|XP_017137618.1| PREDICTED: L-seryl-tRNA(Sec) kinase [Drosophila miranda]
>gi|1036708525|ref|XP_017137701.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036699792|ref|XP_017144438.1| PREDICTED: uncharacterized protein LOC108157068 [Drosophila
>gi|1036708478|ref|XP_017137663.1| PREDICTED: N-alpha-acetyltransferase 20 [Drosophila miranda]
>gi|1036708346|ref|XP_017137583.1| PREDICTED: dual specificity protein phosphatase 12 [Drosophi
>gi|1036708364|ref|XP_017137591.1| PREDICTED: dual specificity protein phosphatase 12 [Drosophi
>gi|1036708414|ref|XP_017137630.1| PREDICTED: pyridoxal phosphate phosphatase PHOSPHO2 [Drosoph
>gi|1036708250|ref|XP_017137521.1| PREDICTED: supporter of activation of yellow protein [Drosop
>gi|1036708266|ref|XP_017137529.1| PREDICTED: supporter of activation of yellow protein [Drosop
>gi|1036714845|ref|XP_017141500.1| PREDICTED: pyridoxal phosphate phosphatase PHOSPHO2 [Drosoph
>gi|1036714809|ref|XP_017141480.1| PREDICTED: cell differentiation protein RCD1 homolog [Drosop
>gi|1036714828|ref|XP_017141489.1| PREDICTED: cell differentiation protein RCD1 homolog [Drosop
>gi|1036714863|ref|XP_017141510.1| PREDICTED: protein transport protein Sec61 gamma-2 subunit
>gi|1036714772|ref|XP_017141455.1| PREDICTED: signal recognition particle receptor subunit alp
>gi|1036714790|ref|XP_017141468.1| PREDICTED: 26S protease regulatory subunit 6B [Drosophila mi
>gi|1036714735|ref|XP_017141434.1| PREDICTED: uncharacterized protein LOC108155248 isoform X1
>gi|1036714753|ref|XP_017141442.1| PREDICTED: uncharacterized protein LOC108155248 isoform X2
>gi|1036742320|ref|XP_017155878.1| PREDICTED: 60 kDa heat shock protein, mitochondrial [Drosop
>gi|1036753814|ref|XP_017156520.1| PREDICTED: uncharacterized protein LOC108165001 [Drosophila
>gi|1036753831|ref|XP_017156521.1| PREDICTED: uncharacterized protein LOC108165001 [Drosophila
>gi|1036753850|ref|XP_017156523.1| PREDICTED: uncharacterized protein LOC108165001 [Drosophila
>gi|1036726131|ref|XP_017147590.1| PREDICTED: protein ELYS homolog [Drosophila miranda]
>gi|1036739261|ref|XP_017154884.1| PREDICTED: exportin-5 [Drosophila miranda]
>gi|1036732555|ref|XP_017151198.1| PREDICTED: actin-related protein 10 [Drosophila miranda]
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>gi|1036732537|ref|XP_017151189.1| PREDICTED: CWF19-like protein 2 homolog [Drosophila miranda]
>gi|1036732572|ref|XP_017151208.1| PREDICTED: 40S ribosomal protein S19a [Drosophila miranda]
>gi|1036726596|ref|XP_017147892.1| PREDICTED: rho-associated protein kinase 1 isoform X1 [Dros
>gi|1036726614|ref|XP_017147902.1| PREDICTED: rho-associated protein kinase 1 isoform X2 [Dros
>gi|1036726635|ref|XP_017147912.1| PREDICTED: rho-associated protein kinase 1 isoform X3 [Dros
>gi|1036730558|ref|XP_017150260.1| PREDICTED: brain acid soluble protein 1 homolog [Drosophila
>gi|1036730522|ref|XP_017150243.1| PREDICTED: nardilysin-like isoform X1 [Drosophila miranda]
>gi|1036730542|ref|XP_017150248.1| PREDICTED: nardilysin-like isoform X2 [Drosophila miranda]
>gi|1036751918|ref|XP_017156404.1| PREDICTED: 40S ribosomal protein S19a-like [Drosophila mira
>gi|1036741362|ref|XP_017155825.1| PREDICTED: myotubularin-related protein 10-B [Drosophila mi
>gi|1036699809|ref|XP_017144563.1| PREDICTED: uncharacterized protein LOC108157134 [Drosophila
>gi|1036752338|ref|XP_017156430.1| PREDICTED: glutaredoxin-related protein 5, mitochondrial [Dr
>gi|1036716487|ref|XP_017142475.1| PREDICTED: flotillin-2 isoform X1 [Drosophila miranda]
>gi|1036716503|ref|XP_017142484.1| PREDICTED: flotillin-2 isoform X1 [Drosophila miranda]
>gi|1036716557|ref|XP_017142492.1| PREDICTED: flotillin-2 isoform X1 [Drosophila miranda]
>gi|1036716576|ref|XP_017142501.1| PREDICTED: flotillin-2 isoform X2 [Drosophila miranda]
>gi|1036716595|ref|XP_017142511.1| PREDICTED: flotillin-2 isoform X3 [Drosophila miranda]
>gi|1036716453|ref|XP_017142450.1| PREDICTED: glucose dehydrogenase [FAD, quinone] isoform X2
>gi|1036716381|ref|XP_017142404.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716434|ref|XP_017142444.1| PREDICTED: glucose dehydrogenase [FAD, quinone] isoform X1
>gi|1036716400|ref|XP_017142420.1| PREDICTED: LOW QUALITY PROTEIN: glucose dehydrogenase [FAD,
>gi|1036716362|ref|XP_017142392.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716651|ref|XP_017142542.1| PREDICTED: uncharacterized protein LOC108155905 [Drosophila
>gi|1036716288|ref|XP_017142339.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716690|ref|XP_017142567.1| PREDICTED: glucose dehydrogenase [FAD, quinone]-like [Drosop
>gi|1036716268|ref|XP_017142328.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716470|ref|XP_017142463.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716325|ref|XP_017142368.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716670|ref|XP_017142555.1| PREDICTED: uncharacterized protein LOC108155915 [Drosophila
>gi|1036716343|ref|XP_017142379.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716306|ref|XP_017142354.1| PREDICTED: LOW QUALITY PROTEIN: glucose dehydrogenase [FAD,
>gi|1036716416|ref|XP_017142433.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716630|ref|XP_017142530.1| PREDICTED: uncharacterized protein LOC108155896 [Drosophila
>gi|1036716611|ref|XP_017142519.1| PREDICTED: flotillin-2 isoform X4 [Drosophila miranda]
>gi|1036737512|ref|XP_017153952.1| PREDICTED: probable 4-coumarate--CoA ligase 1 [Drosophila m
>gi|1036750922|ref|XP_017156348.1| PREDICTED: chromobox protein homolog 5-like [Drosophila mira
>gi|1036750848|ref|XP_017156344.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036750904|ref|XP_017156347.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036750866|ref|XP_017156345.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036750885|ref|XP_017156346.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036755473|ref|XP_017156618.1| PREDICTED: cathepsin E-B-like [Drosophila miranda]
>gi|1036743389|ref|XP_017155941.1| PREDICTED: uncharacterized protein LOC108164612 [Drosophila
>gi|1036743408|ref|XP_017155942.1| PREDICTED: branched-chain-amino-acid aminotransferase, cytos
>gi|1036723989|ref|XP_017146275.1| PREDICTED: mpv17-like protein 2 [Drosophila miranda]
>gi|1036724008|ref|XP_017146288.1| PREDICTED: protein yippee [Drosophila miranda]
>gi|1036724029|ref|XP_017146296.1| PREDICTED: protein yippee [Drosophila miranda]
>gi|1036724048|ref|XP_017146309.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036723970|ref|XP_017146263.1| PREDICTED: RNA-binding protein 1 isoform X2 [Drosophila mira
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>gi|1036723953|ref|XP_017146252.1| PREDICTED: RNA-binding protein 1 isoform X1 [Drosophila mira
>gi|1036743339|ref|XP_017155938.1| PREDICTED: alanine aminotransferase 1 [Drosophila miranda]
>gi|1036743355|ref|XP_017155939.1| PREDICTED: alanine aminotransferase 1 [Drosophila miranda]
>gi|1036706093|ref|XP_017136076.1| PREDICTED: doublesex- and mab-3-related transcription facto
>gi|1036706108|ref|XP_017136083.1| PREDICTED: uncharacterized protein LOC108151748 isoform X2
>gi|1036757699|ref|XP_017156738.1| PREDICTED: uncharacterized protein LOC108165178 [Drosophila
>gi|1036738883|ref|XP_017154657.1| PREDICTED: nuclear speckle splicing regulatory protein 1 [D
>gi|1036738901|ref|XP_017154668.1| PREDICTED: 40S ribosomal protein S15Aa [Drosophila miranda]
>gi|1036715760|ref|XP_017142027.1| PREDICTED: peroxiredoxin 1 [Drosophila miranda]
>gi|1036715712|ref|XP_017141989.1| PREDICTED: uncharacterized protein LOC108155578 isoform X2
>gi|1036715674|ref|XP_017141973.1| PREDICTED: uncharacterized protein LOC108155578 isoform X1
>gi|1036715693|ref|XP_017141982.1| PREDICTED: uncharacterized protein LOC108155578 isoform X1
>gi|1036715730|ref|XP_017142002.1| PREDICTED: uncharacterized protein LOC108155592 [Drosophila
>gi|1036715747|ref|XP_017142015.1| PREDICTED: pre-mRNA-splicing factor 38B [Drosophila miranda]
>gi|1036715640|ref|XP_017141951.1| PREDICTED: raf homolog serine/threonine-protein kinase phl
>gi|1036715657|ref|XP_017141961.1| PREDICTED: raf homolog serine/threonine-protein kinase phl
>gi|1036715779|ref|XP_017142039.1| PREDICTED: 39S ribosomal protein L14, mitochondrial [Drosop
>gi|1036737020|ref|XP_017153684.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036737038|ref|XP_017153692.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036737057|ref|XP_017153701.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036737076|ref|XP_017153710.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036737094|ref|XP_017153719.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036757793|ref|XP_017156742.1| PREDICTED: sporozoite surface protein 2-like [Drosophila mi
>gi|1036757814|ref|XP_017156743.1| PREDICTED: sporozoite surface protein 2-like [Drosophila mi
>gi|1036704513|ref|XP_017134942.1| PREDICTED: putative uncharacterized protein DDB_G0268364 [D
>gi|1036744504|ref|XP_017155993.1| PREDICTED: uncharacterized protein LOC108164652 isoform X1
>gi|1036744523|ref|XP_017155994.1| PREDICTED: uncharacterized protein LOC108164652 isoform X2
>gi|1036704226|ref|XP_017134763.1| PREDICTED: uncharacterized protein LOC108150940 [Drosophila
>gi|1036726833|ref|XP_017148037.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036743927|ref|XP_017155965.1| PREDICTED: inosine-5'-monophosphate dehydrogenase [Drosophi
>gi|1036743946|ref|XP_017155966.1| PREDICTED: inosine-5'-monophosphate dehydrogenase [Drosophi
>gi|1036755210|ref|XP_017156603.1| PREDICTED: rabphilin-3A [Drosophila miranda]
>gi|1036744063|ref|XP_017155972.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036744082|ref|XP_017155973.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036744101|ref|XP_017155974.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036748183|ref|XP_017156209.1| PREDICTED: gamma-aminobutyric acid receptor-associated prote
>gi|1036752829|ref|XP_017156460.1| PREDICTED: transcription initiation factor IIA subunit 2 [D
>gi|1036719530|ref|XP_017144168.1| PREDICTED: integrin beta-PS isoform X1 [Drosophila miranda]
>gi|1036719547|ref|XP_017144176.1| PREDICTED: integrin beta-PS isoform X1 [Drosophila miranda]
>gi|1036719567|ref|XP_017144184.1| PREDICTED: integrin beta-PS isoform X2 [Drosophila miranda]
>gi|1036719511|ref|XP_017144156.1| PREDICTED: regulator of nonsense transcripts 2 [Drosophila
>gi|1036749467|ref|XP_017156266.1| PREDICTED: dynein intermediate chain 3, ciliary [Drosophila
>gi|1036749481|ref|XP_017156268.1| PREDICTED: 17-beta-hydroxysteroid dehydrogenase 13 [Drosoph
>gi|1036699824|ref|XP_017144691.1| PREDICTED: EF-hand calcium-binding domain-containing protein
>gi|1036725087|ref|XP_017146972.1| PREDICTED: ubiquitin-conjugating enzyme E2 H [Drosophila mi
>gi|1036725103|ref|XP_017146979.1| PREDICTED: ubiquitin-conjugating enzyme E2 H [Drosophila mi
>gi|1036725070|ref|XP_017146959.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036725116|ref|XP_017146990.1| PREDICTED: uncharacterized protein LOC108158809 [Drosophila
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>gi|1036725134|ref|XP_017146999.1| PREDICTED: uncharacterized protein LOC108158809 [Drosophila
>gi|1036725152|ref|XP_017147006.1| PREDICTED: uncharacterized protein LOC108158809 [Drosophila
>gi|1036749533|ref|XP_017156271.1| PREDICTED: uncharacterized protein LOC108164853 [Drosophila
>gi|1036699841|ref|XP_017144812.1| PREDICTED: inosine-5'-monophosphate dehydrogenase-like [Dros
>gi|1036730503|ref|XP_017150229.1| PREDICTED: putative folylpolyglutamate synthase [Drosophila
>gi|1036730484|ref|XP_017150217.1| PREDICTED: uncharacterized protein LOC108160603 isoform X2
>gi|1036730450|ref|XP_017150200.1| PREDICTED: uncharacterized protein LOC108160603 isoform X1
>gi|1036730467|ref|XP_017150208.1| PREDICTED: uncharacterized protein LOC108160603 isoform X1
>gi|1036756545|ref|XP_017156676.1| PREDICTED: YEATS domain-containing protein 4 [Drosophila mir
>gi|1036756524|ref|XP_017156675.1| PREDICTED: uncharacterized protein LOC108165123 [Drosophila
>gi|1036717066|ref|XP_017142799.1| PREDICTED: protein kinase shaggy isoform X2 [Drosophila mira
>gi|1036717086|ref|XP_017142807.1| PREDICTED: protein kinase shaggy isoform X3 [Drosophila mira
>gi|1036717047|ref|XP_017142789.1| PREDICTED: protein kinase shaggy isoform X1 [Drosophila mira
>gi|1036732065|ref|XP_017150966.1| PREDICTED: phosphatidylinositol 4-kinase alpha [Drosophila m
>gi|1036732615|ref|XP_017151230.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036732633|ref|XP_017151238.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036732651|ref|XP_017151246.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036732669|ref|XP_017151253.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036737668|ref|XP_017154048.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036737686|ref|XP_017154058.1| PREDICTED: enhancer of yellow 2 transcription factor isoform
>gi|1036737703|ref|XP_017154062.1| PREDICTED: enhancer of yellow 2 transcription factor isoform
>gi|1036737652|ref|XP_017154034.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036744992|ref|XP_017156020.1| PREDICTED: uncharacterized protein LOC108164676 [Drosophila
>gi|1036745011|ref|XP_017156021.1| PREDICTED: transmembrane protein 242 [Drosophila miranda]
>gi|1036744937|ref|XP_017156017.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036744954|ref|XP_017156018.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036744973|ref|XP_017156019.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036723507|ref|XP_017146006.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB1 [Dros
>gi|1036722753|ref|XP_017145636.1| PREDICTED: putative ATP-dependent RNA helicase BoYb [Drosoph
>gi|1036722997|ref|XP_017145742.1| PREDICTED: LOW QUALITY PROTEIN: peptidoglycan-recognition pr
>gi|1036722770|ref|XP_017145646.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722791|ref|XP_017145653.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722812|ref|XP_017145661.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722855|ref|XP_017145669.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722874|ref|XP_017145676.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722889|ref|XP_017145683.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722925|ref|XP_017145698.1| PREDICTED: homeobox protein 5 isoform X3 [Drosophila miranda
>gi|1036722906|ref|XP_017145691.1| PREDICTED: homeobox protein 5 isoform X2 [Drosophila miranda
>gi|1036722943|ref|XP_017145706.1| PREDICTED: homeobox protein 5 isoform X4 [Drosophila miranda
>gi|1036722980|ref|XP_017145727.1| PREDICTED: coiled-coil domain-containing protein 25 [Drosoph
>gi|1036722962|ref|XP_017145716.1| PREDICTED: facilitated trehalose transporter Tret1 [Drosophi
>gi|1036720091|ref|XP_017144450.1| PREDICTED: putative uncharacterized protein DDB_G0286901 [Dr
>gi|1036720053|ref|XP_017144428.1| PREDICTED: protein sex-lethal isoform X10 [Drosophila miranc
>gi|1036719889|ref|XP_017144358.1| PREDICTED: protein sex-lethal isoform X2 [Drosophila miranda
>gi|1036719972|ref|XP_017144392.1| PREDICTED: protein sex-lethal isoform X6 [Drosophila miranda
>gi|1036719869|ref|XP_017144350.1| PREDICTED: protein sex-lethal isoform X1 [Drosophila miranda
>gi|1036720073|ref|XP_017144439.1| PREDICTED: protein sex-lethal isoform X11 [Drosophila miranc
>gi|1036719908|ref|XP_017144368.1| PREDICTED: protein sex-lethal isoform X3 [Drosophila miranda
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>gi|1036720033|ref|XP_017144419.1| PREDICTED: protein sex-lethal isoform X9 [Drosophila miranda
>gi|1036720014|ref|XP_017144408.1| PREDICTED: protein sex-lethal isoform X8 [Drosophila miranda
>gi|1036719928|ref|XP_017144375.1| PREDICTED: protein sex-lethal isoform X4 [Drosophila miranda
>gi|1036719952|ref|XP_017144383.1| PREDICTED: protein sex-lethal isoform X5 [Drosophila miranda
>gi|1036719994|ref|XP_017144400.1| PREDICTED: protein sex-lethal isoform X7 [Drosophila miranda
>gi|1036710707|ref|XP_017139059.1| PREDICTED: protein suppressor of sable isoform X2 [Drosophil
>gi|1036710692|ref|XP_017139050.1| PREDICTED: protein suppressor of sable isoform X1 [Drosophil
>gi|1036710744|ref|XP_017139082.1| PREDICTED: caspase-8 [Drosophila miranda]
>gi|1036710726|ref|XP_017139070.1| PREDICTED: uncharacterized protein LOC108153519 [Drosophila
>gi|1036710776|ref|XP_017139105.1| PREDICTED: tetraspanin-33 [Drosophila miranda]
>gi|1036710793|ref|XP_017139114.1| PREDICTED: tetraspanin-33 [Drosophila miranda]
>gi|1036710760|ref|XP_017139093.1| PREDICTED: seipin [Drosophila miranda]
>gi|1036748948|ref|XP_017156238.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X2 [Dros
>gi|1036748907|ref|XP_017156236.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X1 [Dros
>gi|1036748929|ref|XP_017156237.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X1 [Dros
>gi|1036745978|ref|XP_017156078.1| PREDICTED: 60S ribosomal protein L17 [Drosophila miranda]
>gi|1036744920|ref|XP_017156016.1| PREDICTED: putative metabolite transport protein HI_1104 [Dr
>gi|1036711151|ref|XP_017139364.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711167|ref|XP_017139372.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711184|ref|XP_017139381.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711199|ref|XP_017139389.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711324|ref|XP_017139426.1| PREDICTED: uncharacterized protein LOC108153835 [Drosophila
>gi|1036711290|ref|XP_017139421.1| PREDICTED: uncharacterized protein LOC108153801 [Drosophila
>gi|1036711307|ref|XP_017139422.1| PREDICTED: 5-demethoxyubiquinone hydroxylase, mitochondrial
>gi|1036711273|ref|XP_017139420.1| PREDICTED: protein shifted isoform X4 [Drosophila miranda]
>gi|1036711256|ref|XP_017139415.1| PREDICTED: protein shifted isoform X3 [Drosophila miranda]
>gi|1036711235|ref|XP_017139408.1| PREDICTED: protein shifted isoform X2 [Drosophila miranda]
>gi|1036711216|ref|XP_017139399.1| PREDICTED: protein shifted isoform X1 [Drosophila miranda]
>gi|1036726982|ref|XP_017148114.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X4
>gi|1036727021|ref|XP_017148131.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X6
>gi|1036726963|ref|XP_017148107.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X3
>gi|1036726945|ref|XP_017148099.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X2
>gi|1036726927|ref|XP_017148092.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X1
>gi|1036726999|ref|XP_017148122.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X5
>gi|1036725173|ref|XP_017147019.1| PREDICTED: coronin-7 isoform X1 [Drosophila miranda]
>gi|1036725191|ref|XP_017147027.1| PREDICTED: coronin-7 isoform X2 [Drosophila miranda]
>gi|1036725210|ref|XP_017147038.1| PREDICTED: coronin-7 isoform X3 [Drosophila miranda]
>gi|1036725226|ref|XP_017147047.1| PREDICTED: coronin-7 isoform X4 [Drosophila miranda]
>gi|1036737198|ref|XP_017153777.1| PREDICTED: uncharacterized protein LOC108163163 [Drosophila
>gi|1036737233|ref|XP_017153800.1| PREDICTED: putative uncharacterized protein DDB_G0268364 iso
>gi|1036737213|ref|XP_017153791.1| PREDICTED: putative uncharacterized protein DDB_G0268364 iso
>gi|1036744540|ref|XP_017155995.1| PREDICTED: probable dolichyl pyrophosphate Glc1Man9GlcNAc2 a
>gi|1036744556|ref|XP_017155996.1| PREDICTED: ataxin-1 [Drosophila miranda]
>gi|1036742682|ref|XP_017155900.1| PREDICTED: frizzled-3 [Drosophila miranda]
>gi|1036749239|ref|XP_017156255.1| PREDICTED: protein O-mannosyl-transferase 2 [Drosophila mira
>gi|1036749258|ref|XP_017156256.1| PREDICTED: uncharacterized protein LOC108164841 [Drosophila
>gi|1036749297|ref|XP_017156259.1| PREDICTED: uncharacterized protein LOC108164842 isoform X2
>gi|1036749278|ref|XP_017156258.1| PREDICTED: uncharacterized protein LOC108164842 isoform X1
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>gi|1036749327|ref|XP_017156260.1| PREDICTED: uncharacterized protein LOC108164842 isoform X3
>gi|1036699856|ref|XP_017144927.1| PREDICTED: uncharacterized protein LOC108157372 [Drosophila
>gi|1036749342|ref|XP_017156261.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036749360|ref|XP_017156262.1| PREDICTED: uncharacterized protein LOC108164844 [Drosophila
>gi|1036717849|ref|XP_017143223.1| PREDICTED: histone deacetylase complex subunit SAP30 homolog
>gi|1036717731|ref|XP_017143155.1| PREDICTED: exosome complex component RRP45 [Drosophila miran
>gi|1036717772|ref|XP_017143180.1| PREDICTED: uncharacterized protein LOC108156303 isoform X1
>gi|1036717791|ref|XP_017143187.1| PREDICTED: UPF0472 protein C16orf72 isoform X2 [Drosophila r
>gi|1036717809|ref|XP_017143198.1| PREDICTED: NECAP-like protein CG9132 [Drosophila miranda]
>gi|1036717753|ref|XP_017143167.1| PREDICTED: zinc finger protein 845 [Drosophila miranda]
>gi|1036717830|ref|XP_017143211.1| PREDICTED: 39S ribosomal protein L22, mitochondrial [Drosoph
>gi|1036744327|ref|XP_017155985.1| PREDICTED: uncharacterized protein LOC108164646 [Drosophila
>gi|1036730907|ref|XP_017150464.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730926|ref|XP_017150472.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730940|ref|XP_017150481.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730959|ref|XP_017150488.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730979|ref|XP_017150498.1| PREDICTED: integrin alpha-PS2 isoform X2 [Drosophila miranda
>gi|1036705657|ref|XP_017135737.1| PREDICTED: transmembrane protein 256 homolog [Drosophila mi
>gi|1036728452|ref|XP_017148976.1| PREDICTED: endothelin-converting enzyme 1 isoform X1 [Drosop
>gi|1036728471|ref|XP_017148984.1| PREDICTED: endothelin-converting enzyme 1 isoform X2 [Drosop
>gi|1036728532|ref|XP_017149017.1| PREDICTED: keratin, type I cytoskeletal 10 [Drosophila miran
>gi|1036728585|ref|XP_017149055.1| PREDICTED: uncharacterized protein LOC108159922 [Drosophila
>gi|1036728568|ref|XP_017149042.1| PREDICTED: uncharacterized protein LOC108159916 [Drosophila
>gi|1036728550|ref|XP_017149031.1| PREDICTED: uncharacterized protein LOC108159907 [Drosophila
>gi|1036728509|ref|XP_017149006.1| PREDICTED: uncharacterized protein LOC108159887 isoform X2
>gi|1036728490|ref|XP_017148997.1| PREDICTED: uncharacterized protein LOC108159887 isoform X1
>gi|1036728604|ref|XP_017149068.1| PREDICTED: transforming growth factor beta-1-induced transcr
>gi|1036728621|ref|XP_017149076.1| PREDICTED: transforming growth factor beta-1-induced transcr
>gi|1036747573|ref|XP_017156177.1| PREDICTED: uncharacterized protein LOC108164772 [Drosophila
>gi|1036747631|ref|XP_017156180.1| PREDICTED: carboxypeptidase D-like isoform X3 [Drosophila mi
>gi|1036747612|ref|XP_017156179.1| PREDICTED: carboxypeptidase D-like isoform X2 [Drosophila mi
>gi|1036747591|ref|XP_017156178.1| PREDICTED: carboxypeptidase D-like isoform X1 [Drosophila mi
>gi|1036747666|ref|XP_017156182.1| PREDICTED: carboxypeptidase D-like isoform X5 [Drosophila mi
>gi|1036747649|ref|XP_017156181.1| PREDICTED: carboxypeptidase D-like isoform X4 [Drosophila mi
>gi|1036747722|ref|XP_017156186.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036747760|ref|XP_017156188.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036747685|ref|XP_017156183.1| PREDICTED: trypsin beta-like [Drosophila miranda]
>gi|1036747741|ref|XP_017156187.1| PREDICTED: chymotrypsin-1-like [Drosophila miranda]
>gi|1036747704|ref|XP_017156184.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036717598|ref|XP_017143079.1| PREDICTED: transmembrane GTPase Marf isoform X3 [Drosophila
>gi|1036717579|ref|XP_017143072.1| PREDICTED: transmembrane GTPase Marf isoform X2 [Drosophila
>gi|1036717554|ref|XP_017143062.1| PREDICTED: transmembrane GTPase Marf isoform X1 [Drosophila
>gi|1036717617|ref|XP_017143092.1| PREDICTED: probable citrate synthase, mitochondrial isoform
>gi|1036717628|ref|XP_017143101.1| PREDICTED: probable citrate synthase, mitochondrial isoform
>gi|1036717647|ref|XP_017143110.1| PREDICTED: probable citrate synthase, mitochondrial isoform
>gi|1036717694|ref|XP_017143135.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036717712|ref|XP_017143143.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036717676|ref|XP_017143123.1| PREDICTED: zinc finger protein 624 [Drosophila miranda]
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>gi|1036734492|ref|XP_017152219.1| PREDICTED: retinol dehydrogenase 12 [Drosophila miranda]
>gi|1036734510|ref|XP_017152232.1| PREDICTED: LOW QUALITY PROTEIN: probable myosin light chain
>gi|1036734427|ref|XP_017152198.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036734445|ref|XP_017152206.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036734526|ref|XP_017152246.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036755381|ref|XP_017156613.1| PREDICTED: alpha-tocopherol transfer protein [Drosophila mi
>gi|1036713245|ref|XP_017140494.1| PREDICTED: E3 ubiquitin-protein ligase highwire isoform X2
>gi|1036713228|ref|XP_017140487.1| PREDICTED: E3 ubiquitin-protein ligase highwire isoform X1
>gi|1036713262|ref|XP_017140506.1| PREDICTED: uncharacterized protein LOC108154673 [Drosophila
>gi|1036713278|ref|XP_017140517.1| PREDICTED: uncharacterized protein LOC108154684 [Drosophila
>gi|1036713295|ref|XP_017140527.1| PREDICTED: uncharacterized protein LOC108154684 [Drosophila
>gi|1036746647|ref|XP_017156121.1| PREDICTED: transmembrane emp24 domain-containing protein 5
>gi|1036746666|ref|XP_017156122.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036728171|ref|XP_017148819.1| PREDICTED: probable DNA replication complex GINS protein PSF
>gi|1036728149|ref|XP_017148809.1| PREDICTED: protein FAM57B [Drosophila miranda]
>gi|1036698255|ref|XP_017137402.1| PREDICTED: uncharacterized protein LOC108152276 [Drosophila
>gi|1036728131|ref|XP_017148795.1| PREDICTED: proline-rich protein 36 [Drosophila miranda]
>gi|1036729098|ref|XP_017149359.1| PREDICTED: plexin domain-containing protein 2 [Drosophila mi
>gi|1036729135|ref|XP_017149386.1| PREDICTED: perlucin-like [Drosophila miranda]
>gi|1036729116|ref|XP_017149373.1| PREDICTED: accessory gland protein Acp29AB-like [Drosophila
>gi|1036754428|ref|XP_017156557.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036745181|ref|XP_017156032.1| PREDICTED: putative hydroxypyruvate isomerase [Drosophila mi
>gi|1036745162|ref|XP_017156031.1| PREDICTED: uncharacterized protein LOC108164680 [Drosophila
>gi|1036709629|ref|XP_017138340.1| PREDICTED: serine/threonine-protein phosphatase 4 catalytic
>gi|1036709870|ref|XP_017138492.1| PREDICTED: general odorant-binding protein 19d [Drosophila
>gi|1036709817|ref|XP_017138456.1| PREDICTED: general odorant-binding protein 56a [Drosophila
>gi|1036709851|ref|XP_017138479.1| PREDICTED: uncharacterized protein LOC108153179 [Drosophila
>gi|1036709887|ref|XP_017138504.1| PREDICTED: general odorant-binding protein 19a [Drosophila
>gi|1036709921|ref|XP_017138528.1| PREDICTED: putative 60S ribosomal protein L33 [Drosophila mi
>gi|1036709733|ref|XP_017138401.1| PREDICTED: uncharacterized protein LOC108153120 [Drosophila
>gi|1036709646|ref|XP_017138352.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC2 [Dros
>gi|1036709716|ref|XP_017138388.1| PREDICTED: syntaxin-16 [Drosophila miranda]
>gi|1036709750|ref|XP_017138412.1| PREDICTED: RNA-binding protein pno1 [Drosophila miranda]
>gi|1036709767|ref|XP_017138426.1| PREDICTED: pre-rRNA-processing protein esf2-like [Drosophila
>gi|1036709834|ref|XP_017138467.1| PREDICTED: AN1-type zinc finger protein 6 [Drosophila mirand
>gi|1036709784|ref|XP_017138438.1| PREDICTED: uncharacterized protein LOC108153144 isoform X1
>gi|1036709800|ref|XP_017138445.1| PREDICTED: uncharacterized protein LOC108153144 isoform X2
>gi|1036709904|ref|XP_017138517.1| PREDICTED: sarcocystatin-A [Drosophila miranda]
>gi|1036709698|ref|XP_017138378.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036709665|ref|XP_017138358.1| PREDICTED: dihydropyrimidine dehydrogenase [NADP(+)] isoform
>gi|1036709682|ref|XP_017138366.1| PREDICTED: dihydropyrimidine dehydrogenase [NADP(+)] isoform
>gi|1036737111|ref|XP_017153732.1| PREDICTED: NCK-interacting protein with SH3 domain [Drosophi
>gi|1036730102|ref|XP_017149981.1| PREDICTED: protein will die slowly [Drosophila miranda]
>gi|1036730116|ref|XP_017149989.1| PREDICTED: protein will die slowly [Drosophila miranda]
>gi|1036730083|ref|XP_017149968.1| PREDICTED: pyruvate dehydrogenase E1 component subunit alpha
>gi|1036730062|ref|XP_017149958.1| PREDICTED: probable pyruvate dehydrogenase E1 component subu
>gi|1036730009|ref|XP_017149929.1| PREDICTED: uncharacterized protein LOC108160432 [Drosophila
>gi|1036730024|ref|XP_017149937.1| PREDICTED: uncharacterized protein LOC108160432 [Drosophila
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>gi|1036730043|ref|XP_017149945.1| PREDICTED: uncharacterized protein LOC108160432 [Drosophila
>gi|1036706663|ref|XP_017136472.1| PREDICTED: uncharacterized protein LOC108152006 [Drosophila
>gi|1036703804|ref|XP_017157172.1| PREDICTED: E3 ubiquitin-protein ligase RNF185-like [Drosophi
>gi|1036757871|ref|XP_017156746.1| PREDICTED: putative inorganic phosphate cotransporter [Droso
>gi|1036706143|ref|XP_017136103.1| PREDICTED: uncharacterized protein LOC108151766 [Drosophila
>gi|1036706436|ref|XP_017136302.1| PREDICTED: Krueppel-like factor 6 [Drosophila miranda]
>gi|1036747885|ref|XP_017156195.1| PREDICTED: general transcription factor IIH subunit 3-like
>gi|1036747864|ref|XP_017156194.1| PREDICTED: uncharacterized protein LOC108164786 [Drosophila
>gi|1036747848|ref|XP_017156193.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1036699871|ref|XP_017145044.1| PREDICTED: uncharacterized protein LOC108157453 [Drosophila
>gi|1036739368|ref|XP_017154956.1| PREDICTED: glycine receptor subunit alpha-4 [Drosophila mira
>gi|1036739384|ref|XP_017154969.1| PREDICTED: zinc finger protein 391 isoform X1 [Drosophila m
>gi|1036739400|ref|XP_017154977.1| PREDICTED: gastrula zinc finger protein XlCGF8.2DB isoform X
>gi|1036699886|ref|XP_017145164.1| PREDICTED: nardilysin [Drosophila miranda]
>gi|1036719418|ref|XP_017144118.1| PREDICTED: nardilysin-like isoform X1 [Drosophila miranda]
>gi|1036719436|ref|XP_017144126.1| PREDICTED: nardilysin-like isoform X2 [Drosophila miranda]
>gi|1036719399|ref|XP_017144106.1| PREDICTED: copper-transporting ATPase 1 isoform X2 [Drosophi
>gi|1036719380|ref|XP_017144097.1| PREDICTED: copper-transporting ATPase 1 isoform X1 [Drosophi
>gi|1036741195|ref|XP_017155813.1| PREDICTED: replication protein A 14 kDa subunit A [Drosophil
>gi|1036741177|ref|XP_017155812.1| PREDICTED: ATP-binding cassette sub-family F member 1 [Droso
>gi|1036712680|ref|XP_017140170.1| PREDICTED: circadian locomoter output cycles protein kaput
>gi|1036712646|ref|XP_017140150.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X2 [Dros
>gi|1036712663|ref|XP_017140158.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X3 [Dros
>gi|1036712629|ref|XP_017140142.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036712714|ref|XP_017140190.1| PREDICTED: uncharacterized protein LOC108154429 [Drosophila
>gi|1036712697|ref|XP_017140180.1| PREDICTED: lysophosphatidylcholine acyltransferase [Drosophi
>gi|1036740908|ref|XP_017155798.1| PREDICTED: cubilin homolog [Drosophila miranda]
>gi|1036740924|ref|XP_017155799.1| PREDICTED: lon protease homolog, mitochondrial-like [Drosop
>gi|1036703949|ref|XP_017134599.1| PREDICTED: uncharacterized protein LOC108150803 [Drosophila
>gi|1036707589|ref|XP_017137082.1| PREDICTED: LOW QUALITY PROTEIN: glycine-rich cell wall stru
>gi|1036725363|ref|XP_017147111.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725306|ref|XP_017147084.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725287|ref|XP_017147076.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725269|ref|XP_017147066.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725251|ref|XP_017147058.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725327|ref|XP_017147093.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725346|ref|XP_017147102.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725399|ref|XP_017147141.1| PREDICTED: uncharacterized protein LOC108158881 [Drosophila
>gi|1036699905|ref|XP_017145282.1| PREDICTED: skin secretory protein xP2 [Drosophila miranda]
>gi|1036725382|ref|XP_017147125.1| PREDICTED: ATP synthase subunit delta, mitochondrial [Droso
>gi|1036723542|ref|XP_017146018.1| PREDICTED: LOW QUALITY PROTEIN: nuclear envelope integral me
>gi|1036723561|ref|XP_017146029.1| PREDICTED: uncharacterized protein LOC108158297 [Drosophila
>gi|1036723580|ref|XP_017146039.1| PREDICTED: uncharacterized protein LOC108158307 [Drosophila
>gi|1036707251|ref|XP_017136854.1| PREDICTED: kunitz-type serine protease inhibitor textilinin-
>gi|1036732685|ref|XP_017151265.1| PREDICTED: myc protein [Drosophila miranda]
>gi|1036699920|ref|XP_017145517.1| PREDICTED: uncharacterized protein LOC108157795 [Drosophila
>gi|1036747236|ref|XP_017156157.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X3 [Di
>gi|1036747251|ref|XP_017156158.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X3 [Di
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>gi|1036747167|ref|XP_017156153.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X1 [Di
>gi|1036747184|ref|XP_017156154.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X1 [Di
>gi|1036747203|ref|XP_017156155.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X1 [Di
>gi|1036747272|ref|XP_017156159.1| PREDICTED: uncharacterized protein LOC108164760 [Drosophila
>gi|1036747220|ref|XP_017156156.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X2 [Di
>gi|1036755573|ref|XP_017156620.1| PREDICTED: glutaredoxin domain-containing cysteine-rich prot
>gi|1036705085|ref|XP_017135370.1| PREDICTED: calumenin-B-like [Drosophila miranda]
>gi|1036750995|ref|XP_017156352.1| PREDICTED: sodium-dependent neutral amino acid transporter 
>gi|1036726707|ref|XP_017147957.1| PREDICTED: alanyl-tRNA editing protein Aarsd1-B [Drosophila 
>gi|1036726757|ref|XP_017147996.1| PREDICTED: uncharacterized protein C19orf52 [Drosophila mira
>gi|1036726776|ref|XP_017148004.1| PREDICTED: uncharacterized protein C19orf52 [Drosophila mira
>gi|1036726795|ref|XP_017148013.1| PREDICTED: uncharacterized protein C19orf52 [Drosophila mira
>gi|1036726740|ref|XP_017147981.1| PREDICTED: uncharacterized protein LOC108159314 [Drosophila 
>gi|1036726814|ref|XP_017148024.1| PREDICTED: 40S ribosomal protein S28 [Drosophila miranda]
>gi|1036726723|ref|XP_017147969.1| PREDICTED: translocon-associated protein subunit alpha [Dros
>gi|1036699936|ref|XP_017145583.1| PREDICTED: LOW QUALITY PROTEIN: ran GTPase-activating protei
>gi|1036737740|ref|XP_017154082.1| PREDICTED: uncharacterized protein LOC108163349 [Drosophila 
>gi|1036737720|ref|XP_017154072.1| PREDICTED: uncharacterized protein LOC108163342 [Drosophila 
>gi|1036706351|ref|XP_017136241.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosoph
>gi|1036713311|ref|XP_017140538.1| PREDICTED: low-density lipoprotein receptor-related protein 
>gi|1036745613|ref|XP_017156058.1| PREDICTED: B-cell CLL/lymphoma 7 protein family member A [Di
>gi|1036720303|ref|XP_017144564.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036720350|ref|XP_017144580.1| PREDICTED: palmitoyltransferase ZDHHC5 isoform X3 [Drosophil
>gi|1036720322|ref|XP_017144572.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036720369|ref|XP_017144589.1| PREDICTED: palmitoyltransferase ZDHHC5 isoform X4 [Drosophil
>gi|1036720407|ref|XP_017144605.1| PREDICTED: probable protein S-acyltransferase 2 isoform X6 
>gi|1036720426|ref|XP_017144613.1| PREDICTED: palmitoyltransferase ZDHHC9 isoform X7 [Drosophil
>gi|1036720390|ref|XP_017144597.1| PREDICTED: neurogenic protein mastermind isoform X5 [Drosoph
>gi|1036720285|ref|XP_017144552.1| PREDICTED: von Willebrand factor A domain-containing protei
>gi|1036720507|ref|XP_017144658.1| PREDICTED: protein CWC15 homolog [Drosophila miranda]
>gi|1036720445|ref|XP_017144625.1| PREDICTED: protein lozenge [Drosophila miranda]
>gi|1036720465|ref|XP_017144636.1| PREDICTED: protein lozenge [Drosophila miranda]
>gi|1036720485|ref|XP_017144645.1| PREDICTED: protein lozenge [Drosophila miranda]
>gi|1036731878|ref|XP_017150882.1| PREDICTED: maestro heat-like repeat-containing protein fami
>gi|1036731897|ref|XP_017150893.1| PREDICTED: aladin [Drosophila miranda]
>gi|1036703130|ref|XP_017156793.1| PREDICTED: uncharacterized protein LOC108165265 isoform X1 
>gi|1036703149|ref|XP_017156798.1| PREDICTED: uncharacterized protein LOC108165265 isoform X2 
>gi|1036702601|ref|XP_017156471.1| PREDICTED: CXXC-type zinc finger protein 1-like [Drosophila
>gi|1036702454|ref|XP_017156392.1| PREDICTED: uncharacterized protein LOC108164915 isoform X1 
>gi|1036702470|ref|XP_017156400.1| PREDICTED: uncharacterized protein LOC108164915 isoform X1 
>gi|1036702487|ref|XP_017156408.1| PREDICTED: uncharacterized protein LOC108164915 isoform X2 
>gi|1036702617|ref|XP_017156484.1| PREDICTED: synembryn [Drosophila miranda]
>gi|1036702567|ref|XP_017156450.1| PREDICTED: kelch-like protein 5 isoform X1 [Drosophila mira
>gi|1036702584|ref|XP_017156459.1| PREDICTED: kelch-like protein 5 isoform X2 [Drosophila mira
>gi|1036702649|ref|XP_017156509.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702633|ref|XP_017156497.1| PREDICTED: uncharacterized protein LOC108164982 [Drosophila
>gi|1036699953|ref|XP_017145690.1| PREDICTED: LOW QUALITY PROTEIN: pickpocket protein 28 [Droso
>gi|1036702506|ref|XP_017156419.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X1 [Drosop
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>gi|1036702552|ref|XP_017156436.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X3 [Droso
>gi|1036702535|ref|XP_017156427.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X2 [Droso
>gi|1036702666|ref|XP_017156522.1| PREDICTED: uncharacterized protein LOC108164998 [Drosophila
>gi|1036702437|ref|XP_017156381.1| PREDICTED: sodium/potassium-transporting ATPase subunit alp
>gi|1036702405|ref|XP_017156358.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702390|ref|XP_017156350.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702421|ref|XP_017156368.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702753|ref|XP_017156573.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702772|ref|XP_017156585.1| PREDICTED: probable alpha-aspartyl dipeptidase [Drosophila m
>gi|1036702698|ref|XP_017156540.1| PREDICTED: chondroitin sulfate glucuronyltransferase [Drosop
>gi|1036702787|ref|XP_017156599.1| PREDICTED: soma ferritin [Drosophila miranda]
>gi|1036702717|ref|XP_017156550.1| PREDICTED: actin-related protein 6 [Drosophila miranda]
>gi|1036702838|ref|XP_017156627.1| PREDICTED: iron-sulfur cluster assembly 1 homolog, mitochond
>gi|1036702821|ref|XP_017156619.1| PREDICTED: iron-sulfur cluster assembly 1 homolog, mitochond
>gi|1036702804|ref|XP_017156609.1| PREDICTED: DNL-type zinc finger protein [Drosophila miranda]
>gi|1036702734|ref|XP_017156562.1| PREDICTED: uncharacterized protein LOC108165033 [Drosophila
>gi|1036717282|ref|XP_017142918.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X4 [Dro
>gi|1036717242|ref|XP_017142900.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X2 [Dro
>gi|1036717301|ref|XP_017142927.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X5 [Dro
>gi|1036717263|ref|XP_017142909.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X3 [Dro
>gi|1036717224|ref|XP_017142892.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X1 [Dro
>gi|1036717322|ref|XP_017142938.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu
>gi|1036753514|ref|XP_017156503.1| PREDICTED: uncharacterized protein LOC108164988 [Drosophila
>gi|1036736826|ref|XP_017153559.1| PREDICTED: ATP-binding cassette sub-family F member 2 [Droso
>gi|1036748164|ref|XP_017156208.1| PREDICTED: uncharacterized protein LOC108164795 [Drosophila
>gi|1036734022|ref|XP_017151984.1| PREDICTED: CDP-diacylglycerol--inositol 3-phosphatidyltrans
>gi|1036734041|ref|XP_017151997.1| PREDICTED: mitochondrial inner membrane protease subunit 1
>gi|1036733966|ref|XP_017151952.1| PREDICTED: nudix hydrolase 8 [Drosophila miranda]
>gi|1036733985|ref|XP_017151962.1| PREDICTED: nudix hydrolase 8 [Drosophila miranda]
>gi|1036734003|ref|XP_017151971.1| PREDICTED: nudix hydrolase 8 [Drosophila miranda]
>gi|1036699966|ref|XP_017145814.1| PREDICTED: uncharacterized protein LOC108158109 [Drosophila
>gi|1036755682|ref|XP_017156626.1| PREDICTED: uncharacterized protein LOC108165086 [Drosophila
>gi|1036699987|ref|XP_017145930.1| PREDICTED: visual system homeobox 2, partial [Drosophila mir
>gi|1036750499|ref|XP_017156323.1| PREDICTED: homeotic protein ocelliless isoform X3 [Drosophil
>gi|1036750518|ref|XP_017156324.1| PREDICTED: homeotic protein ocelliless isoform X4 [Drosophil
>gi|1036750537|ref|XP_017156325.1| PREDICTED: uncharacterized protein LOC108164886 [Drosophila
>gi|1036750460|ref|XP_017156320.1| PREDICTED: diencephalon/mesencephalon homeobox protein 1 iso
>gi|1036750477|ref|XP_017156321.1| PREDICTED: diencephalon/mesencephalon homeobox protein 1 iso
>gi|1036750442|ref|XP_017156319.1| PREDICTED: uncharacterized protein LOC108164884 [Drosophila
>gi|1036750425|ref|XP_017156318.1| PREDICTED: N-chimaerin [Drosophila miranda]
>gi|1036736551|ref|XP_017153409.1| PREDICTED: myosin-2 essential light chain isoform X2 [Drosop
>gi|1036736538|ref|XP_017153400.1| PREDICTED: myosin-2 essential light chain isoform X1 [Drosop
>gi|1036736486|ref|XP_017153372.1| PREDICTED: ribonucleases P/MRP protein subunit POP1 [Drosop
>gi|1036736505|ref|XP_017153381.1| PREDICTED: ribonucleases P/MRP protein subunit POP1 [Drosop
>gi|1036736522|ref|XP_017153389.1| PREDICTED: ribonucleases P/MRP protein subunit POP1 [Drosop
>gi|1036737952|ref|XP_017154196.1| PREDICTED: uncharacterized protein LOC108163366 isoform X12
>gi|1036737883|ref|XP_017154162.1| PREDICTED: uncharacterized protein LOC108163366 isoform X8
>gi|1036737936|ref|XP_017154186.1| PREDICTED: uncharacterized protein LOC108163366 isoform X11
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>gi|1036737811|ref|XP_017154129.1| PREDICTED: uncharacterized protein LOC108163366 isoform X4
>gi|1036737849|ref|XP_017154146.1| PREDICTED: uncharacterized protein LOC108163366 isoform X6
>gi|1036737865|ref|XP_017154154.1| PREDICTED: uncharacterized protein LOC108163366 isoform X7
>gi|1036737793|ref|XP_017154122.1| PREDICTED: uncharacterized protein LOC108163366 isoform X3
>gi|1036737829|ref|XP_017154137.1| PREDICTED: uncharacterized protein LOC108163366 isoform X5
>gi|1036737775|ref|XP_017154117.1| PREDICTED: uncharacterized protein LOC108163366 isoform X2
>gi|1036737758|ref|XP_017154110.1| PREDICTED: uncharacterized protein LOC108163366 isoform X1
>gi|1036737970|ref|XP_017154204.1| PREDICTED: uncharacterized protein LOC108163366 isoform X13
>gi|1036737918|ref|XP_017154178.1| PREDICTED: uncharacterized protein LOC108163366 isoform X10
>gi|1036737901|ref|XP_017154170.1| PREDICTED: uncharacterized protein LOC108163366 isoform X9
>gi|1036749952|ref|XP_017156295.1| PREDICTED: uncharacterized protein LOC108164866 isoform X2
>gi|1036749879|ref|XP_017156290.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749897|ref|XP_017156291.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749916|ref|XP_017156293.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749933|ref|XP_017156294.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749971|ref|XP_017156296.1| PREDICTED: uncharacterized protein LOC108164866 isoform X3
>gi|1036753758|ref|XP_017156517.1| PREDICTED: ras-related protein Rab-21 [Drosophila miranda]
>gi|1036729882|ref|XP_017149826.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036729901|ref|XP_017149833.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036729924|ref|XP_017149842.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036729747|ref|XP_017149758.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729765|ref|XP_017149767.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729784|ref|XP_017149775.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729805|ref|XP_017149784.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729821|ref|XP_017149796.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729842|ref|XP_017149805.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729859|ref|XP_017149813.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729730|ref|XP_017149745.1| PREDICTED: protein stoned-B [Drosophila miranda]
>gi|1036756763|ref|XP_017156689.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036756746|ref|XP_017156688.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036704064|ref|XP_017134668.1| PREDICTED: uncharacterized protein LOC108150871 [Drosophila
>gi|1036705831|ref|XP_017135865.1| PREDICTED: 39 kDa FK506-binding nuclear protein-like [Drosop
>gi|1036700002|ref|XP_017146043.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1036751789|ref|XP_017156396.1| PREDICTED: uncharacterized protein LOC108164921 [Drosophila
>gi|1036751771|ref|XP_017156395.1| PREDICTED: uncharacterized protein LOC108164920 [Drosophila
>gi|1036735990|ref|XP_017153071.1| PREDICTED: protein folded gastrulation [Drosophila miranda]
>gi|1036736007|ref|XP_017153079.1| PREDICTED: protein folded gastrulation [Drosophila miranda]
>gi|1036736023|ref|XP_017153087.1| PREDICTED: protein folded gastrulation [Drosophila miranda]
>gi|1036757551|ref|XP_017156731.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036706245|ref|XP_017136170.1| PREDICTED: uncharacterized protein LOC108151818 [Drosophila
>gi|1036754546|ref|XP_017156564.1| PREDICTED: uncharacterized protein LOC108165038 [Drosophila
>gi|1036754565|ref|XP_017156565.1| PREDICTED: uncharacterized protein LOC108165038 [Drosophila
>gi|1036739929|ref|XP_017155271.1| PREDICTED: uncharacterized protein LOC108164168 isoform X1
>gi|1036739946|ref|XP_017155280.1| PREDICTED: uncharacterized protein LOC108164168 isoform X2
>gi|1036739964|ref|XP_017155291.1| PREDICTED: uncharacterized protein LOC108164181 isoform X1
>gi|1036739982|ref|XP_017155298.1| PREDICTED: uncharacterized protein LOC108164181 isoform X2
>gi|1036740000|ref|XP_017155307.1| PREDICTED: uncharacterized protein LOC108164181 isoform X2
>gi|1036740074|ref|XP_017155347.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
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>gi|1036740096|ref|XP_017155355.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
>gi|1036740114|ref|XP_017155363.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
>gi|1036740132|ref|XP_017155369.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
>gi|1036740150|ref|XP_017155373.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
>gi|1036740164|ref|XP_017155383.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
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>gi|1036740216|ref|XP_017155404.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
>gi|1036740022|ref|XP_017155320.1| PREDICTED: probable isoaspartyl peptidase/L-asparaginase GA2
>gi|1036740056|ref|XP_017155339.1| PREDICTED: UPF0454 protein C12orf49 homolog isoform X2 [Dros
>gi|1036740040|ref|XP_017155331.1| PREDICTED: UPF0454 protein C12orf49 homolog isoform X1 [Dros
>gi|1036715291|ref|XP_017141775.1| PREDICTED: dnaJ homolog subfamily C member 25 homolog [Droso
>gi|1036715255|ref|XP_017141751.1| PREDICTED: neurotrophin receptor-interacting factor homolog
>gi|1036715272|ref|XP_017141763.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036715165|ref|XP_017141706.1| PREDICTED: chondroitin sulfate synthase 1 isoform X1 [Drosop
>gi|1036715184|ref|XP_017141716.1| PREDICTED: chondroitin sulfate synthase 1 isoform X1 [Drosop
>gi|1036715203|ref|XP_017141723.1| PREDICTED: chondroitin sulfate synthase 1 isoform X1 [Drosop
>gi|1036715219|ref|XP_017141732.1| PREDICTED: chondroitin sulfate synthase 3 isoform X2 [Drosop
>gi|1036715237|ref|XP_017141740.1| PREDICTED: chondroitin sulfate synthase 3 isoform X3 [Drosop
>gi|1036733508|ref|XP_017151672.1| PREDICTED: dorsal-ventral patterning protein Sog [Drosophila
>gi|1036754675|ref|XP_017156571.1| PREDICTED: uncharacterized protein LOC108165044 [Drosophila
>gi|1036706604|ref|XP_017136411.1| PREDICTED: uncharacterized protein LOC108151968 [Drosophila
>gi|1036700017|ref|XP_017146158.1| PREDICTED: uncharacterized protein LOC108158376 [Drosophila
>gi|1036700032|ref|XP_017146280.1| PREDICTED: endothelin-converting enzyme-like 1 [Drosophila m
>gi|1036753950|ref|XP_017156529.1| PREDICTED: sex peptide receptor [Drosophila miranda]
>gi|1036700047|ref|XP_017146420.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036751051|ref|XP_017156355.1| PREDICTED: hornerin [Drosophila miranda]
>gi|1036751070|ref|XP_017156356.1| PREDICTED: lysozyme [Drosophila miranda]
>gi|1036733076|ref|XP_017151440.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 16 isoform
>gi|1036733060|ref|XP_017151433.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 16 isoform
>gi|1036711610|ref|XP_017139576.1| PREDICTED: serine/threonine-protein kinase Smg1 isoform X1
>gi|1036711629|ref|XP_017139583.1| PREDICTED: serine/threonine-protein kinase Smg1 isoform X2
>gi|1036711646|ref|XP_017139594.1| PREDICTED: beta-1 adrenergic receptor [Drosophila miranda]
>gi|1036711663|ref|XP_017139604.1| PREDICTED: 40S ribosomal protein S23-like [Drosophila miran
>gi|1036711733|ref|XP_017139642.1| PREDICTED: E3 ubiquitin-protein ligase ZNRF2 [Drosophila mi
>gi|1036711748|ref|XP_017139649.1| PREDICTED: E3 ubiquitin-protein ligase ZNRF2 [Drosophila mi
>gi|1036711680|ref|XP_017139615.1| PREDICTED: TATA element modulatory factor isoform X1 [Droso
>gi|1036711697|ref|XP_017139622.1| PREDICTED: TATA element modulatory factor isoform X2 [Droso
>gi|1036711716|ref|XP_017139632.1| PREDICTED: chitinase-3-like protein 1 [Drosophila miranda]
>gi|1036711761|ref|XP_017139663.1| PREDICTED: tRNA 2'-phosphotransferase 1 [Drosophila miranda]
>gi|1036742934|ref|XP_017155914.1| PREDICTED: zinc finger protein 236 [Drosophila miranda]
>gi|1036700064|ref|XP_017146550.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036706469|ref|XP_017136323.1| PREDICTED: probable serine/threonine-protein kinase mps1 [D
>gi|1036727380|ref|XP_017148366.1| PREDICTED: glutathione synthetase-like isoform X1 [Drosophil
>gi|1036727398|ref|XP_017148376.1| PREDICTED: glutathione synthetase-like isoform X1 [Drosophil
>gi|1036727417|ref|XP_017148383.1| PREDICTED: glutathione synthetase-like isoform X2 [Drosophil
>gi|1036727436|ref|XP_017148392.1| PREDICTED: glutathione synthetase-like isoform X3 [Drosophil
>gi|1036727472|ref|XP_017148404.1| PREDICTED: glutathione synthetase-like isoform X4 [Drosophil
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>gi|1036727489|ref|XP_017148418.1| PREDICTED: LOW QUALITY PROTEIN: mitochondrial fission proces
>gi|1036727507|ref|XP_017148434.1| PREDICTED: hsc70-interacting protein 1-like [Drosophila mira
>gi|1036754331|ref|XP_017156551.1| PREDICTED: uncharacterized protein LOC108165026 [Drosophila
>gi|1036754348|ref|XP_017156552.1| PREDICTED: uncharacterized protein LOC108165026 [Drosophila
>gi|1036723616|ref|XP_017146054.1| PREDICTED: uncharacterized protein LOC108158321 isoform X1
>gi|1036723632|ref|XP_017146061.1| PREDICTED: uncharacterized protein LOC108158321 isoform X2
>gi|1036723650|ref|XP_017146070.1| PREDICTED: putative mediator of RNA polymerase II transcrip
>gi|1036700080|ref|XP_017146676.1| PREDICTED: mucin-19-like [Drosophila miranda]
>gi|1036703230|ref|XP_017156829.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036718884|ref|XP_017143802.1| PREDICTED: malate dehydrogenase-like [Drosophila miranda]
>gi|1036718865|ref|XP_017143788.1| PREDICTED: malate dehydrogenase, mitochondrial-like [Drosopl
>gi|1036718827|ref|XP_017143762.1| PREDICTED: uncharacterized protein LOC108156656 [Drosophila
>gi|1036718846|ref|XP_017143776.1| PREDICTED: mitoguardin [Drosophila miranda]
>gi|1036718904|ref|XP_017143816.1| PREDICTED: uncharacterized protein LOC108156683 [Drosophila
>gi|1036711076|ref|XP_017139312.1| PREDICTED: uncharacterized protein LOC108153694 [Drosophila
>gi|1036711106|ref|XP_017139334.1| PREDICTED: GPI transamidase component PIG-T [Drosophila mir
>gi|1036711085|ref|XP_017139323.1| PREDICTED: trithorax group protein osa [Drosophila miranda]
>gi|1036711121|ref|XP_017139346.1| PREDICTED: CTD nuclear envelope phosphatase 1 homolog isofo
>gi|1036711138|ref|XP_017139353.1| PREDICTED: CTD nuclear envelope phosphatase 1 homolog isofo
>gi|1036736902|ref|XP_017153613.1| PREDICTED: SRA stem-loop-interacting RNA-binding protein, mi
>gi|1036736882|ref|XP_017153600.1| PREDICTED: acyl-CoA-binding domain-containing protein 6 [Dr
>gi|1036736864|ref|XP_017153585.1| PREDICTED: 26S protease regulatory subunit 8 [Drosophila mi
>gi|1036736845|ref|XP_017153572.1| PREDICTED: ATP-binding cassette sub-family A member 2 [Dros
>gi|1036737563|ref|XP_017153983.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofo
>gi|1036737579|ref|XP_017153992.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofo
>gi|1036737598|ref|XP_017154001.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofo
>gi|1036737618|ref|XP_017154008.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofo
>gi|1036755855|ref|XP_017156637.1| PREDICTED: ATP-binding cassette sub-family A member 3 [Dros
>gi|1036739225|ref|XP_017154858.1| PREDICTED: loricrin isoform X1 [Drosophila miranda]
>gi|1036739243|ref|XP_017154866.1| PREDICTED: pro-resilin isoform X2 [Drosophila miranda]
>gi|1036705755|ref|XP_017135808.1| PREDICTED: uncharacterized protein LOC108151618 [Drosophila
>gi|1036730379|ref|XP_017150158.1| PREDICTED: ATP-binding cassette sub-family A member 7 isofo
>gi|1036730342|ref|XP_017150144.1| PREDICTED: ATP-binding cassette sub-family A member 13 isof
>gi|1036730360|ref|XP_017150150.1| PREDICTED: ATP-binding cassette sub-family A member 13 isof
>gi|1036730396|ref|XP_017150171.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1036730413|ref|XP_017150179.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1036730429|ref|XP_017150188.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1036748852|ref|XP_017156233.1| PREDICTED: uncharacterized protein LOC108164823 [Drosophila
>gi|1036748871|ref|XP_017156234.1| PREDICTED: uncharacterized protein LOC108164823 [Drosophila
>gi|1036700097|ref|XP_017146813.1| PREDICTED: uncharacterized protein LOC108158723 [Drosophila
>gi|1036700111|ref|XP_017146945.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036748890|ref|XP_017156235.1| PREDICTED: NEDD8-specific protease 2 [Drosophila miranda]
>gi|1036727782|ref|XP_017148596.1| PREDICTED: blood vessel epicardial substance isoform X3 [Dr
>gi|1036727764|ref|XP_017148587.1| PREDICTED: blood vessel epicardial substance isoform X2 [Dr
>gi|1036727725|ref|XP_017148570.1| PREDICTED: blood vessel epicardial substance isoform X1 [Dr
>gi|1036727747|ref|XP_017148578.1| PREDICTED: blood vessel epicardial substance isoform X1 [Dr
>gi|1036727707|ref|XP_017148558.1| PREDICTED: transcription factor kayak [Drosophila miranda]
>gi|1036735348|ref|XP_017152720.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
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>gi|1036735369|ref|XP_017152733.1| PREDICTED: uncharacterized protein LOC108162488 isoform X1
>gi|1036735388|ref|XP_017152741.1| PREDICTED: uncharacterized protein LOC108162488 isoform X2
>gi|1036744643|ref|XP_017156001.1| PREDICTED: cystathionine beta-synthase [Drosophila miranda]
>gi|1036734725|ref|XP_017152381.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X2
>gi|1036734706|ref|XP_017152371.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X1
>gi|1036734670|ref|XP_017152351.1| PREDICTED: uncharacterized protein LOC108162231 isoform X1
>gi|1036734688|ref|XP_017152360.1| PREDICTED: uncharacterized protein LOC108162231 isoform X2
>gi|1036734742|ref|XP_017152393.1| PREDICTED: probable nuclear transport factor 2 [Drosophila m
>gi|1036739193|ref|XP_017154835.1| PREDICTED: zinc finger protein 134 [Drosophila miranda]
>gi|1036739209|ref|XP_017154848.1| PREDICTED: NAD(P)H-hydrate epimerase [Drosophila miranda]
>gi|1036739173|ref|XP_017154822.1| PREDICTED: RNA-binding protein NOB1 [Drosophila miranda]
>gi|1036720170|ref|XP_017144493.1| PREDICTED: trithorax group protein osa isoform X1 [Drosophil
>gi|1036720189|ref|XP_017144502.1| PREDICTED: collagen alpha-1(I) chain isoform X2 [Drosophila
>gi|1036720209|ref|XP_017144512.1| PREDICTED: collagen alpha-1(I) chain isoform X3 [Drosophila
>gi|1036720266|ref|XP_017144539.1| PREDICTED: uncharacterized protein LOC108157132 isoform X3
>gi|1036720229|ref|XP_017144523.1| PREDICTED: uncharacterized protein LOC108157132 isoform X1
>gi|1036720247|ref|XP_017144531.1| PREDICTED: uncharacterized protein LOC108157132 isoform X2
>gi|1036698274|ref|XP_017138640.1| PREDICTED: putative gustatory receptor 9a [Drosophila mirand
>gi|1036755801|ref|XP_017156634.1| PREDICTED: vitellogenin-1 [Drosophila miranda]
>gi|1036756202|ref|XP_017156656.1| PREDICTED: vitellogenin-2 [Drosophila miranda]
>gi|1036742628|ref|XP_017155897.1| PREDICTED: mucin-5AC isoform X7 [Drosophila miranda]
>gi|1036742610|ref|XP_017155895.1| PREDICTED: mucin-5AC isoform X6 [Drosophila miranda]
>gi|1036742538|ref|XP_017155891.1| PREDICTED: uncharacterized protein LOC108164581 isoform X2
>gi|1036742520|ref|XP_017155890.1| PREDICTED: uncharacterized protein LOC108164581 isoform X1
>gi|1036742592|ref|XP_017155894.1| PREDICTED: mucin-5AC isoform X5 [Drosophila miranda]
>gi|1036742557|ref|XP_017155892.1| PREDICTED: mucin-5AC isoform X3 [Drosophila miranda]
>gi|1036742644|ref|XP_017155898.1| PREDICTED: serine-rich adhesin for platelets isoform X8 [Dro
>gi|1036742576|ref|XP_017155893.1| PREDICTED: mucin-5AC isoform X4 [Drosophila miranda]
>gi|1036739678|ref|XP_017155131.1| PREDICTED: DNA replication ATP-dependent helicase/nuclease
>gi|1036705629|ref|XP_017135713.1| PREDICTED: basic helix-loop-helix transcription factor scle
>gi|1036752664|ref|XP_017156449.1| PREDICTED: keratin, type I cytoskeletal 10 isoform X3 [Drosc
>gi|1036752628|ref|XP_017156447.1| PREDICTED: keratin, type I cytoskeletal 10 isoform X1 [Drosc
>gi|1036752701|ref|XP_017156452.1| PREDICTED: uncharacterized protein LOC108164957 isoform X5
>gi|1036752719|ref|XP_017156453.1| PREDICTED: uncharacterized protein LOC108164957 isoform X5
>gi|1036752645|ref|XP_017156448.1| PREDICTED: uncharacterized protein DDB_G0280205 isoform X2
>gi|1036752682|ref|XP_017156451.1| PREDICTED: uncharacterized protein LOC108164957 isoform X4
>gi|1036752735|ref|XP_017156454.1| PREDICTED: uncharacterized protein LOC108164957 isoform X6
>gi|1036700126|ref|XP_017147072.1| PREDICTED: uncharacterized protein LOC108158833, partial [D
>gi|1036755125|ref|XP_017156598.1| PREDICTED: uncharacterized protein LOC108165061 isoform X2
>gi|1036755087|ref|XP_017156596.1| PREDICTED: uncharacterized protein LOC108165061 isoform X1
>gi|1036755106|ref|XP_017156597.1| PREDICTED: uncharacterized protein LOC108165061 isoform X1
>gi|1036755069|ref|XP_017156595.1| PREDICTED: microfibril-associated glycoprotein 4 isoform X2
>gi|1036755050|ref|XP_017156594.1| PREDICTED: microfibril-associated glycoprotein 4 isoform X1
>gi|1036755014|ref|XP_017156592.1| PREDICTED: fibrinogen-like protein A isoform X1 [Drosophila
>gi|1036755031|ref|XP_017156593.1| PREDICTED: fibrinogen-like protein A isoform X2 [Drosophila
>gi|1036746190|ref|XP_017156091.1| PREDICTED: chitinase-like protein Idgf4 [Drosophila miranda]
>gi|1036700141|ref|XP_017147205.1| PREDICTED: lipoprotein lipase [Drosophila miranda]
>gi|1036756109|ref|XP_017156650.1| PREDICTED: endonuclease G, mitochondrial-like isoform X2 [D
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>gi|1036756090|ref|XP_017156649.1| PREDICTED: endonuclease G, mitochondrial-like isoform X1 [Dr
>gi|1036748145|ref|XP_017156207.1| PREDICTED: uncharacterized protein LOC108164794 [Drosophila
>gi|1036748126|ref|XP_017156205.1| PREDICTED: uncharacterized protein LOC108164793 [Drosophila
>gi|1036703416|ref|XP_017156936.1| PREDICTED: uncharacterized protein LOC108165405 [Drosophila
>gi|1036704401|ref|XP_017134870.1| PREDICTED: uncharacterized protein LOC108151018 [Drosophila
>gi|1036704907|ref|XP_017135229.1| PREDICTED: uncharacterized protein LOC108151237 [Drosophila
>gi|1036703114|ref|XP_017156773.1| PREDICTED: uncharacterized protein LOC108165243 [Drosophila
>gi|1036756583|ref|XP_017156679.1| PREDICTED: PERQ amino acid-rich with GYF domain-containing p
>gi|1036704530|ref|XP_017134954.1| PREDICTED: uncharacterized protein LOC108151079 [Drosophila
>gi|1036754395|ref|XP_017156555.1| PREDICTED: uncharacterized protein LOC108165029 [Drosophila
>gi|1036754412|ref|XP_017156556.1| PREDICTED: uncharacterized protein LOC108165030 [Drosophila
>gi|1036754380|ref|XP_017156554.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1036702984|ref|XP_017156713.1| PREDICTED: uncharacterized protein LOC108165154 isoform X1
>gi|1036703002|ref|XP_017156721.1| PREDICTED: uncharacterized protein LOC108165154 isoform X1
>gi|1036703018|ref|XP_017156728.1| PREDICTED: uncharacterized protein LOC108165154 isoform X2
>gi|1036751844|ref|XP_017156399.1| PREDICTED: carbonic anhydrase-related protein 10 [Drosophila
>gi|1036751880|ref|XP_017156402.1| PREDICTED: N-alpha-acetyltransferase 38, NatC auxiliary subu
>gi|1036751863|ref|XP_017156401.1| PREDICTED: N-alpha-acetyltransferase 38, NatC auxiliary subu
>gi|1036741596|ref|XP_017155839.1| PREDICTED: obg-like ATPase 1 [Drosophila miranda]
>gi|1036741615|ref|XP_017155840.1| PREDICTED: obg-like ATPase 1 [Drosophila miranda]
>gi|1036749185|ref|XP_017156252.1| PREDICTED: transcription factor Sp9 isoform X2 [Drosophila m
>gi|1036749152|ref|XP_017156250.1| PREDICTED: transcription factor Sp9 isoform X1 [Drosophila m
>gi|1036749167|ref|XP_017156251.1| PREDICTED: transcription factor Sp9 isoform X1 [Drosophila m
>gi|1036745200|ref|XP_017156033.1| PREDICTED: transcription factor btd [Drosophila miranda]
>gi|1036720130|ref|XP_017144471.1| PREDICTED: histone acetyltransferase p300 isoform X1 [Droso
>gi|1036720150|ref|XP_017144480.1| PREDICTED: histone acetyltransferase p300 isoform X2 [Droso
>gi|1036754470|ref|XP_017156559.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036724778|ref|XP_017146786.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 9
>gi|1036724743|ref|XP_017146764.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 9
>gi|1036724761|ref|XP_017146775.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 9
>gi|1036731645|ref|XP_017150820.1| PREDICTED: E3 ubiquitin-protein ligase HECW2 isoform X1 [Dr
>gi|1036731681|ref|XP_017150825.1| PREDICTED: E3 ubiquitin-protein ligase HECW2 isoform X3 [Dr
>gi|1036731664|ref|XP_017150821.1| PREDICTED: E3 ubiquitin-protein ligase HECW2 isoform X2 [Dr
>gi|1036750209|ref|XP_017156308.1| PREDICTED: uncharacterized protein LOC108164874 [Drosophila
>gi|1036745509|ref|XP_017156052.1| PREDICTED: lysine--tRNA ligase isoform X1 [Drosophila miran
>gi|1036745525|ref|XP_017156053.1| PREDICTED: lysine--tRNA ligase isoform X2 [Drosophila miran
>gi|1036717368|ref|XP_017142962.1| PREDICTED: probable histone-lysine N-methyltransferase CG17
>gi|1036717387|ref|XP_017142973.1| PREDICTED: fatty acid hydroxylase domain-containing protein
>gi|1036717406|ref|XP_017142985.1| PREDICTED: glutathione S-transferase theta-1 [Drosophila mi
>gi|1036744381|ref|XP_017155989.1| PREDICTED: kinesin-like protein Nod [Drosophila miranda]
>gi|1036744400|ref|XP_017155990.1| PREDICTED: uncharacterized protein LOC108164649 [Drosophila
>gi|1036753741|ref|XP_017156516.1| PREDICTED: nuclease [Drosophila miranda]
>gi|1036753650|ref|XP_017156511.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753668|ref|XP_017156512.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753686|ref|XP_017156513.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753705|ref|XP_017156514.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753724|ref|XP_017156515.1| PREDICTED: uncharacterized protein LOC108164995 isoform X2
>gi|1036724391|ref|XP_017146525.1| PREDICTED: uncharacterized protein LOC108158553 isoform X4
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>gi|1036724356|ref|XP_017146508.1| PREDICTED: uncharacterized protein LOC108158553 isoform X2
>gi|1036724373|ref|XP_017146516.1| PREDICTED: uncharacterized protein LOC108158553 isoform X3
>gi|1036724250|ref|XP_017146459.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724270|ref|XP_017146469.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724286|ref|XP_017146477.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724305|ref|XP_017146487.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724321|ref|XP_017146495.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036742663|ref|XP_017155899.1| PREDICTED: uncharacterized protein LOC108164583 [Drosophila
>gi|1036710962|ref|XP_017139235.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X2
>gi|1036710945|ref|XP_017139228.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X1
>gi|1036710996|ref|XP_017139251.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X4
>gi|1036710979|ref|XP_017139243.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X3
>gi|1036711030|ref|XP_017139276.1| PREDICTED: protein bangles and beads [Drosophila miranda]
>gi|1036711046|ref|XP_017139287.1| PREDICTED: autophagy-related protein 101 [Drosophila miranda
>gi|1036711013|ref|XP_017139265.1| PREDICTED: uncharacterized protein LOC108153661 [Drosophila
>gi|1036711065|ref|XP_017139300.1| PREDICTED: coactosin-like protein [Drosophila miranda]
>gi|1036756147|ref|XP_017156652.1| PREDICTED: gastrin/cholecystokinin type B receptor [Drosophi
>gi|1036705770|ref|XP_017135820.1| PREDICTED: cofilin/actin-depolymerizing factor homolog [Dros
>gi|1036700156|ref|XP_017147331.1| PREDICTED: gastrin/cholecystokinin type B receptor [Drosophi
>gi|1036754941|ref|XP_017156588.1| PREDICTED: probable cytochrome P450 308a1 isoform X1 [Drosop
>gi|1036754959|ref|XP_017156589.1| PREDICTED: probable cytochrome P450 308a1 isoform X2 [Drosop
>gi|1036754978|ref|XP_017156590.1| PREDICTED: probable cytochrome P450 308a1 isoform X3 [Drosop
>gi|1036744903|ref|XP_017156015.1| PREDICTED: cytochrome P450 18a1 [Drosophila miranda]
>gi|1036744886|ref|XP_017156013.1| PREDICTED: cytochrome P450 306a1 [Drosophila miranda]
>gi|1036700171|ref|XP_017147467.1| PREDICTED: zinc metalloproteinase nas-7 [Drosophila miranda]
>gi|1036727073|ref|XP_017148167.1| PREDICTED: serine/threonine-protein kinase fused [Drosophila
>gi|1036727039|ref|XP_017148142.1| PREDICTED: C-mannosyltransferase dpy-19 homolog [Drosophila
>gi|1036727056|ref|XP_017148152.1| PREDICTED: C-mannosyltransferase dpy-19 homolog [Drosophila
>gi|1036746173|ref|XP_017156090.1| PREDICTED: inhibitor of growth protein 3 [Drosophila miranda
>gi|1036728732|ref|XP_017149133.1| PREDICTED: glucose-induced degradation protein 8 homolog [Dr
>gi|1036728713|ref|XP_017149120.1| PREDICTED: nucleoporin NUP53 [Drosophila miranda]
>gi|1036728639|ref|XP_017149086.1| PREDICTED: rab11 family-interacting protein 1 isoform X1 [Dr
>gi|1036728675|ref|XP_017149098.1| PREDICTED: rab11 family-interacting protein 1 isoform X2 [Dr
>gi|1036728694|ref|XP_017149108.1| PREDICTED: uncharacterized protein LOC108159941 isoform X3
>gi|1036746156|ref|XP_017156089.1| PREDICTED: protein valois-like [Drosophila miranda]
>gi|1036742880|ref|XP_017155911.1| PREDICTED: tumor necrosis factor receptor superfamily membei
>gi|1036742899|ref|XP_017155912.1| PREDICTED: tumor necrosis factor receptor superfamily membei
>gi|1036742917|ref|XP_017155913.1| PREDICTED: tumor necrosis factor receptor superfamily membei
>gi|1036753121|ref|XP_017156479.1| PREDICTED: uncharacterized protein LOC108164975 isoform X1
>gi|1036753136|ref|XP_017156480.1| PREDICTED: uncharacterized protein LOC108164975 isoform X1
>gi|1036753153|ref|XP_017156481.1| PREDICTED: uncharacterized protein LOC108164975 isoform X1
>gi|1036753170|ref|XP_017156482.1| PREDICTED: uncharacterized protein LOC108164975 isoform X2
>gi|1036728284|ref|XP_017148876.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036728303|ref|XP_017148899.1| PREDICTED: uncharacterized protein LOC108159829 [Drosophila
>gi|1036728247|ref|XP_017148856.1| PREDICTED: histidine--tRNA ligase, cytoplasmic isoform X2 [I
>gi|1036728265|ref|XP_017148864.1| PREDICTED: histidine--tRNA ligase, cytoplasmic isoform X3 [I
>gi|1036728225|ref|XP_017148847.1| PREDICTED: histidine--tRNA ligase, cytoplasmic isoform X1 [I
>gi|1036752173|ref|XP_017156420.1| PREDICTED: gamma-glutamyltranspeptidase 1 [Drosophila mirand
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>gi|1036738919|ref|XP_017154680.1| PREDICTED: N-acetylgalactosaminyltransferase 7 [Drosophila m
>gi|1036756273|ref|XP_017156660.1| PREDICTED: uncharacterized protein LOC108165112 [Drosophila
>gi|1036756290|ref|XP_017156661.1| PREDICTED: uncharacterized protein LOC108165113 [Drosophila
>gi|1036746772|ref|XP_017156129.1| PREDICTED: uncharacterized protein LOC108164744 isoform X1
>gi|1036746791|ref|XP_017156130.1| PREDICTED: uncharacterized protein LOC108164744 isoform X2
>gi|1036746756|ref|XP_017156128.1| PREDICTED: uncharacterized protein LOC108164743 [Drosophila
>gi|1036746722|ref|XP_017156126.1| PREDICTED: serine protease persephone-like isoform X1 [Droso
>gi|1036746739|ref|XP_017156127.1| PREDICTED: serine protease persephone-like isoform X2 [Droso
>gi|1036746685|ref|XP_017156123.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036746706|ref|XP_017156124.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036700188|ref|XP_017147595.1| PREDICTED: uncharacterized protein LOC108159096 [Drosophila
>gi|1036757062|ref|XP_017156703.1| PREDICTED: uncharacterized protein LOC108165146 [Drosophila
>gi|1036757043|ref|XP_017156702.1| PREDICTED: transcription factor kayak [Drosophila miranda]
>gi|1036703565|ref|XP_017157020.1| PREDICTED: cuticle protein 21.3 isoform X2 [Drosophila miran
>gi|1036703548|ref|XP_017157013.1| PREDICTED: cuticle protein 16.5 isoform X1 [Drosophila miran
>gi|1036733150|ref|XP_017151479.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X1 [Dros
>gi|1036733169|ref|XP_017151487.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X1 [Dros
>gi|1036733204|ref|XP_017151500.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X3 [Dros
>gi|1036733221|ref|XP_017151508.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X4 [Dros
>gi|1036733188|ref|XP_017151494.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X2 [Dros
>gi|1036753777|ref|XP_017156518.1| PREDICTED: transferrin [Drosophila miranda]
>gi|1036739630|ref|XP_017155095.1| PREDICTED: coatomer subunit beta [Drosophila miranda]
>gi|1036736349|ref|XP_017153284.1| PREDICTED: nitric oxide synthase-interacting protein homolog
>gi|1036736333|ref|XP_017153272.1| PREDICTED: protein-serine O-palmitoleoyltransferase porcupin
>gi|1036731200|ref|XP_017150599.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731088|ref|XP_017150547.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731107|ref|XP_017150555.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731127|ref|XP_017150565.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731145|ref|XP_017150573.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731163|ref|XP_017150583.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731181|ref|XP_017150591.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731218|ref|XP_017150608.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731072|ref|XP_017150535.1| PREDICTED: allantoinase [Drosophila miranda]
>gi|1036731015|ref|XP_017150515.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036703264|ref|XP_017156851.1| PREDICTED: uncharacterized protein LOC108165325 [Drosophila
>gi|1036746880|ref|XP_017156135.1| PREDICTED: uncharacterized protein LOC108164749 isoform X2
>gi|1036746863|ref|XP_017156134.1| PREDICTED: uncharacterized protein LOC108164749 isoform X1
>gi|1036705738|ref|XP_017135798.1| PREDICTED: uncharacterized protein LOC108151612 [Drosophila
>gi|1036705969|ref|XP_017135969.1| PREDICTED: uncharacterized protein LOC108151706 [Drosophila
>gi|1036755886|ref|XP_017156638.1| PREDICTED: uncharacterized protein LOC108165097 isoform X1
>gi|1036755905|ref|XP_017156639.1| PREDICTED: uncharacterized protein LOC108165097 isoform X2
>gi|1036745905|ref|XP_017156073.1| PREDICTED: protein Exd1 homolog isoform X1 [Drosophila miran
>gi|1036745923|ref|XP_017156074.1| PREDICTED: protein Exd1 homolog isoform X2 [Drosophila miran
>gi|1036724408|ref|XP_017146537.1| PREDICTED: SWI5-dependent HO expression protein 3-like [Dros
>gi|1036701212|ref|XP_017154222.1| PREDICTED: uncharacterized protein LOC108163424 [Drosophila
>gi|1036701229|ref|XP_017154347.1| PREDICTED: TPR-containing protein DDB_G0280363 [Drosophila m
>gi|1036758205|ref|XP_017156763.1| PREDICTED: cytochrome c oxidase subunit 4 isoform 2, mitocho
>gi|1036816707|ref|XP_017137410.1| PREDICTED: protein yippee-like CG15309 [Drosophila miranda]
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>gi|1036816726|ref|XP_017137411.1| PREDICTED: protein yippee-like CG15309 [Drosophila miranda]
>gi|1036816649|ref|XP_017137406.1| PREDICTED: kelch domain-containing protein 3 isoform X1 [Dro
>gi|1036816667|ref|XP_017137408.1| PREDICTED: kelch domain-containing protein 3 isoform X2 [Dro
>gi|1036816688|ref|XP_017137409.1| PREDICTED: kelch domain-containing protein 3 isoform X2 [Dro
>gi|1036816745|ref|XP_017137412.1| PREDICTED: INO80 complex subunit C [Drosophila miranda]
>gi|1036830533|ref|XP_017138182.1| PREDICTED: mannosyl-oligosaccharide glucosidase [Drosophila
>gi|1036799288|ref|XP_017136446.1| PREDICTED: transmembrane protein 68 [Drosophila miranda]
>gi|1036799253|ref|XP_017136444.1| PREDICTED: transport and Golgi organization protein 2 isofor
>gi|1036799271|ref|XP_017136445.1| PREDICTED: transport and Golgi organization protein 2 isofor
>gi|1036799234|ref|XP_017136443.1| PREDICTED: RRP12-like protein [Drosophila miranda]
>gi|1036784211|ref|XP_017135592.1| PREDICTED: uncharacterized protein LOC108151473 isoform X3
>gi|1036784247|ref|XP_017135594.1| PREDICTED: uncharacterized protein LOC108151473 isoform X5
>gi|1036784192|ref|XP_017135591.1| PREDICTED: uncharacterized protein LOC108151473 isoform X2
>gi|1036784173|ref|XP_017135590.1| PREDICTED: uncharacterized protein LOC108151473 isoform X1
>gi|1036784230|ref|XP_017135593.1| PREDICTED: uncharacterized protein LOC108151473 isoform X4
>gi|1036832323|ref|XP_017138279.1| PREDICTED: DNA ligase 4 [Drosophila miranda]
>gi|1036838628|ref|XP_017138644.1| PREDICTED: ribonuclease H2 subunit B [Drosophila miranda]
>gi|1036832475|ref|XP_017138289.1| PREDICTED: uncharacterized protein LOC108153041 isoform X1
>gi|1036832495|ref|XP_017138290.1| PREDICTED: uncharacterized protein LOC108153041 isoform X2
>gi|1036827231|ref|XP_017137990.1| PREDICTED: fatty acid hydroxylase domain-containing protein
>gi|1036827212|ref|XP_017137989.1| PREDICTED: RNA-binding protein FUS [Drosophila miranda]
>gi|1036827248|ref|XP_017137991.1| PREDICTED: selenoprotein BthD [Drosophila miranda]
>gi|1036795987|ref|XP_017136259.1| PREDICTED: actin cytoskeleton-regulatory complex protein pan
>gi|1036796006|ref|XP_017136260.1| PREDICTED: actin cytoskeleton-regulatory complex protein pan
>gi|1036795931|ref|XP_017136257.1| PREDICTED: neuroglian isoform X1 [Drosophila miranda]
>gi|1036795968|ref|XP_017136258.1| PREDICTED: neuroglian isoform X2 [Drosophila miranda]
>gi|1036826162|ref|XP_017137928.1| PREDICTED: integrator complex subunit 4 [Drosophila miranda]
>gi|1036775700|ref|XP_017135090.1| PREDICTED: uncharacterized protein LOC108151171 isoform X2
>gi|1036775683|ref|XP_017135089.1| PREDICTED: uncharacterized protein LOC108151171 isoform X1
>gi|1036775719|ref|XP_017135091.1| PREDICTED: uncharacterized protein LOC108151171 isoform X3
>gi|1036775740|ref|XP_017135092.1| PREDICTED: uncharacterized protein LOC108151171 isoform X4
>gi|1036775664|ref|XP_017135088.1| PREDICTED: LOW QUALITY PROTEIN: E3 ubiquitin-protein ligase
>gi|1036758220|ref|XP_017156764.1| PREDICTED: uncharacterized protein LOC108165236 [Drosophila
>gi|1036832871|ref|XP_017138312.1| PREDICTED: uncharacterized protein LOC108153050 [Drosophila
>gi|1036832836|ref|XP_017138310.1| PREDICTED: uncharacterized protein LOC108153049 [Drosophila
>gi|1036832854|ref|XP_017138311.1| PREDICTED: uncharacterized protein LOC108153049 [Drosophila
>gi|1036832817|ref|XP_017138309.1| PREDICTED: zinc finger C4H2 domain-containing protein [Droso
>gi|1036812587|ref|XP_017137180.1| PREDICTED: dnaJ homolog subfamily C member 22 [Drosophila mi
>gi|1036812606|ref|XP_017137181.1| PREDICTED: dnaJ homolog subfamily C member 22 [Drosophila mi
>gi|1036812549|ref|XP_017137178.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036812568|ref|XP_017137179.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036812625|ref|XP_017137183.1| PREDICTED: homeotic protein female sterile-like [Drosophila
>gi|1036812644|ref|XP_017137184.1| PREDICTED: homeotic protein female sterile-like [Drosophila
>gi|1036812662|ref|XP_017137185.1| PREDICTED: uncharacterized protein LOC108152395 isoform X1
>gi|1036812683|ref|XP_017137186.1| PREDICTED: uncharacterized protein LOC108152395 isoform X2
>gi|1036838429|ref|XP_017138629.1| PREDICTED: uncharacterized protein LOC108153271 [Drosophila
>gi|1036789975|ref|XP_017135921.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC9 [Dro
>gi|1036789919|ref|XP_017135918.1| PREDICTED: density-regulated protein homolog [Drosophila mir
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>gi|1036789883|ref|XP_017135916.1| PREDICTED: nitrogen permease regulator 2-like protein [Droso
>gi|1036789901|ref|XP_017135917.1| PREDICTED: ketimine reductase mu-crystallin [Drosophila mira
>gi|1036789938|ref|XP_017135919.1| PREDICTED: uncharacterized protein LOC108151686 isoform X1
>gi|1036789956|ref|XP_017135920.1| PREDICTED: uncharacterized protein LOC108151686 isoform X2
>gi|1036789831|ref|XP_017135913.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036789850|ref|XP_017135914.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036789867|ref|XP_017135915.1| PREDICTED: regulator of G-protein signaling 7 [Drosophila mi
>gi|1036839647|ref|XP_017138697.1| PREDICTED: uncharacterized protein LOC108153304 isoform X1
>gi|1036839662|ref|XP_017138698.1| PREDICTED: uncharacterized protein LOC108153304 isoform X2
>gi|1036839680|ref|XP_017138699.1| PREDICTED: uncharacterized protein LOC108153304 isoform X2
>gi|1036778909|ref|XP_017135282.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036778776|ref|XP_017135273.1| PREDICTED: uncharacterized protein LOC108151262 isoform X5
>gi|1036778758|ref|XP_017135272.1| PREDICTED: uncharacterized protein LOC108151262 isoform X4
>gi|1036778704|ref|XP_017135269.1| PREDICTED: uncharacterized protein LOC108151262 isoform X1
>gi|1036778740|ref|XP_017135271.1| PREDICTED: uncharacterized protein LOC108151262 isoform X3
>gi|1036778722|ref|XP_017135270.1| PREDICTED: uncharacterized protein LOC108151262 isoform X2
>gi|1036778855|ref|XP_017135278.1| PREDICTED: LOW QUALITY PROTEIN: proteasome inhibitor PI31 su
>gi|1036778796|ref|XP_017135274.1| PREDICTED: uncharacterized protein LOC108151264 [Drosophila
>gi|1036778815|ref|XP_017135275.1| PREDICTED: uncharacterized protein LOC108151265 [Drosophila
>gi|1036778834|ref|XP_017135277.1| PREDICTED: uncharacterized protein LOC108151266 [Drosophila
>gi|1036778874|ref|XP_017135279.1| PREDICTED: 40S ribosomal protein S19a-like [Drosophila mira
>gi|1036778892|ref|XP_017135280.1| PREDICTED: uncharacterized protein LOC108151269 [Drosophila
>gi|1036767279|ref|XP_017134623.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036851147|ref|XP_017139330.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036767393|ref|XP_017134627.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036845649|ref|XP_017139012.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036845667|ref|XP_017139014.1| PREDICTED: 60 kDa heat shock protein, mitochondrial-like [Dr
>gi|1036758238|ref|XP_017156765.1| PREDICTED: 60 kDa heat shock protein, mitochondrial-like [Dr
>gi|1036845632|ref|XP_017139011.1| PREDICTED: calcitonin gene-related peptide type 1 receptor
>gi|1036843301|ref|XP_017138895.1| PREDICTED: ELAV-like protein 2 isoform X2 [Drosophila mirand
>gi|1036843283|ref|XP_017138894.1| PREDICTED: ELAV-like protein 1 isoform X1 [Drosophila mirand
>gi|1036838410|ref|XP_017138627.1| PREDICTED: protein YIPF1 [Drosophila miranda]
>gi|1036835152|ref|XP_017138433.1| PREDICTED: breast cancer metastasis-suppressor 1 homolog [Dr
>gi|1036835186|ref|XP_017138435.1| PREDICTED: probable 39S ribosomal protein L49, mitochondrial
>gi|1036835167|ref|XP_017138434.1| PREDICTED: probable 39S ribosomal protein L49, mitochondrial
>gi|1036849963|ref|XP_017139262.1| PREDICTED: uncharacterized protein LOC108153662 isoform X1
>gi|1036849983|ref|XP_017139263.1| PREDICTED: uncharacterized protein LOC108153662 isoform X2
>gi|1036827395|ref|XP_017138001.1| PREDICTED: FAD synthase [Drosophila miranda]
>gi|1036827413|ref|XP_017138002.1| PREDICTED: FAD synthase [Drosophila miranda]
>gi|1036827378|ref|XP_017137999.1| PREDICTED: aarF domain-containing protein kinase 4 [Drosophi
>gi|1036811059|ref|XP_017137099.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036811019|ref|XP_017137097.1| PREDICTED: uncharacterized protein LOC108152344 [Drosophila
>gi|1036811038|ref|XP_017137098.1| PREDICTED: neprilysin-11 [Drosophila miranda]
>gi|1036839828|ref|XP_017138707.1| PREDICTED: uncharacterized protein LOC108153308 [Drosophila
>gi|1036839846|ref|XP_017138708.1| PREDICTED: uncharacterized protein LOC108153308 [Drosophila
>gi|1036842877|ref|XP_017138871.1| PREDICTED: polyhomeotic-proximal chromatin protein-like [Dro
>gi|1036758253|ref|XP_017156766.1| PREDICTED: bromodomain-containing factor 1-like [Drosophila
>gi|1036790512|ref|XP_017135950.1| PREDICTED: LOW QUALITY PROTEIN: polyhomeotic-proximal chroma
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>gi|1036790419|ref|XP_017135945.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790435|ref|XP_017135946.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790455|ref|XP_017135947.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790474|ref|XP_017135948.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790493|ref|XP_017135949.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036812222|ref|XP_017137160.1| PREDICTED: 6-phosphogluconate dehydrogenase, decarboxylating
>gi|1036812297|ref|XP_017137164.1| PREDICTED: protein bcn92 [Drosophila miranda]
>gi|1036812239|ref|XP_017137161.1| PREDICTED: MAP kinase-activated protein kinase 2 [Drosophila
>gi|1036812260|ref|XP_017137162.1| PREDICTED: MAP kinase-activated protein kinase 2 [Drosophila
>gi|1036812279|ref|XP_017137163.1| PREDICTED: MAP kinase-activated protein kinase 2 [Drosophila
>gi|1036800012|ref|XP_017136486.1| PREDICTED: N-acylneuraminate-9-phosphatase [Drosophila miran
>gi|1036800050|ref|XP_017136488.1| PREDICTED: protein lin-52 homolog [Drosophila miranda]
>gi|1036800032|ref|XP_017136487.1| PREDICTED: alpha-protein kinase 1 [Drosophila miranda]
>gi|1036799994|ref|XP_017136485.1| PREDICTED: regulator of telomere elongation helicase 1 homol
>gi|1036805859|ref|XP_017136816.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805878|ref|XP_017136818.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805897|ref|XP_017136819.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805916|ref|XP_017136820.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805934|ref|XP_017136821.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805951|ref|XP_017136822.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036784600|ref|XP_017135614.1| PREDICTED: sorting nexin-27 isoform X1 [Drosophila miranda]
>gi|1036784638|ref|XP_017135616.1| PREDICTED: uncharacterized protein LOC108151488 [Drosophila
>gi|1036784657|ref|XP_017135618.1| PREDICTED: 60S ribosomal protein L28 [Drosophila miranda]
>gi|1036784621|ref|XP_017135615.1| PREDICTED: sorting nexin-27 isoform X2 [Drosophila miranda]
>gi|1036784485|ref|XP_017135608.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784506|ref|XP_017135609.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784525|ref|XP_017135610.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784544|ref|XP_017135611.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784563|ref|XP_017135612.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784449|ref|XP_017135606.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784412|ref|XP_017135604.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784430|ref|XP_017135605.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784466|ref|XP_017135607.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784582|ref|XP_017135613.1| PREDICTED: NADH-ubiquinone oxidoreductase 75 kDa subunit, m
>gi|1036758272|ref|XP_017156767.1| PREDICTED: protein trapped in endoderm-1-like, partial [Dros
>gi|1036819313|ref|XP_017137554.1| PREDICTED: thioredoxin reductase 1, mitochondrial isoform X
>gi|1036819294|ref|XP_017137553.1| PREDICTED: thioredoxin reductase 1, mitochondrial isoform X
>gi|1036819332|ref|XP_017137555.1| PREDICTED: C-factor isoform X1 [Drosophila miranda]
>gi|1036819351|ref|XP_017137556.1| PREDICTED: C-factor isoform X2 [Drosophila miranda]
>gi|1036819374|ref|XP_017137557.1| PREDICTED: uncharacterized protein LOC108152606 [Drosophila
>gi|1036837757|ref|XP_017138589.1| PREDICTED: troponin I isoform X1 [Drosophila miranda]
>gi|1036837813|ref|XP_017138592.1| PREDICTED: troponin I isoform X4 [Drosophila miranda]
>gi|1036837795|ref|XP_017138591.1| PREDICTED: troponin I isoform X3 [Drosophila miranda]
>gi|1036837776|ref|XP_017138590.1| PREDICTED: troponin I isoform X2 [Drosophila miranda]
>gi|1036837868|ref|XP_017138595.1| PREDICTED: troponin I isoform X7 [Drosophila miranda]
>gi|1036837922|ref|XP_017138599.1| PREDICTED: troponin I isoform X10 [Drosophila miranda]
>gi|1036837903|ref|XP_017138598.1| PREDICTED: troponin I isoform X9 [Drosophila miranda]
>gi|1036837885|ref|XP_017138597.1| PREDICTED: troponin I isoform X8 [Drosophila miranda]
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>gi|1036837831|ref|XP_017138593.1| PREDICTED: troponin I isoform X5 [Drosophila miranda]
>gi|1036837850|ref|XP_017138594.1| PREDICTED: troponin I isoform X6 [Drosophila miranda]
>gi|1036837941|ref|XP_017138600.1| PREDICTED: troponin I isoform X11 [Drosophila miranda]
>gi|1036837960|ref|XP_017138601.1| PREDICTED: troponin I isoform X12 [Drosophila miranda]
>gi|1036848353|ref|XP_017139165.1| PREDICTED: casein kinase II subunit beta'-like [Drosophila m
>gi|1036808771|ref|XP_017136975.1| PREDICTED: interference hedgehog isoform X2 [Drosophila mira
>gi|1036808655|ref|XP_017136969.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808674|ref|XP_017136970.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808694|ref|XP_017136971.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808713|ref|XP_017136972.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808732|ref|XP_017136973.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808753|ref|XP_017136974.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808788|ref|XP_017136976.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808806|ref|XP_017136977.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808827|ref|XP_017136978.1| PREDICTED: regulatory protein zeste [Drosophila miranda]
>gi|1036803920|ref|XP_017136705.1| PREDICTED: basement membrane-specific heparan sulfate proteo
>gi|1036803937|ref|XP_017136706.1| PREDICTED: basement membrane-specific heparan sulfate proteo
>gi|1036846112|ref|XP_017139040.1| PREDICTED: PDF receptor [Drosophila miranda]
>gi|1036850930|ref|XP_017139318.1| PREDICTED: probable V-type proton ATPase subunit D 2 [Droso
>gi|1036828261|ref|XP_017138050.1| PREDICTED: dopamine N-acetyltransferase isoform X1 [Drosophi
>gi|1036828280|ref|XP_017138052.1| PREDICTED: dopamine N-acetyltransferase isoform X2 [Drosophi
>gi|1036828299|ref|XP_017138053.1| PREDICTED: dopamine N-acetyltransferase isoform X2 [Drosophi
>gi|1036828338|ref|XP_017138055.1| PREDICTED: protein GUCD1 isoform X2 [Drosophila miranda]
>gi|1036828319|ref|XP_017138054.1| PREDICTED: protein GUCD1 isoform X1 [Drosophila miranda]
>gi|1036828243|ref|XP_017138049.1| PREDICTED: zinc finger and BTB domain-containing protein 49
>gi|1036833065|ref|XP_017138323.1| PREDICTED: PHD finger protein 12 [Drosophila miranda]
>gi|1036815826|ref|XP_017137357.1| PREDICTED: uncharacterized protein LOC108152492 [Drosophila
>gi|1036815807|ref|XP_017137356.1| PREDICTED: uncharacterized protein LOC108152491 [Drosophila
>gi|1036815784|ref|XP_017137354.1| PREDICTED: mitochondrial ribonuclease P protein 3 [Drosophil
>gi|1036815763|ref|XP_017137353.1| PREDICTED: DNA repair endonuclease XPF [Drosophila miranda]
>gi|1036851487|ref|XP_017139351.1| PREDICTED: uncharacterized protein LOC108153733 [Drosophila
>gi|1036767666|ref|XP_017134640.1| PREDICTED: uncharacterized protein LOC108150845 [Drosophila
>gi|1036758291|ref|XP_017156768.1| PREDICTED: uncharacterized protein LOC108165242 [Drosophila
>gi|1036765327|ref|XP_017157164.1| PREDICTED: uncharacterized protein LOC108165601 [Drosophila
>gi|1036769627|ref|XP_017134750.1| PREDICTED: uncharacterized protein LOC108150931 [Drosophila
>gi|1036766542|ref|XP_017134580.1| PREDICTED: uncharacterized protein LOC108150789 isoform X2
>gi|1036766523|ref|XP_017134579.1| PREDICTED: uncharacterized protein LOC108150789 isoform X1
>gi|1036852340|ref|XP_017139405.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036763406|ref|XP_017157064.1| PREDICTED: translation initiation factor IF-2 [Drosophila m
>gi|1036851345|ref|XP_017139342.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1036766958|ref|XP_017134604.1| PREDICTED: uncharacterized protein LOC108150810 [Drosophila
>gi|1036797845|ref|XP_017136363.1| PREDICTED: fasciclin-2 isoform X1 [Drosophila miranda]
>gi|1036797879|ref|XP_017136365.1| PREDICTED: fasciclin-2 isoform X3 [Drosophila miranda]
>gi|1036797862|ref|XP_017136364.1| PREDICTED: fasciclin-2 isoform X2 [Drosophila miranda]
>gi|1036797898|ref|XP_017136366.1| PREDICTED: fasciclin-2 isoform X4 [Drosophila miranda]
>gi|1036797917|ref|XP_017136367.1| PREDICTED: fasciclin-2 isoform X5 [Drosophila miranda]
>gi|1036797936|ref|XP_017136369.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucuro
>gi|1036846146|ref|XP_017139041.1| PREDICTED: translation machinery-associated protein 7 homolo
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>gi|1036827266|ref|XP_017137993.1| PREDICTED: mitochondrial tRNA-specific 2-thiouridylase 1 [Di
>gi|1036827284|ref|XP_017137994.1| PREDICTED: monocyte to macrophage differentiation factor [Di
>gi|1036827303|ref|XP_017137995.1| PREDICTED: monocyte to macrophage differentiation factor [Di
>gi|1036801742|ref|XP_017136587.1| PREDICTED: HEAT repeat-containing protein 6 [Drosophila mira
>gi|1036801760|ref|XP_017136588.1| PREDICTED: retinoblastoma family protein [Drosophila miranda
>gi|1036852545|ref|XP_017139417.1| PREDICTED: uncharacterized protein LOC108153784 [Drosophila
>gi|1036758307|ref|XP_017156769.1| PREDICTED: ESX-1 secretion-associated protein EspI [Drosophi
>gi|1036791737|ref|XP_017136020.1| PREDICTED: vesicle-trafficking protein SEC22b-B [Drosophila
>gi|1036791635|ref|XP_017136013.1| PREDICTED: uncharacterized protein CG43867 isoform X7 [Droso
>gi|1036791652|ref|XP_017136014.1| PREDICTED: uncharacterized protein CG43867 isoform X8 [Droso
>gi|1036791616|ref|XP_017136012.1| PREDICTED: uncharacterized protein CG43867 isoform X6 [Droso
>gi|1036791579|ref|XP_017136010.1| PREDICTED: uncharacterized protein CG43867 isoform X4 [Droso
>gi|1036791539|ref|XP_017136008.1| PREDICTED: uncharacterized protein CG43867 isoform X2 [Droso
>gi|1036791558|ref|XP_017136009.1| PREDICTED: uncharacterized protein CG43867 isoform X3 [Droso
>gi|1036791520|ref|XP_017136007.1| PREDICTED: uncharacterized protein CG43867 isoform X1 [Droso
>gi|1036791598|ref|XP_017136011.1| PREDICTED: uncharacterized protein CG43867 isoform X5 [Droso
>gi|1036791671|ref|XP_017136015.1| PREDICTED: uncharacterized protein CG43867 isoform X9 [Droso
>gi|1036791703|ref|XP_017136016.1| PREDICTED: uncharacterized protein CG43867 isoform X10 [Dros
>gi|1036791720|ref|XP_017136017.1| PREDICTED: uncharacterized protein CG43867 isoform X11 [Dros
>gi|1036791774|ref|XP_017136022.1| PREDICTED: uncharacterized protein LOC108151733 [Drosophila
>gi|1036791755|ref|XP_017136021.1| PREDICTED: uncharacterized protein LOC108151732 [Drosophila
>gi|1036846346|ref|XP_017139052.1| PREDICTED: protein lunapark-B-like isoform X2 [Drosophila mi
>gi|1036846363|ref|XP_017139053.1| PREDICTED: protein lunapark-B-like isoform X3 [Drosophila mi
>gi|1036846327|ref|XP_017139051.1| PREDICTED: protein lunapark-B-like isoform X1 [Drosophila mi
>gi|1036827152|ref|XP_017137986.1| PREDICTED: leucine-zipper-like transcriptional regulator 1
>gi|1036843707|ref|XP_017138920.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036843688|ref|XP_017138919.1| PREDICTED: nucleosome assembly protein 1-like 1-A [Drosophil
>gi|1036824926|ref|XP_017137860.1| PREDICTED: uncharacterized protein LOC108152787 [Drosophila
>gi|1036824944|ref|XP_017137861.1| PREDICTED: cell wall integrity and stress response component
>gi|1036824963|ref|XP_017137862.1| PREDICTED: translocating chain-associated membrane protein
>gi|1036842197|ref|XP_017138834.1| PREDICTED: crossover junction endonuclease MUS81 [Drosophila
>gi|1036842181|ref|XP_017138833.1| PREDICTED: LOW QUALITY PROTEIN: RUN domain-containing prote
>gi|1036846681|ref|XP_017139072.1| PREDICTED: 3-oxoacyl-[acyl-carrier-protein] reductase FabG
>gi|1036846658|ref|XP_017139071.1| PREDICTED: LOW QUALITY PROTEIN: synaptic vesicle glycoprote
>gi|1036758325|ref|XP_017156770.1| PREDICTED: NADH-quinone oxidoreductase subunit F 1-like [Dr
>gi|1036758343|ref|XP_017156771.1| PREDICTED: NADH-quinone oxidoreductase subunit G-like [Dros
>gi|1036846381|ref|XP_017139054.1| PREDICTED: uncharacterized protein LOC108153509 [Drosophila
>gi|1036833321|ref|XP_017138338.1| PREDICTED: 60S ribosomal protein L13a-like [Drosophila mira
>gi|1036758359|ref|XP_017156774.1| PREDICTED: ERI1 exoribonuclease 2-like [Drosophila miranda]
>gi|1036765347|ref|XP_017157165.1| PREDICTED: uncharacterized protein DDB_G0283357 [Drosophila
>gi|1036821039|ref|XP_017137644.1| PREDICTED: probable calcium-binding protein CML12 [Drosophil
>gi|1036821057|ref|XP_017137645.1| PREDICTED: probable calcium-binding protein CML12 [Drosophil
>gi|1036821000|ref|XP_017137641.1| PREDICTED: adenylate cyclase, terminal-differentiation speci
>gi|1036821113|ref|XP_017137647.1| PREDICTED: uncharacterized protein LOC108152662 [Drosophila
>gi|1036820981|ref|XP_017137640.1| PREDICTED: cell division control protein 45 homolog [Drosop
>gi|1036821094|ref|XP_017137646.1| PREDICTED: ribonuclease P protein subunit p21 [Drosophila mi
>gi|1036821018|ref|XP_017137643.1| PREDICTED: gamma-butyrobetaine dioxygenase [Drosophila miran
>gi|1036824082|ref|XP_017137814.1| PREDICTED: protein suppressor of white apricot isoform X1 [
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>gi|1036824101|ref|XP_017137815.1| PREDICTED: protein suppressor of white apricot isoform X2 [
>gi|1036824136|ref|XP_017137817.1| PREDICTED: protein suppressor of white apricot isoform X3 [
>gi|1036824155|ref|XP_017137818.1| PREDICTED: protein suppressor of white apricot isoform X4 [
>gi|1036767334|ref|XP_017134624.1| PREDICTED: chymotrypsin-elastase inhibitor ixodidin [Drosopl
>gi|1036848879|ref|XP_017139196.1| PREDICTED: 27 kDa hemolymph protein [Drosophila miranda]
>gi|1036765926|ref|XP_017134548.1| PREDICTED: uncharacterized protein LOC108150765 [Drosophila
>gi|1036852071|ref|XP_017139388.1| PREDICTED: succinyl-CoA ligase subunit alpha, mitochondrial
>gi|1036812434|ref|XP_017137172.1| PREDICTED: uncharacterized protein LOC108152386 [Drosophila
>gi|1036812491|ref|XP_017137175.1| PREDICTED: uncharacterized protein LOC108152388 [Drosophila
>gi|1036812452|ref|XP_017137173.1| PREDICTED: uncharacterized protein LOC108152387 isoform X1
>gi|1036812470|ref|XP_017137174.1| PREDICTED: uncharacterized protein LOC108152387 isoform X2
>gi|1036765745|ref|XP_017134537.1| PREDICTED: uncharacterized protein LOC108150754 [Drosophila
>gi|1036758375|ref|XP_017156775.1| PREDICTED: pollen-specific leucine-rich repeat extensin-like
>gi|1036764635|ref|XP_017157127.1| PREDICTED: uncharacterized protein LOC108165564 [Drosophila
>gi|1036849174|ref|XP_017139213.1| PREDICTED: 27 kDa hemolymph protein [Drosophila miranda]
>gi|1036765996|ref|XP_017134551.1| PREDICTED: uncharacterized protein LOC108150768 [Drosophila
>gi|1036765975|ref|XP_017134550.1| PREDICTED: uncharacterized protein LOC108150767 [Drosophila
>gi|1036770401|ref|XP_017134793.1| PREDICTED: probable basic-leucine zipper transcription facto
>gi|1036758394|ref|XP_017156776.1| PREDICTED: uncharacterized protein LOC108165249 [Drosophila
>gi|1036852162|ref|XP_017139394.1| PREDICTED: alpha-protein kinase 1 [Drosophila miranda]
>gi|1036758413|ref|XP_017156777.1| PREDICTED: uncharacterized protein LOC108165250 [Drosophila
>gi|1036852090|ref|XP_017139390.1| PREDICTED: GATA zinc finger domain-containing protein 10 [Dr
>gi|1036758432|ref|XP_017156778.1| PREDICTED: uncharacterized protein LOC108165251 [Drosophila
>gi|1036764243|ref|XP_017157110.1| PREDICTED: hornerin [Drosophila miranda]
>gi|1036846417|ref|XP_017139056.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1036801064|ref|XP_017136546.1| PREDICTED: LOW QUALITY PROTEIN: polyhomeotic-proximal chroma
>gi|1036835060|ref|XP_017138428.1| PREDICTED: probable ATP-dependent DNA helicase DDX11 [Drosop
>gi|1036832108|ref|XP_017138270.1| PREDICTED: activating transcription factor 3 [Drosophila mi
>gi|1036764738|ref|XP_017157133.1| PREDICTED: calcineurin B homologous protein 1 [Drosophila m
>gi|1036758451|ref|XP_017156779.1| PREDICTED: uncharacterized protein LOC108165252 [Drosophila
>gi|1036850799|ref|XP_017139310.1| PREDICTED: uncharacterized protein LOC108153696 [Drosophila
>gi|1036796526|ref|XP_017136288.1| PREDICTED: disheveled-associated activator of morphogenesis
>gi|1036796507|ref|XP_017136287.1| PREDICTED: disheveled-associated activator of morphogenesis
>gi|1036796545|ref|XP_017136289.1| PREDICTED: golgin subfamily A member 6C-like [Drosophila mi
>gi|1036849821|ref|XP_017139253.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036831922|ref|XP_017138259.1| PREDICTED: N-alpha-acetyltransferase 30 isoform X1 [Drosoph
>gi|1036831941|ref|XP_017138260.1| PREDICTED: N-alpha-acetyltransferase 30 isoform X2 [Drosoph
>gi|1036837347|ref|XP_017138563.1| PREDICTED: poly(A) RNA polymerase, mitochondrial [Drosophila
>gi|1036837366|ref|XP_017138565.1| PREDICTED: CD9 antigen [Drosophila miranda]
>gi|1036807353|ref|XP_017136897.1| PREDICTED: solute carrier family 12 member 8 [Drosophila mi
>gi|1036807335|ref|XP_017136896.1| PREDICTED: ESF1 homolog [Drosophila miranda]
>gi|1036807389|ref|XP_017136898.1| PREDICTED: serine/threonine-protein kinase Nek8 [Drosophila
>gi|1036767411|ref|XP_017134628.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036811422|ref|XP_017137116.1| PREDICTED: sex-lethal homolog isoform X2 [Drosophila miranda
>gi|1036811403|ref|XP_017137115.1| PREDICTED: sex-lethal homolog isoform X1 [Drosophila miranda
>gi|1036811441|ref|XP_017137117.1| PREDICTED: RNA-binding protein FUS-like [Drosophila miranda]
>gi|1036801929|ref|XP_017136598.1| PREDICTED: 5'-AMP-activated protein kinase catalytic subunit
>gi|1036801949|ref|XP_017136599.1| PREDICTED: thioredoxin, mitochondrial [Drosophila miranda]
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>gi|1036801781|ref|XP_017136589.1| PREDICTED: rho GTPase-activating protein gacZ isoform X1 [Di
>gi|1036801800|ref|XP_017136590.1| PREDICTED: rho GTPase-activating protein gacZ isoform X2 [Di
>gi|1036801817|ref|XP_017136591.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [Di
>gi|1036801836|ref|XP_017136592.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [Di
>gi|1036801855|ref|XP_017136593.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [Di
>gi|1036801874|ref|XP_017136594.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [Di
>gi|1036801893|ref|XP_017136595.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [Di
>gi|1036801911|ref|XP_017136597.1| PREDICTED: rho GTPase-activating protein gacZ isoform X4 [Di
>gi|1036830077|ref|XP_017138153.1| PREDICTED: RILP-like protein homolog isoform X2 [Drosophila
>gi|1036830058|ref|XP_017138152.1| PREDICTED: RILP-like protein homolog isoform X1 [Drosophila
>gi|1036825259|ref|XP_017137880.1| PREDICTED: LOW QUALITY PROTEIN: microtubule-associated prote
>gi|1036758470|ref|XP_017156780.1| PREDICTED: putative gustatory receptor 2a [Drosophila mirand
>gi|1036849335|ref|XP_017139222.1| PREDICTED: uncharacterized protein LOC108153629 [Drosophila
>gi|1036849349|ref|XP_017139223.1| PREDICTED: uncharacterized protein LOC108153630 [Drosophila
>gi|1036826589|ref|XP_017137953.1| PREDICTED: leucine-rich PPR motif-containing protein, mitoch
>gi|1036834104|ref|XP_017138374.1| PREDICTED: enoyl-CoA delta isomerase 2, mitochondrial [Dros
>gi|1036834085|ref|XP_017138373.1| PREDICTED: large subunit GTPase 1 homolog [Drosophila mirand
>gi|1036783797|ref|XP_017135569.1| PREDICTED: GDP-fucose protein O-fucosyltransferase 2 [Drosop
>gi|1036784078|ref|XP_017135585.1| PREDICTED: PXMP2/4 family protein 4 [Drosophila miranda]
>gi|1036784097|ref|XP_017135586.1| PREDICTED: PXMP2/4 family protein 4 [Drosophila miranda]
>gi|1036783816|ref|XP_017135570.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036784021|ref|XP_017135582.1| PREDICTED: protein sym1 isoform X1 [Drosophila miranda]
>gi|1036784040|ref|XP_017135583.1| PREDICTED: protein sym1 isoform X1 [Drosophila miranda]
>gi|1036784059|ref|XP_017135584.1| PREDICTED: protein sym1 isoform X2 [Drosophila miranda]
>gi|1036784002|ref|XP_017135581.1| PREDICTED: uncharacterized protein LOC108151467 [Drosophila
>gi|1036783889|ref|XP_017135574.1| PREDICTED: trypsin eta [Drosophila miranda]
>gi|1036783927|ref|XP_017135577.1| PREDICTED: ras-related protein Rab-27A isoform X1 [Drosophil
>gi|1036783946|ref|XP_017135578.1| PREDICTED: ras-related protein Rab-27A isoform X2 [Drosophil
>gi|1036783965|ref|XP_017135579.1| PREDICTED: ras-related protein Rab-27A isoform X2 [Drosophil
>gi|1036783983|ref|XP_017135580.1| PREDICTED: ras-related protein Rab-27A isoform X3 [Drosophil
>gi|1036783834|ref|XP_017135571.1| PREDICTED: uncharacterized protein LOC108151462 [Drosophila
>gi|1036783853|ref|XP_017135572.1| PREDICTED: pleckstrin homology domain-containing family F me
>gi|1036783872|ref|XP_017135573.1| PREDICTED: pleckstrin homology domain-containing family F me
>gi|1036783908|ref|XP_017135576.1| PREDICTED: 40S ribosomal protein SA [Drosophila miranda]
>gi|1036849305|ref|XP_017139220.1| PREDICTED: glutamate receptor ionotropic, NMDA 2B isoform X
>gi|1036849270|ref|XP_017139218.1| PREDICTED: glutamate receptor ionotropic, NMDA 2B isoform X
>gi|1036849289|ref|XP_017139219.1| PREDICTED: glutamate receptor ionotropic, NMDA 2B isoform X
>gi|1036764581|ref|XP_017157124.1| PREDICTED: uncharacterized protein LOC108165560 [Drosophila
>gi|1036820681|ref|XP_017137623.1| PREDICTED: BTB/POZ domain-containing protein KCTD5 [Drosophi
>gi|1036758489|ref|XP_017156781.1| PREDICTED: LOW QUALITY PROTEIN: mucin-5AC [Drosophila mirand
>gi|1036770545|ref|XP_017134801.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770578|ref|XP_017134803.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770512|ref|XP_017134799.1| PREDICTED: broad-complex core protein isoform X1 [Drosophila
>gi|1036770529|ref|XP_017134800.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770561|ref|XP_017134802.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770494|ref|XP_017134798.1| PREDICTED: vacuolar protein sorting-associated protein 18 ho
>gi|1036770656|ref|XP_017134808.1| PREDICTED: protein C19orf12 homolog [Drosophila miranda]
>gi|1036770597|ref|XP_017134804.1| PREDICTED: protein halfway isoform X1 [Drosophila miranda]
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>gi|1036770616|ref|XP_017134805.1| PREDICTED: protein halfway isoform X2 [Drosophila miranda]
>gi|1036770438|ref|XP_017134795.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770420|ref|XP_017134794.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770475|ref|XP_017134797.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770456|ref|XP_017134796.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770676|ref|XP_017134809.1| PREDICTED: uncharacterized protein LOC108150978 [Drosophila
>gi|1036770637|ref|XP_017134807.1| PREDICTED: uncharacterized protein LOC108150976 [Drosophila
>gi|1036834752|ref|XP_017138409.1| PREDICTED: protein a6 [Drosophila miranda]
>gi|1036758508|ref|XP_017156782.1| PREDICTED: trypsin-3 [Drosophila miranda]
>gi|1036850967|ref|XP_017139319.1| PREDICTED: delta-sarcoglycan [Drosophila miranda]
>gi|1036766628|ref|XP_017134584.1| PREDICTED: uncharacterized protein LOC108150793 [Drosophila
>gi|1036765469|ref|XP_017157173.1| PREDICTED: uncharacterized protein LOC108165607 [Drosophila
>gi|1036803038|ref|XP_017136656.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802982|ref|XP_017136653.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802961|ref|XP_017136652.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802942|ref|XP_017136651.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802924|ref|XP_017136650.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036803000|ref|XP_017136654.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036803019|ref|XP_017136655.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036803059|ref|XP_017136657.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036758526|ref|XP_017156783.1| PREDICTED: LOW QUALITY PROTEIN: TRAF3-interacting protein 1-
>gi|1036776046|ref|XP_017135108.1| PREDICTED: neurobeachin isoform X5 [Drosophila miranda]
>gi|1036776063|ref|XP_017135109.1| PREDICTED: neurobeachin isoform X6 [Drosophila miranda]
>gi|1036776084|ref|XP_017135110.1| PREDICTED: neurobeachin isoform X7 [Drosophila miranda]
>gi|1036776196|ref|XP_017135116.1| PREDICTED: neurobeachin isoform X13 [Drosophila miranda]
>gi|1036776141|ref|XP_017135113.1| PREDICTED: neurobeachin isoform X10 [Drosophila miranda]
>gi|1036776234|ref|XP_017135118.1| PREDICTED: neurobeachin isoform X15 [Drosophila miranda]
>gi|1036776308|ref|XP_017135122.1| PREDICTED: proteasomal ubiquitin receptor ADRM1 homolog [Dr
>gi|1036776327|ref|XP_017135123.1| PREDICTED: proteasome subunit alpha type-3-like [Drosophila
>gi|1036776346|ref|XP_017135124.1| PREDICTED: uncharacterized protein LOC108151189 [Drosophila
>gi|1036776270|ref|XP_017135120.1| PREDICTED: uncharacterized protein LOC108151185 [Drosophila
>gi|1036776289|ref|XP_017135121.1| PREDICTED: uncharacterized protein LOC108151185 [Drosophila
>gi|1036776251|ref|XP_017135119.1| PREDICTED: uncharacterized abhydrolase domain-containing pr
>gi|1036775966|ref|XP_017135104.1| PREDICTED: neurobeachin isoform X1 [Drosophila miranda]
>gi|1036776160|ref|XP_017135114.1| PREDICTED: neurobeachin isoform X11 [Drosophila miranda]
>gi|1036775985|ref|XP_017135105.1| PREDICTED: neurobeachin isoform X2 [Drosophila miranda]
>gi|1036776004|ref|XP_017135106.1| PREDICTED: neurobeachin isoform X3 [Drosophila miranda]
>gi|1036776023|ref|XP_017135107.1| PREDICTED: neurobeachin isoform X4 [Drosophila miranda]
>gi|1036776179|ref|XP_017135115.1| PREDICTED: neurobeachin isoform X12 [Drosophila miranda]
>gi|1036776103|ref|XP_017135111.1| PREDICTED: neurobeachin isoform X8 [Drosophila miranda]
>gi|1036776120|ref|XP_017135112.1| PREDICTED: neurobeachin isoform X9 [Drosophila miranda]
>gi|1036776215|ref|XP_017135117.1| PREDICTED: neurobeachin isoform X14 [Drosophila miranda]
>gi|1036776383|ref|XP_017135127.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1036776365|ref|XP_017135125.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1036767718|ref|XP_017134642.1| PREDICTED: uncharacterized protein LOC108150849 [Drosophila
>gi|1036852578|ref|XP_017139419.1| PREDICTED: dnaJ homolog subfamily C member 7-like [Drosophil
>gi|1036764462|ref|XP_017157119.1| PREDICTED: uncharacterized protein LOC108165557 [Drosophila
>gi|1036773580|ref|XP_017134973.1| PREDICTED: uncharacterized protein LOC108151094 isoform X1
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>gi|1036773597|ref|XP_017134974.1| PREDICTED: uncharacterized protein LOC108151094 isoform X2
>gi|1036773394|ref|XP_017134962.1| PREDICTED: uncharacterized protein LOC108151087 [Drosophila
>gi|1036773671|ref|XP_017134979.1| PREDICTED: uncharacterized protein LOC108151096 [Drosophila
>gi|1036773564|ref|XP_017134972.1| PREDICTED: cyclin-dependent kinase 7 [Drosophila miranda]
>gi|1036773690|ref|XP_017134980.1| PREDICTED: U1 small nuclear ribonucleoprotein A [Drosophila
>gi|1036773709|ref|XP_017134981.1| PREDICTED: thioredoxin-2 [Drosophila miranda]
>gi|1036773728|ref|XP_017134982.1| PREDICTED: thioredoxin-1 [Drosophila miranda]
>gi|1036773431|ref|XP_017134964.1| PREDICTED: uncharacterized protein LOC108151090 [Drosophila
>gi|1036773413|ref|XP_017134963.1| PREDICTED: uncharacterized protein LOC108151089 [Drosophila
>gi|1036773542|ref|XP_017134971.1| PREDICTED: something about silencing protein 10 [Drosophila
>gi|1036773375|ref|XP_017134961.1| PREDICTED: RNA-binding protein 4F [Drosophila miranda]
>gi|1036773469|ref|XP_017134966.1| PREDICTED: acid phosphatase type 7 isoform X2 [Drosophila mi
>gi|1036773523|ref|XP_017134970.1| PREDICTED: acid phosphatase type 7 isoform X5 [Drosophila mi
>gi|1036773486|ref|XP_017134968.1| PREDICTED: acid phosphatase type 7 isoform X3 [Drosophila mi
>gi|1036773505|ref|XP_017134969.1| PREDICTED: acid phosphatase type 7 isoform X4 [Drosophila mi
>gi|1036773450|ref|XP_017134965.1| PREDICTED: acid phosphatase type 7 isoform X1 [Drosophila mi
>gi|1036773652|ref|XP_017134978.1| PREDICTED: shematrin-like protein 2 isoform X3 [Drosophila m
>gi|1036773634|ref|XP_017134977.1| PREDICTED: glycine-rich cell wall structural protein 2 isofo
>gi|1036773616|ref|XP_017134975.1| PREDICTED: glycine-rich RNA-binding protein 3, mitochondrial
>gi|1036824695|ref|XP_017137850.1| PREDICTED: RNA-binding protein 33 [Drosophila miranda]
>gi|1036763208|ref|XP_017157053.1| PREDICTED: uncharacterized protein LOC108165508 [Drosophila
>gi|1036790788|ref|XP_017135965.1| PREDICTED: zinc finger protein 12 [Drosophila miranda]
>gi|1036790846|ref|XP_017135968.1| PREDICTED: probable 6-phosphogluconolactonase [Drosophila mi
>gi|1036790807|ref|XP_017135966.1| PREDICTED: nuclear RNA export factor 1 [Drosophila miranda]
>gi|1036790825|ref|XP_017135967.1| PREDICTED: putative sodium-dependent multivitamin transporte
>gi|1036790917|ref|XP_017135972.1| PREDICTED: uncharacterized protein LOC108151711 [Drosophila
>gi|1036790934|ref|XP_017135973.1| PREDICTED: uncharacterized protein LOC108151712 [Drosophila
>gi|1036790863|ref|XP_017135970.1| PREDICTED: uncharacterized protein LOC108151710 isoform X1
>gi|1036790882|ref|XP_017135971.1| PREDICTED: uncharacterized protein LOC108151710 isoform X2
>gi|1036812105|ref|XP_017137153.1| PREDICTED: uncharacterized protein LOC108152374 [Drosophila
>gi|1036812026|ref|XP_017137149.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036812047|ref|XP_017137150.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036812066|ref|XP_017137151.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036812087|ref|XP_017137152.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036832144|ref|XP_017138273.1| PREDICTED: pyruvate dehydrogenase phosphatase regulatory sub
>gi|1036762241|ref|XP_017157000.1| PREDICTED: optomotor-blind protein [Drosophila miranda]
>gi|1036762365|ref|XP_017157008.1| PREDICTED: uncharacterized protein CG3556 [Drosophila mirand
>gi|1036762403|ref|XP_017157010.1| PREDICTED: protein TonB [Drosophila miranda]
>gi|1036762223|ref|XP_017156999.1| PREDICTED: AP-3 complex subunit beta-2 [Drosophila miranda]
>gi|1036762437|ref|XP_017157012.1| PREDICTED: protein SREK1IP1 [Drosophila miranda]
>gi|1036762384|ref|XP_017157009.1| PREDICTED: uncharacterized protein LOC108165469 [Drosophila
>gi|1036762418|ref|XP_017157011.1| PREDICTED: serine--pyruvate aminotransferase, mitochondrial
>gi|1036762455|ref|XP_017157014.1| PREDICTED: 60S ribosomal protein L7a [Drosophila miranda]
>gi|1036762259|ref|XP_017157002.1| PREDICTED: protein deltex [Drosophila miranda]
>gi|1036762014|ref|XP_017156992.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762026|ref|XP_017156993.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762131|ref|XP_017156994.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762148|ref|XP_017156995.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
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>gi|1036762166|ref|XP_017156996.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036761940|ref|XP_017156988.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036761959|ref|XP_017156989.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036761998|ref|XP_017156991.1| PREDICTED: uncharacterized protein LOC108165461 isoform X3
>gi|1036762204|ref|XP_017156998.1| PREDICTED: uncharacterized protein LOC108165461 isoform X10
>gi|1036762186|ref|XP_017156997.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036762560|ref|XP_017157019.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036761977|ref|XP_017156990.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036762493|ref|XP_017157016.1| PREDICTED: uncharacterized protein LOC108165476 [Drosophila
>gi|1036762511|ref|XP_017157017.1| PREDICTED: 40S ribosomal protein S12, mitochondrial [Drosoph
>gi|1036758542|ref|XP_017156784.1| PREDICTED: protein giant [Drosophila miranda]
>gi|1036762524|ref|XP_017157018.1| PREDICTED: uncharacterized protein LOC108165478 [Drosophila
>gi|1036758559|ref|XP_017156785.1| PREDICTED: LOW QUALITY PROTEIN: protein regulator of cytokin
>gi|1036758578|ref|XP_017156786.1| PREDICTED: protein argonaute-2-like, partial [Drosophila mi
>gi|1036762474|ref|XP_017157015.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1036762347|ref|XP_017157007.1| PREDICTED: uncharacterized protein LOC108165467 isoform X4
>gi|1036762314|ref|XP_017157005.1| PREDICTED: uncharacterized protein LOC108165467 isoform X2
>gi|1036762295|ref|XP_017157004.1| PREDICTED: uncharacterized protein LOC108165467 isoform X1
>gi|1036762332|ref|XP_017157006.1| PREDICTED: uncharacterized protein LOC108165467 isoform X3
>gi|1036758596|ref|XP_017156787.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036762277|ref|XP_017157003.1| PREDICTED: zinc finger and BTB domain-containing protein 48
>gi|1036817445|ref|XP_017137449.1| PREDICTED: casein kinase I [Drosophila miranda]
>gi|1036811137|ref|XP_017137103.1| PREDICTED: syntaxin-binding protein 5 isoform X3 [Drosophila
>gi|1036811117|ref|XP_017137102.1| PREDICTED: syntaxin-binding protein 5 isoform X2 [Drosophila
>gi|1036811155|ref|XP_017137104.1| PREDICTED: syntaxin-binding protein 5 isoform X4 [Drosophila
>gi|1036811078|ref|XP_017137100.1| PREDICTED: syntaxin-binding protein 5 isoform X1 [Drosophila
>gi|1036811097|ref|XP_017137101.1| PREDICTED: syntaxin-binding protein 5 isoform X1 [Drosophila
>gi|1036811213|ref|XP_017137107.1| PREDICTED: syntaxin-binding protein 5 isoform X7 [Drosophila
>gi|1036811194|ref|XP_017137106.1| PREDICTED: syntaxin-binding protein 5 isoform X6 [Drosophila
>gi|1036811173|ref|XP_017137105.1| PREDICTED: syntaxin-binding protein 5 isoform X5 [Drosophila
>gi|1036811232|ref|XP_017137108.1| PREDICTED: putative glutathione-specific gamma-glutamylcyclo
>gi|1036845029|ref|XP_017138984.1| PREDICTED: uncharacterized protein LOC108153465 [Drosophila
>gi|1036765005|ref|XP_017157146.1| PREDICTED: uncharacterized protein LOC108165585 [Drosophila
>gi|1036766684|ref|XP_017134587.1| PREDICTED: sperm flagellar protein 1 [Drosophila miranda]
>gi|1036825751|ref|XP_017137904.1| PREDICTED: NK1 transcription factor-related protein 2 [Dros
>gi|1036825694|ref|XP_017137900.1| PREDICTED: folylpolyglutamate synthase, mitochondrial [Dros
>gi|1036825713|ref|XP_017137901.1| PREDICTED: folylpolyglutamate synthase, mitochondrial [Dros
>gi|1036825732|ref|XP_017137903.1| PREDICTED: box C/D snoRNA protein 1 [Drosophila miranda]
>gi|1036800658|ref|XP_017136523.1| PREDICTED: uncharacterized protein LOC108152037 isoform X3
>gi|1036800675|ref|XP_017136524.1| PREDICTED: uncharacterized protein LOC108152037 isoform X4
>gi|1036800584|ref|XP_017136519.1| PREDICTED: uncharacterized protein LOC108152037 isoform X1
>gi|1036800603|ref|XP_017136520.1| PREDICTED: uncharacterized protein LOC108152037 isoform X2
>gi|1036800621|ref|XP_017136521.1| PREDICTED: uncharacterized protein LOC108152037 isoform X2
>gi|1036800639|ref|XP_017136522.1| PREDICTED: uncharacterized protein LOC108152037 isoform X2
>gi|1036837275|ref|XP_017138559.1| PREDICTED: ATP-binding cassette sub-family B member 8, mito
>gi|1036823056|ref|XP_017137756.1| PREDICTED: LIM domain kinase 1 isoform X1 [Drosophila miran
>gi|1036823075|ref|XP_017137757.1| PREDICTED: LIM domain kinase 1 isoform X1 [Drosophila miran
>gi|1036823094|ref|XP_017137758.1| PREDICTED: LIM domain kinase 1 isoform X2 [Drosophila miran
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>gi|1036819003|ref|XP_017137539.1| PREDICTED: interferon regulatory factor 2-binding protein-l
>gi|1036819022|ref|XP_017137540.1| PREDICTED: interferon regulatory factor 2-binding protein-l
>gi|1036833557|ref|XP_017138351.1| PREDICTED: uncharacterized protein LOC108153079 [Drosophila
>gi|1036833576|ref|XP_017138353.1| PREDICTED: larval cuticle protein LCP-17 [Drosophila miranda
>gi|1036837219|ref|XP_017138555.1| PREDICTED: rho GTPase-activating protein gacZ [Drosophila mi
>gi|1036849748|ref|XP_017139248.1| PREDICTED: Meckel syndrome type 1 protein [Drosophila mirand
>gi|1036765417|ref|XP_017157169.1| PREDICTED: pupal cuticle protein 36 [Drosophila miranda]
>gi|1036831303|ref|XP_017138226.1| PREDICTED: uncharacterized protein LOC108152999 isoform X1
>gi|1036831322|ref|XP_017138227.1| PREDICTED: uncharacterized protein LOC108152999 isoform X2
>gi|1036831338|ref|XP_017138228.1| PREDICTED: uncharacterized protein LOC108152999 isoform X2
>gi|1036832992|ref|XP_017138319.1| PREDICTED: ras-related protein Rab-40C [Drosophila miranda]
>gi|1036758615|ref|XP_017156788.1| PREDICTED: glycine-rich cell wall structural protein 1.8 [D
>gi|1036758634|ref|XP_017156789.1| PREDICTED: uncharacterized protein LOC108165264 [Drosophila
>gi|1036837488|ref|XP_017138572.1| PREDICTED: putative protein kinase C delta type homolog iso
>gi|1036837506|ref|XP_017138573.1| PREDICTED: putative protein kinase C delta type homolog iso
>gi|1036837522|ref|XP_017138574.1| PREDICTED: putative protein kinase C delta type homolog iso
>gi|1036837472|ref|XP_017138571.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1036837454|ref|XP_017138570.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1036764720|ref|XP_017157132.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036851255|ref|XP_017139337.1| PREDICTED: actin-57B-like [Drosophila miranda]
>gi|1036765890|ref|XP_017134545.1| PREDICTED: ubiquitin-conjugating enzyme E2 D2 [Drosophila mi
>gi|1036766155|ref|XP_017134561.1| PREDICTED: protein arginine N-methyltransferase 1-B-like [D
>gi|1036851579|ref|XP_017139357.1| PREDICTED: 14-3-3 protein epsilon-like [Drosophila miranda]
>gi|1036848183|ref|XP_017139155.1| PREDICTED: midasin-like [Drosophila miranda]
>gi|1036814388|ref|XP_017137278.1| PREDICTED: uncharacterized protein LOC108152443 [Drosophila
>gi|1036764670|ref|XP_017157129.1| PREDICTED: casein kinase I [Drosophila miranda]
>gi|1036758650|ref|XP_017156790.1| PREDICTED: uncharacterized protein LOC108165266 [Drosophila
>gi|1036850217|ref|XP_017139278.1| PREDICTED: uncharacterized protein LOC108153673 isoform X2
>gi|1036850198|ref|XP_017139277.1| PREDICTED: uncharacterized protein LOC108153673 isoform X1
>gi|1036758668|ref|XP_017156791.1| PREDICTED: LOW QUALITY PROTEIN: chromosomal serine/threonine
>gi|1036784116|ref|XP_017135587.1| PREDICTED: teneurin-a [Drosophila miranda]
>gi|1036784135|ref|XP_017135588.1| PREDICTED: uncharacterized protein LOC108151472 isoform X1
>gi|1036784154|ref|XP_017135589.1| PREDICTED: uncharacterized protein LOC108151472 isoform X2
>gi|1036839595|ref|XP_017138694.1| PREDICTED: cell surface glycoprotein 1 isoform X5 [Drosophil
>gi|1036839577|ref|XP_017138693.1| PREDICTED: cell surface glycoprotein 1 isoform X4 [Drosophil
>gi|1036839558|ref|XP_017138691.1| PREDICTED: cell surface glycoprotein 1 isoform X3 [Drosophil
>gi|1036839611|ref|XP_017138695.1| PREDICTED: cell surface glycoprotein 1 isoform X6 [Drosophil
>gi|1036839540|ref|XP_017138690.1| PREDICTED: cell surface glycoprotein 1 isoform X2 [Drosophil
>gi|1036839521|ref|XP_017138689.1| PREDICTED: cell surface glycoprotein 1 isoform X1 [Drosophil
>gi|1036758687|ref|XP_017156792.1| PREDICTED: ADP-ribosylation factor-like protein 3 [Drosophil
>gi|1036764599|ref|XP_017157125.1| PREDICTED: uncharacterized protein LOC108165561 [Drosophila
>gi|1036842018|ref|XP_017138823.1| PREDICTED: sporozoite surface protein 2 isoform X2 [Drosoph
>gi|1036841804|ref|XP_017138822.1| PREDICTED: proteoglycan 4 isoform X1 [Drosophila miranda]
>gi|1036758706|ref|XP_017156794.1| PREDICTED: protein cutoff-like [Drosophila miranda]
>gi|1036842214|ref|XP_017138835.1| PREDICTED: flightin [Drosophila miranda]
>gi|1036837630|ref|XP_017138581.1| PREDICTED: protein commissureless [Drosophila miranda]
>gi|1036848387|ref|XP_017139167.1| PREDICTED: uncharacterized protein LOC108153597 [Drosophila
>gi|1036848420|ref|XP_017139169.1| PREDICTED: uncharacterized protein LOC108153598 [Drosophila
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>gi|1036848434|ref|XP_017139170.1| PREDICTED: uncharacterized protein LOC108153598 [Drosophila
>gi|1036840797|ref|XP_017138761.1| PREDICTED: uncharacterized protein LOC108153336 [Drosophila
>gi|1036767145|ref|XP_017134615.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036765814|ref|XP_017134541.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036768422|ref|XP_017134681.1| PREDICTED: uncharacterized protein LOC108150883 [Drosophila
>gi|1036768703|ref|XP_017134696.1| PREDICTED: uncharacterized protein LOC108150897 [Drosophila
>gi|1036758723|ref|XP_017156795.1| PREDICTED: uncharacterized protein LOC108165271 [Drosophila
>gi|1036758741|ref|XP_017156796.1| PREDICTED: gamete and mating-type specific protein A-like [D
>gi|1036805803|ref|XP_017136813.1| PREDICTED: uncharacterized protein LOC108152183 isoform X2
>gi|1036805765|ref|XP_017136811.1| PREDICTED: uncharacterized protein LOC108152183 isoform X1
>gi|1036805784|ref|XP_017136812.1| PREDICTED: uncharacterized protein LOC108152183 isoform X1
>gi|1036805822|ref|XP_017136814.1| PREDICTED: uncharacterized protein LOC108152183 isoform X3
>gi|1036805746|ref|XP_017136810.1| PREDICTED: zinc finger and BTB domain-containing protein 49
>gi|1036770123|ref|XP_017134776.1| PREDICTED: uncharacterized protein LOC108150953 [Drosophila
>gi|1036768344|ref|XP_017134678.1| PREDICTED: phosducin-like protein [Drosophila miranda]
>gi|1036768153|ref|XP_017134666.1| PREDICTED: uncharacterized protein LOC108150872 isoform X3
>gi|1036768117|ref|XP_017134664.1| PREDICTED: uncharacterized protein LOC108150872 isoform X1
>gi|1036768134|ref|XP_017134665.1| PREDICTED: uncharacterized protein LOC108150872 isoform X2
>gi|1036768172|ref|XP_017134667.1| PREDICTED: rho GTPase-activating protein 20 isoform X4 [Dros
>gi|1036768327|ref|XP_017134676.1| PREDICTED: ubiquitin-conjugating enzyme E2 R2 [Drosophila mi
>gi|1036768267|ref|XP_017134673.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768246|ref|XP_017134672.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768306|ref|XP_017134675.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768286|ref|XP_017134674.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768189|ref|XP_017134669.1| PREDICTED: U3 small nucleolar RNA-associated protein 14 homo
>gi|1036768227|ref|XP_017134671.1| PREDICTED: general transcription factor IIH subunit 4 [Droso
>gi|1036768208|ref|XP_017134670.1| PREDICTED: probable Ufm1-specific protease 2 [Drosophila mir
>gi|1036768098|ref|XP_017134663.1| PREDICTED: microtubule-associated serine/threonine-protein k
>gi|1036768078|ref|XP_017134662.1| PREDICTED: microtubule-associated serine/threonine-protein k
>gi|1036772210|ref|XP_017134895.1| PREDICTED: DCN1-like protein [Drosophila miranda]
>gi|1036771984|ref|XP_017134882.1| PREDICTED: T-cell immunomodulatory protein [Drosophila miran
>gi|1036772002|ref|XP_017134883.1| PREDICTED: protein argonaute-2 [Drosophila miranda]
>gi|1036772367|ref|XP_017134903.1| PREDICTED: LOW QUALITY PROTEIN: protein disulfide-isomerase
>gi|1036772406|ref|XP_017134905.1| PREDICTED: LOW QUALITY PROTEIN: protein cutoff-like [Drosoph
>gi|1036772040|ref|XP_017134885.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772021|ref|XP_017134884.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772057|ref|XP_017134886.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772075|ref|XP_017134887.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772092|ref|XP_017134888.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772115|ref|XP_017134889.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772134|ref|XP_017134890.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772153|ref|XP_017134891.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772172|ref|XP_017134892.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036758761|ref|XP_017156797.1| PREDICTED: uncharacterized protein LOC108165274 [Drosophila
>gi|1036772327|ref|XP_017134901.1| PREDICTED: uncharacterized protein LOC108151041 [Drosophila
>gi|1036772306|ref|XP_017134900.1| PREDICTED: uncharacterized protein LOC108151040 [Drosophila
>gi|1036758779|ref|XP_017156799.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036772287|ref|XP_017134899.1| PREDICTED: BAG family molecular chaperone regulator 2 isofor
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>gi|1036772230|ref|XP_017134896.1| PREDICTED: BAG family molecular chaperone regulator 2 isofo
>gi|1036772248|ref|XP_017134897.1| PREDICTED: BAG family molecular chaperone regulator 2 isofo
>gi|1036772268|ref|XP_017134898.1| PREDICTED: BAG family molecular chaperone regulator 2 isofo
>gi|1036772193|ref|XP_017134893.1| PREDICTED: uncharacterized protein LOC108151037 [Drosophila
>gi|1036772385|ref|XP_017134904.1| PREDICTED: protein midgut expression 1 [Drosophila miranda]
>gi|1036772341|ref|XP_017134902.1| PREDICTED: uncharacterized protein LOC108151042 [Drosophila
>gi|1036834954|ref|XP_017138421.1| PREDICTED: peptide methionine sulfoxide reductase isoform X2
>gi|1036834935|ref|XP_017138420.1| PREDICTED: peptide methionine sulfoxide reductase isoform X1
>gi|1036834970|ref|XP_017138422.1| PREDICTED: peptide methionine sulfoxide reductase isoform X3
>gi|1036834989|ref|XP_017138423.1| PREDICTED: gonadal protein gdl isoform X1 [Drosophila mirand
>gi|1036835004|ref|XP_017138424.1| PREDICTED: gonadal protein gdl isoform X2 [Drosophila mirand
>gi|1036763476|ref|XP_017157068.1| PREDICTED: protein Z600 [Drosophila miranda]
>gi|1036841769|ref|XP_017138820.1| PREDICTED: uncharacterized protein LOC108153364 [Drosophila
>gi|1036822908|ref|XP_017137747.1| PREDICTED: OTU domain-containing protein 6B [Drosophila mira
>gi|1036822925|ref|XP_017137748.1| PREDICTED: sugar transporter SWEET1 [Drosophila miranda]
>gi|1036822889|ref|XP_017137746.1| PREDICTED: DDB1- and CUL4-associated factor 13 [Drosophila m
>gi|1036835719|ref|XP_017138466.1| PREDICTED: uncharacterized protein LOC108153172 [Drosophila
>gi|1036835738|ref|XP_017138468.1| PREDICTED: mpv17-like protein isoform X1 [Drosophila miranda
>gi|1036835756|ref|XP_017138469.1| PREDICTED: mpv17-like protein isoform X2 [Drosophila miranda
>gi|1036835776|ref|XP_017138470.1| PREDICTED: mpv17-like protein isoform X2 [Drosophila miranda
>gi|1036829575|ref|XP_017138124.1| PREDICTED: protein toll [Drosophila miranda]
>gi|1036823723|ref|XP_017137793.1| PREDICTED: cationic amino acid transporter 2 isoform X2 [Dro
>gi|1036823665|ref|XP_017137790.1| PREDICTED: high affinity cationic amino acid transporter 1 i
>gi|1036823683|ref|XP_017137791.1| PREDICTED: high affinity cationic amino acid transporter 1 i
>gi|1036823704|ref|XP_017137792.1| PREDICTED: high affinity cationic amino acid transporter 1 i
>gi|1036823740|ref|XP_017137794.1| PREDICTED: bestrophin-2 [Drosophila miranda]
>gi|1036852454|ref|XP_017139412.1| PREDICTED: bestrophin-4 [Drosophila miranda]
>gi|1036813967|ref|XP_017137253.1| PREDICTED: protein toll [Drosophila miranda]
>gi|1036782493|ref|XP_017135490.1| PREDICTED: endoplasmic reticulum-Golgi intermediate compartm
>gi|1036782507|ref|XP_017135491.1| PREDICTED: reactive oxygen species modulator 1 [Drosophila m
>gi|1036782455|ref|XP_017135488.1| PREDICTED: U4/U6 small nuclear ribonucleoprotein Prp31 [Dros
>gi|1036782379|ref|XP_017135484.1| PREDICTED: probable DNA mismatch repair protein Msh6 [Drosop
>gi|1036782397|ref|XP_017135485.1| PREDICTED: glycoprotein 3-alpha-L-fucosyltransferase A [Dros
>gi|1036782419|ref|XP_017135486.1| PREDICTED: glycoprotein 3-alpha-L-fucosyltransferase A [Dros
>gi|1036782437|ref|XP_017135487.1| PREDICTED: glycoprotein 3-alpha-L-fucosyltransferase A [Dros
>gi|1036782474|ref|XP_017135489.1| PREDICTED: peroxisomal biogenesis factor 3 [Drosophila miran
>gi|1036815904|ref|XP_017137361.1| PREDICTED: pyroglutamyl-peptidase 1 [Drosophila miranda]
>gi|1036815923|ref|XP_017137362.1| PREDICTED: uncharacterized protein LOC108152498 [Drosophila
>gi|1036815883|ref|XP_017137360.1| PREDICTED: protein disulfide-isomerase [Drosophila miranda]
>gi|1036790178|ref|XP_017135932.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X1
>gi|1036790197|ref|XP_017135933.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X2
>gi|1036790215|ref|XP_017135934.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X2
>gi|1036790236|ref|XP_017135935.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X2
>gi|1036790159|ref|XP_017135931.1| PREDICTED: centrosomal protein of 135 kDa isoform X4 [Drosop
>gi|1036790064|ref|XP_017135926.1| PREDICTED: uncharacterized protein LOC108151691 isoform X1
>gi|1036790083|ref|XP_017135927.1| PREDICTED: uncharacterized protein LOC108151691 isoform X2
>gi|1036790140|ref|XP_017135930.1| PREDICTED: centrosomal protein of 135 kDa isoform X3 [Drosop
>gi|1036790121|ref|XP_017135929.1| PREDICTED: centrosomal protein of 135 kDa isoform X2 [Drosop
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>gi|1036790102|ref|XP_017135928.1| PREDICTED: centrosomal protein of 135 kDa isoform X1 [Droso
>gi|1036790252|ref|XP_017135936.1| PREDICTED: uncharacterized protein LOC108151695 isoform X1
>gi|1036790269|ref|XP_017135937.1| PREDICTED: uncharacterized protein LOC108151695 isoform X2
>gi|1036852144|ref|XP_017139393.1| PREDICTED: uncharacterized protein LOC108153765 [Drosophila
>gi|1036813503|ref|XP_017137226.1| PREDICTED: CTP synthase isoform X1 [Drosophila miranda]
>gi|1036813557|ref|XP_017137229.1| PREDICTED: CTP synthase isoform X3 [Drosophila miranda]
>gi|1036813522|ref|XP_017137227.1| PREDICTED: uncharacterized protein LOC108152422 isoform X2
>gi|1036813537|ref|XP_017137228.1| PREDICTED: uncharacterized protein LOC108152422 isoform X2
>gi|1036813597|ref|XP_017137231.1| PREDICTED: uncharacterized protein LOC108152423 isoform X2
>gi|1036813578|ref|XP_017137230.1| PREDICTED: uncharacterized protein LOC108152423 isoform X1
>gi|1036833393|ref|XP_017138343.1| PREDICTED: glycine--tRNA ligase [Drosophila miranda]
>gi|1036777693|ref|XP_017135206.1| PREDICTED: putative uncharacterized protein DDB_G0271606 [Dr
>gi|1036777453|ref|XP_017135192.1| PREDICTED: A-kinase anchor protein 9 isoform X2 [Drosophila
>gi|1036777509|ref|XP_017135196.1| PREDICTED: golgin subfamily A member 4 isoform X5 [Drosophil
>gi|1036777528|ref|XP_017135197.1| PREDICTED: golgin subfamily A member 4 isoform X6 [Drosophil
>gi|1036777566|ref|XP_017135199.1| PREDICTED: pericentrin isoform X8 [Drosophila miranda]
>gi|1036777472|ref|XP_017135193.1| PREDICTED: A-kinase anchor protein 9 isoform X3 [Drosophila
>gi|1036777547|ref|XP_017135198.1| PREDICTED: golgin subfamily A member 4 isoform X7 [Drosophil
>gi|1036777582|ref|XP_017135200.1| PREDICTED: pericentrin isoform X9 [Drosophila miranda]
>gi|1036777601|ref|XP_017135201.1| PREDICTED: pericentrin isoform X10 [Drosophila miranda]
>gi|1036777620|ref|XP_017135202.1| PREDICTED: golgin subfamily A member 4 isoform X11 [Drosoph
>gi|1036777434|ref|XP_017135191.1| PREDICTED: A-kinase anchor protein 9 isoform X1 [Drosophila
>gi|1036777490|ref|XP_017135195.1| PREDICTED: A-kinase anchor protein 9 isoform X4 [Drosophila
>gi|1036777637|ref|XP_017135203.1| PREDICTED: A-kinase anchor protein 9 isoform X12 [Drosophila
>gi|1036777786|ref|XP_017135212.1| PREDICTED: putative odorant receptor 71a [Drosophila miranda
>gi|1036777656|ref|XP_017135204.1| PREDICTED: pericentrin isoform X13 [Drosophila miranda]
>gi|1036777675|ref|XP_017135205.1| PREDICTED: trichohyalin isoform X14 [Drosophila miranda]
>gi|1036777824|ref|XP_017135214.1| PREDICTED: uncharacterized protein LOC108151228 [Drosophila
>gi|1036777842|ref|XP_017135215.1| PREDICTED: uncharacterized protein LOC108151229 [Drosophila
>gi|1036777905|ref|XP_017135219.1| PREDICTED: prostatic spermine-binding protein [Drosophila mi
>gi|1036777870|ref|XP_017135218.1| PREDICTED: probable DNA-directed RNA polymerase subunit delt
>gi|1036777857|ref|XP_017135216.1| PREDICTED: uncharacterized protein LOC108151230 [Drosophila
>gi|1036777767|ref|XP_017135211.1| PREDICTED: synaptotagmin-5 isoform X2 [Drosophila miranda]
>gi|1036777805|ref|XP_017135213.1| PREDICTED: GS homeobox 1 [Drosophila miranda]
>gi|1036777710|ref|XP_017135207.1| PREDICTED: synaptotagmin-3 isoform X1 [Drosophila miranda]
>gi|1036777729|ref|XP_017135208.1| PREDICTED: synaptotagmin-3 isoform X1 [Drosophila miranda]
>gi|1036777748|ref|XP_017135210.1| PREDICTED: synaptotagmin-3 isoform X1 [Drosophila miranda]
>gi|1036765796|ref|XP_017134540.1| PREDICTED: uncharacterized protein LOC108150757 [Drosophila
>gi|1036829999|ref|XP_017138148.1| PREDICTED: reversion-inducing cysteine-rich protein with Ka
>gi|1036830018|ref|XP_017138149.1| PREDICTED: reversion-inducing cysteine-rich protein with Ka
>gi|1036830037|ref|XP_017138150.1| PREDICTED: proteasome subunit beta type-7 [Drosophila miran
>gi|1036833027|ref|XP_017138321.1| PREDICTED: 39S ribosomal protein L39, mitochondrial [Drosoph
>gi|1036833010|ref|XP_017138320.1| PREDICTED: zinc transporter ZIP10 [Drosophila miranda]
>gi|1036821311|ref|XP_017137658.1| PREDICTED: Y+L amino acid transporter 2 [Drosophila miranda]
>gi|1036821330|ref|XP_017137659.1| PREDICTED: Y+L amino acid transporter 2 [Drosophila miranda]
>gi|1036821349|ref|XP_017137660.1| PREDICTED: angio-associated migratory cell protein [Drosophi
>gi|1036848860|ref|XP_017139195.1| PREDICTED: uncharacterized protein LOC108153608 isoform X2
>gi|1036848841|ref|XP_017139194.1| PREDICTED: uncharacterized protein LOC108153608 isoform X1
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>gi|1036763155|ref|XP_017157052.1| PREDICTED: enhancer of split malpha protein [Drosophila mira
>gi|1036763331|ref|XP_017157059.1| PREDICTED: uncharacterized protein LOC108165514 [Drosophila
>gi|1036846698|ref|XP_017139073.1| PREDICTED: enhancer of split malpha protein [Drosophila mira
>gi|1036850162|ref|XP_017139274.1| PREDICTED: uridine phosphorylase 1-like [Drosophila miranda]
>gi|1036763315|ref|XP_017157058.1| PREDICTED: uncharacterized protein LOC108165513 [Drosophila
>gi|1036758797|ref|XP_017156800.1| PREDICTED: uncharacterized protein LOC108165276 [Drosophila
>gi|1036849104|ref|XP_017139209.1| PREDICTED: echinoderm microtubule-associated protein-like CO
>gi|1036767087|ref|XP_017134612.1| PREDICTED: uncharacterized protein LOC108150817 isoform X1
>gi|1036767108|ref|XP_017134613.1| PREDICTED: uncharacterized protein LOC108150817 isoform X2
>gi|1036830096|ref|XP_017138154.1| PREDICTED: Ig-like and fibronectin type-III domain-containin
>gi|1036812414|ref|XP_017137171.1| PREDICTED: uncharacterized protein LOC108152385 [Drosophila
>gi|1036812336|ref|XP_017137166.1| PREDICTED: probable helicase with zinc finger domain [Droso
>gi|1036812359|ref|XP_017137167.1| PREDICTED: probable helicase with zinc finger domain [Droso
>gi|1036812376|ref|XP_017137168.1| PREDICTED: probable helicase with zinc finger domain [Droso
>gi|1036812395|ref|XP_017137169.1| PREDICTED: probable helicase with zinc finger domain [Droso
>gi|1036838573|ref|XP_017138637.1| PREDICTED: H/ACA ribonucleoprotein complex subunit 2-like p
>gi|1036833046|ref|XP_017138322.1| PREDICTED: tudor domain-containing protein 3 [Drosophila mi
>gi|1036827923|ref|XP_017138033.1| PREDICTED: patatin-like phospholipase domain-containing prot
>gi|1036811742|ref|XP_017137134.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 23
>gi|1036825053|ref|XP_017137867.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylgl
>gi|1036825071|ref|XP_017137868.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylgl
>gi|1036825088|ref|XP_017137870.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylgl
>gi|1036825107|ref|XP_017137871.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylgl
>gi|1036774850|ref|XP_017135049.1| PREDICTED: uncharacterized protein LOC108151135 [Drosophila
>gi|1036774781|ref|XP_017135044.1| PREDICTED: transcription factor GAGA isoform X4 [Drosophila
>gi|1036774727|ref|XP_017135042.1| PREDICTED: transcription factor GAGA isoform X2 [Drosophila
>gi|1036774762|ref|XP_017135043.1| PREDICTED: transcription factor GAGA isoform X3 [Drosophila
>gi|1036774708|ref|XP_017135041.1| PREDICTED: transcription factor GAGA isoform X1 [Drosophila
>gi|1036774816|ref|XP_017135046.1| PREDICTED: protein roadkill [Drosophila miranda]
>gi|1036774632|ref|XP_017135037.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X1 [Drosoph
>gi|1036774651|ref|XP_017135038.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X2 [Drosoph
>gi|1036774670|ref|XP_017135039.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X3 [Drosoph
>gi|1036774903|ref|XP_017135052.1| PREDICTED: thioredoxin-2 [Drosophila miranda]
>gi|1036774885|ref|XP_017135051.1| PREDICTED: uncharacterized protein LOC108151137 [Drosophila
>gi|1036774689|ref|XP_017135040.1| PREDICTED: myosin heavy chain, striated muscle [Drosophila
>gi|1036774922|ref|XP_017135053.1| PREDICTED: uncharacterized protein LOC108151139 [Drosophila
>gi|1036774868|ref|XP_017135050.1| PREDICTED: uncharacterized protein LOC108151136 [Drosophila
>gi|1036774943|ref|XP_017135054.1| PREDICTED: uncharacterized protein LOC108151140 [Drosophila
>gi|1036774834|ref|XP_017135048.1| PREDICTED: protein PIH1D3 [Drosophila miranda]
>gi|1036774799|ref|XP_017135045.1| PREDICTED: venom protease [Drosophila miranda]
>gi|1036815942|ref|XP_017137363.1| PREDICTED: glypican-4 [Drosophila miranda]
>gi|1036834338|ref|XP_017138384.1| PREDICTED: polycystic kidney disease 2-like 2 protein [Droso
>gi|1036834375|ref|XP_017138386.1| PREDICTED: nucleoside-triphosphatase THEP1 [Drosophila miran
>gi|1036834356|ref|XP_017138385.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC5 [Dro
>gi|1036835252|ref|XP_017138439.1| PREDICTED: methyltransferase-like protein 22 [Drosophila mi
>gi|1036835236|ref|XP_017138437.1| PREDICTED: decaprenyl-diphosphate synthase subunit 2 [Drosop
>gi|1036801320|ref|XP_017136561.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036801339|ref|XP_017136562.1| PREDICTED: mediator of RNA polymerase II transcription subu
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>gi|1036801358|ref|XP_017136563.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036801377|ref|XP_017136565.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036764686|ref|XP_017157130.1| PREDICTED: uncharacterized protein LOC108165567 [Drosophila
>gi|1036851440|ref|XP_017139348.1| PREDICTED: uncharacterized protein LOC108153730 [Drosophila
>gi|1036843488|ref|XP_017138907.1| PREDICTED: uncharacterized protein LOC108153417 isoform X1
>gi|1036843507|ref|XP_017138908.1| PREDICTED: uncharacterized protein LOC108153417 isoform X2
>gi|1036843526|ref|XP_017138909.1| PREDICTED: uncharacterized protein LOC108153417 isoform X3
>gi|1036843471|ref|XP_017138906.1| PREDICTED: uncharacterized protein LOC108153416 [Drosophila
>gi|1036843545|ref|XP_017138910.1| PREDICTED: trypsin alpha-3 [Drosophila miranda]
>gi|1036840814|ref|XP_017138762.1| PREDICTED: fringe glycosyltransferase [Drosophila miranda]
>gi|1036840831|ref|XP_017138763.1| PREDICTED: uncharacterized protein LOC108153339 [Drosophila
>gi|1036758816|ref|XP_017156801.1| PREDICTED: uncharacterized protein LOC108165277 [Drosophila
>gi|1036830224|ref|XP_017138163.1| PREDICTED: uncharacterized protein LOC108152966 isoform X1
>gi|1036830241|ref|XP_017138164.1| PREDICTED: uncharacterized protein LOC108152966 isoform X2
>gi|1036830316|ref|XP_017138168.1| PREDICTED: proteasome subunit alpha type-1-like [Drosophila
>gi|1036830260|ref|XP_017138165.1| PREDICTED: zinc finger protein 48 isoform X1 [Drosophila mi
>gi|1036830279|ref|XP_017138166.1| PREDICTED: zinc finger protein 48 isoform X1 [Drosophila mi
>gi|1036830298|ref|XP_017138167.1| PREDICTED: zinc finger protein 48 isoform X2 [Drosophila mi
>gi|1036840407|ref|XP_017138739.1| PREDICTED: transcriptional protein SWT1 isoform X1 [Drosophi
>gi|1036840423|ref|XP_017138740.1| PREDICTED: transcriptional protein SWT1 isoform X2 [Drosophi
>gi|1036798087|ref|XP_017136377.1| PREDICTED: ADP-ribosylation factor 1-like isoform X2 [Drosop
>gi|1036798068|ref|XP_017136376.1| PREDICTED: ADP-ribosylation factor 1-like isoform X1 [Drosop
>gi|1036798105|ref|XP_017136378.1| PREDICTED: ADP-ribosylation factor 1-like isoform X1 [Drosop
>gi|1036797978|ref|XP_017136371.1| PREDICTED: COP9 signalosome complex subunit 3 isoform X1 [Di
>gi|1036797997|ref|XP_017136372.1| PREDICTED: COP9 signalosome complex subunit 3 isoform X2 [Di
>gi|1036798161|ref|XP_017136381.1| PREDICTED: uncharacterized protein LOC108151954 [Drosophila
>gi|1036798052|ref|XP_017136375.1| PREDICTED: uncharacterized protein LOC108151949 [Drosophila
>gi|1036798014|ref|XP_017136373.1| PREDICTED: cAMP-dependent protein kinase type I regulatory s
>gi|1036798033|ref|XP_017136374.1| PREDICTED: cAMP-dependent protein kinase type I regulatory s
>gi|1036798143|ref|XP_017136380.1| PREDICTED: uncharacterized protein LOC108151953 [Drosophila
>gi|1036798124|ref|XP_017136379.1| PREDICTED: uncharacterized protein LOC108151952 [Drosophila
>gi|1036797960|ref|XP_017136370.1| PREDICTED: uncharacterized protein LOC108151946 [Drosophila
>gi|1036800529|ref|XP_017136514.1| PREDICTED: neurofilament heavy polypeptide isoform X2 [Droso
>gi|1036800510|ref|XP_017136513.1| PREDICTED: uncharacterized protein LOC108152034 isoform X1
>gi|1036800565|ref|XP_017136518.1| PREDICTED: protein ST7 homolog [Drosophila miranda]
>gi|1036800548|ref|XP_017136515.1| PREDICTED: uncharacterized protein LOC108152035 [Drosophila
>gi|1036806686|ref|XP_017136858.1| PREDICTED: translocation protein SEC63 homolog [Drosophila r
>gi|1036806723|ref|XP_017136860.1| PREDICTED: neurofilament medium polypeptide isoform X2 [Dros
>gi|1036806704|ref|XP_017136859.1| PREDICTED: methyl-CpG-binding domain-containing protein 10 i
>gi|1036811459|ref|XP_017137118.1| PREDICTED: akirin [Drosophila miranda]
>gi|1036811478|ref|XP_017137120.1| PREDICTED: sphingomyelin synthase-related 1 [Drosophila mira
>gi|1036822326|ref|XP_017137714.1| PREDICTED: nuclear valosin-containing protein-like isoform X
>gi|1036822345|ref|XP_017137715.1| PREDICTED: nuclear valosin-containing protein-like isoform X
>gi|1036822364|ref|XP_017137716.1| PREDICTED: nuclear valosin-containing protein-like isoform X
>gi|1036822383|ref|XP_017137717.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036851560|ref|XP_017139356.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036764838|ref|XP_017157139.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036764003|ref|XP_017157099.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
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>gi|1036851726|ref|XP_017139366.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036850303|ref|XP_017139281.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036831754|ref|XP_017138250.1| PREDICTED: ras-related protein Rac2 [Drosophila miranda]
>gi|1036831716|ref|XP_017138248.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036831735|ref|XP_017138249.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036766702|ref|XP_017134589.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1036766270|ref|XP_017134563.1| PREDICTED: protein late bloomer isoform X1 [Drosophila mirar
>gi|1036766289|ref|XP_017134564.1| PREDICTED: protein late bloomer isoform X2 [Drosophila mirar
>gi|1036766308|ref|XP_017134565.1| PREDICTED: protein late bloomer isoform X3 [Drosophila mirar
>gi|1036769646|ref|XP_017134752.1| PREDICTED: cuticle protein 76 [Drosophila miranda]
>gi|1036768775|ref|XP_017134700.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
>gi|1036768517|ref|XP_017134686.1| PREDICTED: uncharacterized protein LOC108150888 isoform X1
>gi|1036768536|ref|XP_017134687.1| PREDICTED: uncharacterized protein LOC108150888 isoform X2
>gi|1036768558|ref|XP_017134688.1| PREDICTED: uncharacterized protein LOC108150888 isoform X3
>gi|1036787543|ref|XP_017135780.1| PREDICTED: probable cytochrome P450 316a1 [Drosophila mirano
>gi|1036787505|ref|XP_017135778.1| PREDICTED: cytochrome P450 4d8 isoform X1 [Drosophila mirano
>gi|1036787524|ref|XP_017135779.1| PREDICTED: cytochrome P450 4d8 isoform X2 [Drosophila mirano
>gi|1036787559|ref|XP_017135781.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036758835|ref|XP_017156802.1| PREDICTED: zinc carboxypeptidase-like [Drosophila miranda]
>gi|1036787577|ref|XP_017135782.1| PREDICTED: cuticle protein LPCP-23 [Drosophila miranda]
>gi|1036787372|ref|XP_017135770.1| PREDICTED: large proline-rich protein BAG6 isoform X1 [Droso
>gi|1036787391|ref|XP_017135772.1| PREDICTED: large proline-rich protein BAG6 isoform X1 [Droso
>gi|1036787410|ref|XP_017135773.1| PREDICTED: large proline-rich protein BAG6 isoform X1 [Droso
>gi|1036787446|ref|XP_017135775.1| PREDICTED: large proline-rich protein BAG6 isoform X3 [Droso
>gi|1036787429|ref|XP_017135774.1| PREDICTED: large proline-rich protein BAG6 isoform X2 [Droso
>gi|1036787467|ref|XP_017135776.1| PREDICTED: large proline-rich protein BAG6 isoform X4 [Droso
>gi|1036787486|ref|XP_017135777.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036814640|ref|XP_017137292.1| PREDICTED: epsin-1 isoform X1 [Drosophila miranda]
>gi|1036814661|ref|XP_017137293.1| PREDICTED: epsin-1 isoform X1 [Drosophila miranda]
>gi|1036814698|ref|XP_017137295.1| PREDICTED: epsin-1 isoform X3 [Drosophila miranda]
>gi|1036814717|ref|XP_017137296.1| PREDICTED: epsin-1 isoform X4 [Drosophila miranda]
>gi|1036814736|ref|XP_017137297.1| PREDICTED: epsin-1 isoform X5 [Drosophila miranda]
>gi|1036814680|ref|XP_017137294.1| PREDICTED: epsin-1 isoform X2 [Drosophila miranda]
>gi|1036764616|ref|XP_017157126.1| PREDICTED: trypsin [Drosophila miranda]
>gi|1036758851|ref|XP_017156803.1| PREDICTED: trypsin beta-like [Drosophila miranda]
>gi|1036758870|ref|XP_017156804.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036846846|ref|XP_017139081.1| PREDICTED: WD repeat-containing protein 63 [Drosophila mirar
>gi|1036850818|ref|XP_017139311.1| PREDICTED: gamma-tubulin complex component 2 homolog [Drosop
>gi|1036814926|ref|XP_017137308.1| PREDICTED: uncharacterized protein LOC108152455 [Drosophila
>gi|1036814850|ref|XP_017137304.1| PREDICTED: uncharacterized protein LOC108152453 isoform X1
>gi|1036814888|ref|XP_017137306.1| PREDICTED: protein late bloomer-like isoform X1 [Drosophila
>gi|1036814907|ref|XP_017137307.1| PREDICTED: uncharacterized protein LOC108152454 isoform X2
>gi|1036814869|ref|XP_017137305.1| PREDICTED: pleckstrin homology domain-containing family G me
>gi|1036814966|ref|XP_017137309.1| PREDICTED: ubiquitin-related modifier 1 homolog [Drosophila
>gi|1036836279|ref|XP_017138500.1| PREDICTED: uncharacterized protein LOC108153193 [Drosophila
>gi|1036836298|ref|XP_017138501.1| PREDICTED: transmembrane protein 70 homolog, mitochondrial
>gi|1036758889|ref|XP_017156805.1| PREDICTED: peptidoglycan-recognition protein SD [Drosophila
>gi|1036836598|ref|XP_017138520.1| PREDICTED: uncharacterized protein LOC108153209 [Drosophila
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>gi|1036798880|ref|XP_017136422.1| PREDICTED: ankyrin-3 isoform X5 [Drosophila miranda]
>gi|1036798842|ref|XP_017136420.1| PREDICTED: ankyrin-2 isoform X3 [Drosophila miranda]
>gi|1036798823|ref|XP_017136419.1| PREDICTED: ankyrin-2 isoform X2 [Drosophila miranda]
>gi|1036798787|ref|XP_017136417.1| PREDICTED: ankyrin-2 isoform X1 [Drosophila miranda]
>gi|1036798804|ref|XP_017136418.1| PREDICTED: ankyrin-2 isoform X1 [Drosophila miranda]
>gi|1036798961|ref|XP_017136427.1| PREDICTED: ankyrin-2 isoform X9 [Drosophila miranda]
>gi|1036798941|ref|XP_017136426.1| PREDICTED: ankyrin-2 isoform X8 [Drosophila miranda]
>gi|1036798998|ref|XP_017136429.1| PREDICTED: ankyrin-2 isoform X11 [Drosophila miranda]
>gi|1036798979|ref|XP_017136428.1| PREDICTED: ankyrin-2 isoform X10 [Drosophila miranda]
>gi|1036798922|ref|XP_017136425.1| PREDICTED: ankyrin-2 isoform X7 [Drosophila miranda]
>gi|1036798861|ref|XP_017136421.1| PREDICTED: ankyrin-2 isoform X4 [Drosophila miranda]
>gi|1036799016|ref|XP_017136430.1| PREDICTED: uncharacterized protein LOC108151976 [Drosophila
>gi|1036798899|ref|XP_017136423.1| PREDICTED: ankyrin-3 isoform X6 [Drosophila miranda]
>gi|1036794192|ref|XP_017136157.1| PREDICTED: probable medium-chain specific acyl-CoA dehydroge
>gi|1036794140|ref|XP_017136154.1| PREDICTED: tonsoku-like protein [Drosophila miranda]
>gi|1036794156|ref|XP_017136155.1| PREDICTED: zinc finger RNA-binding protein isoform X1 [Droso
>gi|1036794174|ref|XP_017136156.1| PREDICTED: zinc finger RNA-binding protein isoform X2 [Droso
>gi|1036819780|ref|XP_017137580.1| PREDICTED: integrator complex subunit 9 [Drosophila miranda]
>gi|1036819761|ref|XP_017137579.1| PREDICTED: isoleucine--tRNA ligase, mitochondrial [Drosophi
>gi|1036849676|ref|XP_017139244.1| PREDICTED: ATP synthase subunit beta, mitochondrial [Drosoph
>gi|1036778461|ref|XP_017135254.1| PREDICTED: uncharacterized protein LOC108151250 [Drosophila
>gi|1036778443|ref|XP_017135253.1| PREDICTED: H(+)/Cl(-) exchange transporter 3 [Drosophila mi
>gi|1036778479|ref|XP_017135255.1| PREDICTED: threonine aspartase 1 [Drosophila miranda]
>gi|1036778497|ref|XP_017135257.1| PREDICTED: uncharacterized protein LOC108151252 [Drosophila
>gi|1036778423|ref|XP_017135252.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778388|ref|XP_017135250.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778367|ref|XP_017135249.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A iso
>gi|1036778405|ref|XP_017135251.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778348|ref|XP_017135248.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778183|ref|XP_017135238.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A iso
>gi|1036778166|ref|XP_017135237.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778311|ref|XP_017135246.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778255|ref|XP_017135243.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778236|ref|XP_017135241.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778217|ref|XP_017135240.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A iso
>gi|1036778200|ref|XP_017135239.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778274|ref|XP_017135244.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778149|ref|XP_017135236.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778293|ref|XP_017135245.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778329|ref|XP_017135247.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036813691|ref|XP_017137237.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813616|ref|XP_017137232.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813635|ref|XP_017137233.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813653|ref|XP_017137234.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813672|ref|XP_017137236.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036830115|ref|XP_017138155.1| PREDICTED: uncharacterized protein LOC108152963 [Drosophila
>gi|1036830170|ref|XP_017138159.1| PREDICTED: uncharacterized protein LOC108152965 isoform X1
>gi|1036830205|ref|XP_017138162.1| PREDICTED: uncharacterized protein LOC108152965 isoform X3
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>gi|1036830187|ref|XP_017138160.1| PREDICTED: uncharacterized protein LOC108152965 isoform X2
>gi|1036830133|ref|XP_017138157.1| PREDICTED: uncharacterized protein LOC108152964 [Drosophila
>gi|1036830152|ref|XP_017138158.1| PREDICTED: uncharacterized protein LOC108152964 [Drosophila
>gi|1036824294|ref|XP_017137825.1| PREDICTED: uncharacterized protein LOC108152764 isoform X1
>gi|1036824313|ref|XP_017137827.1| PREDICTED: putative dipeptidase CPC735_014430 isoform X2 [Dr
>gi|1036824452|ref|XP_017137836.1| PREDICTED: mpv17-like protein 2 isoform X1 [Drosophila mira
>gi|1036824469|ref|XP_017137837.1| PREDICTED: mpv17-like protein 2 isoform X2 [Drosophila mira
>gi|1036824415|ref|XP_017137833.1| PREDICTED: uncharacterized protein LOC108152769 isoform X1
>gi|1036824434|ref|XP_017137835.1| PREDICTED: uncharacterized protein LOC108152769 isoform X2
>gi|1036824379|ref|XP_017137831.1| PREDICTED: uncharacterized protein LOC108152767 [Drosophila
>gi|1036824396|ref|XP_017137832.1| PREDICTED: peritrophin-44 [Drosophila miranda]
>gi|1036824362|ref|XP_017137830.1| PREDICTED: succinate dehydrogenase [ubiquinone] flavoprotei
>gi|1036824329|ref|XP_017137828.1| PREDICTED: T-related protein isoform X1 [Drosophila miranda]
>gi|1036824346|ref|XP_017137829.1| PREDICTED: T-related protein isoform X2 [Drosophila miranda]
>gi|1036851218|ref|XP_017139335.1| PREDICTED: endonuclease G, mitochondrial [Drosophila miranda
>gi|1036849544|ref|XP_017139236.1| PREDICTED: 26S protease regulatory subunit 10B [Drosophila n
>gi|1036849561|ref|XP_017139237.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036771620|ref|XP_017134861.1| PREDICTED: chondroitin proteoglycan 1 [Drosophila miranda]
>gi|1036771637|ref|XP_017134862.1| PREDICTED: chondroitin proteoglycan 2 [Drosophila miranda]
>gi|1036771748|ref|XP_017134868.1| PREDICTED: chondroitin proteoglycan 2 [Drosophila miranda]
>gi|1036771565|ref|XP_017134857.1| PREDICTED: uncharacterized protein LOC108151010 isoform X1
>gi|1036771601|ref|XP_017134860.1| PREDICTED: uncharacterized protein LOC108151010 isoform X3
>gi|1036771584|ref|XP_017134858.1| PREDICTED: uncharacterized protein LOC108151010 isoform X2
>gi|1036771767|ref|XP_017134869.1| PREDICTED: uncharacterized protein LOC108151021 [Drosophila
>gi|1036758905|ref|XP_017156806.1| PREDICTED: uncharacterized protein LOC108165282 [Drosophila
>gi|1036771471|ref|XP_017134852.1| PREDICTED: glutenin, high molecular weight subunit DX5 isofc
>gi|1036771452|ref|XP_017134851.1| PREDICTED: uncharacterized PE-PGRS family protein PE_PGRS54
>gi|1036771545|ref|XP_017134856.1| PREDICTED: flocculation protein FLO11 isoform X3 [Drosophila
>gi|1036771508|ref|XP_017134854.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036771526|ref|XP_017134855.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036771821|ref|XP_017134873.1| PREDICTED: peritrophin-1-like [Drosophila miranda]
>gi|1036771802|ref|XP_017134872.1| PREDICTED: integumentary mucin B.1-like [Drosophila miranda]
>gi|1036771269|ref|XP_017134840.1| PREDICTED: uncharacterized protein LOC108151001 [Drosophila
>gi|1036771489|ref|XP_017134853.1| PREDICTED: anoctamin-4 [Drosophila miranda]
>gi|1036771729|ref|XP_017134867.1| PREDICTED: tubulin polyglutamylase complex subunit 2 [Drosop
>gi|1036771786|ref|XP_017134871.1| PREDICTED: viral IAP-associated factor homolog [Drosophila n
>gi|1036771435|ref|XP_017134850.1| PREDICTED: uncharacterized protein LOC108151005 [Drosophila
>gi|1036771872|ref|XP_017134875.1| PREDICTED: cAMP-regulated phosphoprotein 19 [Drosophila mira
>gi|1036771287|ref|XP_017134841.1| PREDICTED: tuberin [Drosophila miranda]
>gi|1036771710|ref|XP_017134866.1| PREDICTED: uncharacterized protein LOC108151016 [Drosophila
>gi|1036771656|ref|XP_017134863.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036771692|ref|XP_017134865.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036771400|ref|XP_017134848.1| PREDICTED: uncharacterized protein LOC108151004 isoform X4
>gi|1036771342|ref|XP_017134844.1| PREDICTED: uncharacterized protein LOC108151004 isoform X3
>gi|1036771361|ref|XP_017134845.1| PREDICTED: uncharacterized protein LOC108151004 isoform X3
>gi|1036771380|ref|XP_017134847.1| PREDICTED: uncharacterized protein LOC108151004 isoform X3
>gi|1036771323|ref|XP_017134843.1| PREDICTED: uncharacterized protein LOC108151004 isoform X2
>gi|1036771304|ref|XP_017134842.1| PREDICTED: uncharacterized protein LOC108151004 isoform X1
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>gi|1036771418|ref|XP_017134849.1| PREDICTED: uncharacterized protein LOC108151004 isoform X5
>gi|1036771840|ref|XP_017134874.1| PREDICTED: uncharacterized protein LOC108151025 [Drosophila
>gi|1036771891|ref|XP_017134876.1| PREDICTED: uncharacterized protein LOC108151027 [Drosophila
>gi|1036771674|ref|XP_017134864.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036764876|ref|XP_017157141.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036849232|ref|XP_017139216.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036850651|ref|XP_017139302.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036765507|ref|XP_017157175.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036764281|ref|XP_017157112.1| PREDICTED: peritrophin-44 isoform X2 [Drosophila miranda]
>gi|1036764261|ref|XP_017157111.1| PREDICTED: peritrophin-44 isoform X1 [Drosophila miranda]
>gi|1036764965|ref|XP_017157144.1| PREDICTED: peritrophin-44 [Drosophila miranda]
>gi|1036824848|ref|XP_017137857.1| PREDICTED: peritrophin-44 [Drosophila miranda]
>gi|1036824808|ref|XP_017137855.1| PREDICTED: possible lysine-specific histone demethylase 1 [D
>gi|1036824829|ref|XP_017137856.1| PREDICTED: possible lysine-specific histone demethylase 1 [D
>gi|1036800896|ref|XP_017136536.1| PREDICTED: uncharacterized protein LOC108152044 [Drosophila
>gi|1036800861|ref|XP_017136534.1| PREDICTED: neurogenic protein mastermind isoform X3 [Drosoph
>gi|1036800842|ref|XP_017136533.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036800804|ref|XP_017136531.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036800823|ref|XP_017136532.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036800880|ref|XP_017136535.1| PREDICTED: myb-like protein Q isoform X4 [Drosophila miranda
>gi|1036800936|ref|XP_017136538.1| PREDICTED: probable deoxycytidylate deaminase [Drosophila mi
>gi|1036800917|ref|XP_017136537.1| PREDICTED: clathrin light chain [Drosophila miranda]
>gi|1036758923|ref|XP_017156807.1| PREDICTED: spore coat protein SP96-like [Drosophila miranda]
>gi|1036845837|ref|XP_017139023.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036845800|ref|XP_017139021.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036845818|ref|XP_017139022.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036845856|ref|XP_017139024.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036845875|ref|XP_017139025.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036845892|ref|XP_017139027.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036845911|ref|XP_017139028.1| PREDICTED: serine/threonine-protein phosphatase rdgC isoform
>gi|1036786846|ref|XP_017135742.1| PREDICTED: uncharacterized protein LOC108151577 [Drosophila
>gi|1036786864|ref|XP_017135743.1| PREDICTED: uncharacterized protein C9orf117 homolog [Drosoph
>gi|1036786917|ref|XP_017135746.1| PREDICTED: protein FRG1 homolog [Drosophila miranda]
>gi|1036786827|ref|XP_017135741.1| PREDICTED: tectonin beta-propeller repeat-containing protei
>gi|1036786932|ref|XP_017135747.1| PREDICTED: DCN1-like protein 5 isoform X1 [Drosophila miran
>gi|1036786968|ref|XP_017135749.1| PREDICTED: DCN1-like protein 5 isoform X1 [Drosophila miran
>gi|1036786901|ref|XP_017135745.1| PREDICTED: lysosome-associated membrane glycoprotein 5 isof
>gi|1036786883|ref|XP_017135744.1| PREDICTED: lysosome-associated membrane glycoprotein 5 isof
>gi|1036786987|ref|XP_017135750.1| PREDICTED: DCN1-like protein 4 isoform X2 [Drosophila miran
>gi|1036788166|ref|XP_017135816.1| PREDICTED: alpha-soluble NSF attachment protein [Drosophila
>gi|1036788111|ref|XP_017135813.1| PREDICTED: leukocyte elastase inhibitor [Drosophila miranda]
>gi|1036788094|ref|XP_017135812.1| PREDICTED: retinoblastoma-binding protein 5 homolog [Drosoph
>gi|1036788130|ref|XP_017135814.1| PREDICTED: eukaryotic peptide chain release factor subunit
>gi|1036788148|ref|XP_017135815.1| PREDICTED: eukaryotic peptide chain release factor subunit
>gi|1036788075|ref|XP_017135811.1| PREDICTED: cysteine sulfinic acid decarboxylase [Drosophila
>gi|1036822962|ref|XP_017137750.1| PREDICTED: DNA/RNA-binding protein KIN17 [Drosophila miranda
>gi|1036822981|ref|XP_017137751.1| PREDICTED: LOW QUALITY PROTEIN: vitellogenin-3 [Drosophila m
>gi|1036822944|ref|XP_017137749.1| PREDICTED: DNA primase large subunit [Drosophila miranda]
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>gi|1036832401|ref|XP_017138284.1| PREDICTED: toll-like receptor 1 isoform X4 [Drosophila mira
>gi|1036832361|ref|XP_017138282.1| PREDICTED: toll-like receptor 1 isoform X2 [Drosophila mira
>gi|1036832382|ref|XP_017138283.1| PREDICTED: toll-like receptor 1 isoform X3 [Drosophila mira
>gi|1036832342|ref|XP_017138280.1| PREDICTED: toll-like receptor 1 isoform X1 [Drosophila mira
>gi|1036832419|ref|XP_017138285.1| PREDICTED: protein inturned [Drosophila miranda]
>gi|1036832456|ref|XP_017138288.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036832438|ref|XP_017138287.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036846436|ref|XP_017139057.1| PREDICTED: serine protease 3 [Drosophila miranda]
>gi|1036810377|ref|XP_017137060.1| PREDICTED: cyclin-T [Drosophila miranda]
>gi|1036828764|ref|XP_017138079.1| PREDICTED: ecdysone-induced protein 74EF isoform X2 [Drosoph
>gi|1036828745|ref|XP_017138078.1| PREDICTED: ecdysone-induced protein 74EF isoform X1 [Drosoph
>gi|1036828781|ref|XP_017138080.1| PREDICTED: rho GTPase-activating protein gacII-like [Drosoph
>gi|1036845048|ref|XP_017138985.1| PREDICTED: charged multivesicular body protein 5 [Drosophila
>gi|1036802166|ref|XP_017136610.1| PREDICTED: uncharacterized protein LOC108152084 isoform X1
>gi|1036802185|ref|XP_017136611.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila m
>gi|1036802242|ref|XP_017136614.1| PREDICTED: protein melted isoform X2 [Drosophila miranda]
>gi|1036802204|ref|XP_017136612.1| PREDICTED: protein melted isoform X1 [Drosophila miranda]
>gi|1036802223|ref|XP_017136613.1| PREDICTED: protein melted isoform X1 [Drosophila miranda]
>gi|1036802263|ref|XP_017136616.1| PREDICTED: acyl-CoA-binding protein homolog [Drosophila mira
>gi|1036765957|ref|XP_017134549.1| PREDICTED: acyl-CoA-binding protein-like [Drosophila miranda
>gi|1036849714|ref|XP_017139246.1| PREDICTED: acyl-CoA-binding protein [Drosophila miranda]
>gi|1036849731|ref|XP_017139247.1| PREDICTED: acyl-CoA-binding protein homolog [Drosophila mira
>gi|1036849695|ref|XP_017139245.1| PREDICTED: Bardet-Biedl syndrome 1 protein [Drosophila miran
>gi|1036825035|ref|XP_017137866.1| PREDICTED: protein male-specific lethal-3 [Drosophila mirand
>gi|1036825020|ref|XP_017137865.1| PREDICTED: FH2 domain-containing protein 1 [Drosophila miran
>gi|1036758939|ref|XP_017156808.1| PREDICTED: uncharacterized protein LOC108165284 [Drosophila
>gi|1036768403|ref|XP_017134679.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036767849|ref|XP_017134649.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036767870|ref|XP_017134650.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036848986|ref|XP_017139202.1| PREDICTED: uncharacterized protein LOC108153615 isoform X1
>gi|1036849005|ref|XP_017139203.1| PREDICTED: uncharacterized protein LOC108153615 isoform X2
>gi|1036849045|ref|XP_017139205.1| PREDICTED: uncharacterized protein LOC108153616 isoform X2
>gi|1036849024|ref|XP_017139204.1| PREDICTED: uncharacterized protein LOC108153616 isoform X1
>gi|1036848967|ref|XP_017139201.1| PREDICTED: GTP-binding protein Rhes [Drosophila miranda]
>gi|1036838898|ref|XP_017138661.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X9 [Drosoph
>gi|1036838861|ref|XP_017138658.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X7 [Drosoph
>gi|1036838880|ref|XP_017138659.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X8 [Drosoph
>gi|1036838842|ref|XP_017138657.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X6 [Drosoph
>gi|1036838823|ref|XP_017138656.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X5 [Drosoph
>gi|1036838806|ref|XP_017138655.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X4 [Drosoph
>gi|1036838787|ref|XP_017138654.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X3 [Drosoph
>gi|1036838768|ref|XP_017138653.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X2 [Drosoph
>gi|1036838730|ref|XP_017138651.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X1 [Drosoph
>gi|1036838749|ref|XP_017138652.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X1 [Drosoph
>gi|1036838971|ref|XP_017138665.1| PREDICTED: collagen alpha-1(XV) chain isoform X13 [Drosophil
>gi|1036838953|ref|XP_017138664.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X12 [Drosop
>gi|1036838934|ref|XP_017138663.1| PREDICTED: collagen alpha-1(XV) chain isoform X11 [Drosophil
>gi|1036838915|ref|XP_017138662.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X10 [Drosop
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>gi|1036838990|ref|XP_017138666.1| PREDICTED: uncharacterized protein LOC108153289 [Drosophila
>gi|1036812163|ref|XP_017137156.1| PREDICTED: differentially expressed in FDCP 8 homolog [Drosc
>gi|1036812123|ref|XP_017137154.1| PREDICTED: protein KRI1 homolog [Drosophila miranda]
>gi|1036812142|ref|XP_017137155.1| PREDICTED: protein KRI1 homolog [Drosophila miranda]
>gi|1036812201|ref|XP_017137159.1| PREDICTED: autophagy protein 12-like [Drosophila miranda]
>gi|1036812182|ref|XP_017137158.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036774472|ref|XP_017135027.1| PREDICTED: rho GTPase-activating protein 68F isoform X1 [Dro
>gi|1036774491|ref|XP_017135028.1| PREDICTED: rho GTPase-activating protein 68F isoform X1 [Dro
>gi|1036774510|ref|XP_017135029.1| PREDICTED: rho GTPase-activating protein 68F isoform X2 [Dro
>gi|1036774432|ref|XP_017135025.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036774340|ref|XP_017135019.1| PREDICTED: neurexin-4 isoform X3 [Drosophila miranda]
>gi|1036774359|ref|XP_017135020.1| PREDICTED: neurexin-4 isoform X4 [Drosophila miranda]
>gi|1036774302|ref|XP_017135017.1| PREDICTED: neurexin-4 isoform X1 [Drosophila miranda]
>gi|1036774321|ref|XP_017135018.1| PREDICTED: neurexin-4 isoform X2 [Drosophila miranda]
>gi|1036774565|ref|XP_017135033.1| PREDICTED: L-aminoadipate-semialdehyde dehydrogenase-phospho
>gi|1036774597|ref|XP_017135036.1| PREDICTED: uncharacterized protein LOC108151126 [Drosophila
>gi|1036774451|ref|XP_017135026.1| PREDICTED: opsin, ultraviolet-sensitive [Drosophila miranda]
>gi|1036774547|ref|XP_017135032.1| PREDICTED: carbonic anhydrase 2 [Drosophila miranda]
>gi|1036774583|ref|XP_017135034.1| PREDICTED: inositol oxygenase [Drosophila miranda]
>gi|1036774414|ref|XP_017135024.1| PREDICTED: esterase-5C [Drosophila miranda]
>gi|1036774397|ref|XP_017135022.1| PREDICTED: esterase-5B [Drosophila miranda]
>gi|1036774378|ref|XP_017135021.1| PREDICTED: esterase-5A isoform X5 [Drosophila miranda]
>gi|1036774529|ref|XP_017135030.1| PREDICTED: THO complex subunit 6 [Drosophila miranda]
>gi|1036851872|ref|XP_017139375.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036827881|ref|XP_017138030.1| PREDICTED: adenylate kinase isoenzyme 1-like isoform X2 [Dro
>gi|1036827863|ref|XP_017138029.1| PREDICTED: adenylate kinase isoenzyme 1-like isoform X1 [Dro
>gi|1036827904|ref|XP_017138032.1| PREDICTED: adenylate kinase isoenzyme 1-like [Drosophila mi
>gi|1036827768|ref|XP_017138023.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827787|ref|XP_017138024.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827806|ref|XP_017138025.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827825|ref|XP_017138027.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827844|ref|XP_017138028.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036804750|ref|XP_017136753.1| PREDICTED: probable protein S-acyltransferase 4 [Drosophila
>gi|1036804768|ref|XP_017136754.1| PREDICTED: dnaJ protein ERDJ2A-like [Drosophila miranda]
>gi|1036830663|ref|XP_017138189.1| PREDICTED: uncharacterized protein LOC108152974 [Drosophila
>gi|1036830684|ref|XP_017138190.1| PREDICTED: delta-aminolevulinic acid dehydratase [Drosophila
>gi|1036824752|ref|XP_017137853.1| PREDICTED: spermine synthase isoform X1 [Drosophila miranda]
>gi|1036824771|ref|XP_017137854.1| PREDICTED: spermine synthase isoform X2 [Drosophila miranda]
>gi|1036840223|ref|XP_017138729.1| PREDICTED: 72 kDa inositol polyphosphate 5-phosphatase [Dros
>gi|1036764987|ref|XP_017157145.1| PREDICTED: LIM homeobox transcription factor 1-beta [Drosoph
>gi|1036849511|ref|XP_017139233.1| PREDICTED: LIM homeobox transcription factor 1-beta [Drosoph
>gi|1036842423|ref|XP_017138844.1| PREDICTED: uncharacterized protein LOC108153383 [Drosophila
>gi|1036842441|ref|XP_017138845.1| PREDICTED: uncharacterized protein LOC108153383 [Drosophila
>gi|1036819818|ref|XP_017137582.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso:
>gi|1036819892|ref|XP_017137585.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso:
>gi|1036819873|ref|XP_017137584.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso:
>gi|1036819911|ref|XP_017137586.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso:
>gi|1036827671|ref|XP_017138017.1| PREDICTED: ras-GEF domain-containing family member 1B-B [Dro
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>gi|1036841699|ref|XP_017138815.1| PREDICTED: growth arrest and DNA damage-inducible proteins-
>gi|1036841685|ref|XP_017138814.1| PREDICTED: uncharacterized protein LOC108153359 [Drosophila
>gi|1036842894|ref|XP_017138872.1| PREDICTED: neuropeptides capa receptor isoform X1 [Drosophil
>gi|1036842913|ref|XP_017138873.1| PREDICTED: neuropeptides capa receptor isoform X1 [Drosophil
>gi|1036842950|ref|XP_017138876.1| PREDICTED: neuropeptides capa receptor isoform X3 [Drosophil
>gi|1036842931|ref|XP_017138875.1| PREDICTED: neuropeptides capa receptor isoform X2 [Drosophil
>gi|1036851128|ref|XP_017139329.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036817942|ref|XP_017137479.1| PREDICTED: hemicentin-1 isoform X7 [Drosophila miranda]
>gi|1036817965|ref|XP_017137480.1| PREDICTED: uncharacterized protein LOC108152559 isoform X8
>gi|1036817923|ref|XP_017137478.1| PREDICTED: hemicentin-1 isoform X6 [Drosophila miranda]
>gi|1036817906|ref|XP_017137477.1| PREDICTED: uncharacterized protein LOC108152559 isoform X5
>gi|1036817871|ref|XP_017137475.1| PREDICTED: uncharacterized protein LOC108152559 isoform X3
>gi|1036817855|ref|XP_017137473.1| PREDICTED: uncharacterized protein LOC108152559 isoform X2
>gi|1036817840|ref|XP_017137472.1| PREDICTED: uncharacterized protein LOC108152559 isoform X1
>gi|1036817888|ref|XP_017137476.1| PREDICTED: uncharacterized protein LOC108152559 isoform X4
>gi|1036758957|ref|XP_017156809.1| PREDICTED: muscle M-line assembly protein unc-89-like [Droso
>gi|1036817984|ref|XP_017137481.1| PREDICTED: uncharacterized protein LOC108152560 [Drosophila
>gi|1036827432|ref|XP_017138003.1| PREDICTED: uncharacterized protein LOC108152881 [Drosophila
>gi|1036851309|ref|XP_017139340.1| PREDICTED: cathepsin L1 [Drosophila miranda]
>gi|1036846093|ref|XP_017139038.1| PREDICTED: uncharacterized protein LOC108153498 [Drosophila
>gi|1036767596|ref|XP_017134638.1| PREDICTED: uncharacterized protein LOC108150840 [Drosophila
>gi|1036782830|ref|XP_017135511.1| PREDICTED: cationic amino acid transporter 3 isoform X1 [Dro
>gi|1036782849|ref|XP_017135512.1| PREDICTED: cationic amino acid transporter 3 isoform X2 [Dro
>gi|1036782867|ref|XP_017135513.1| PREDICTED: cationic amino acid transporter 3 isoform X2 [Dro
>gi|1036782884|ref|XP_017135514.1| PREDICTED: cationic amino acid transporter 3 isoform X2 [Dro
>gi|1036782902|ref|XP_017135515.1| PREDICTED: general transcription factor IIH subunit 2 [Droso
>gi|1036782795|ref|XP_017135509.1| PREDICTED: uncharacterized protein LOC108151428 [Drosophila
>gi|1036782813|ref|XP_017135510.1| PREDICTED: uncharacterized protein LOC108151428 [Drosophila
>gi|1036782921|ref|XP_017135516.1| PREDICTED: putative methyltransferase NSUN6 [Drosophila mira
>gi|1036782938|ref|XP_017135517.1| PREDICTED: ADP-ribosylation factor 1 [Drosophila miranda]
>gi|1036782957|ref|XP_017135518.1| PREDICTED: ADP-ribosylation factor 1 [Drosophila miranda]
>gi|1036831611|ref|XP_017138243.1| PREDICTED: ER membrane protein complex subunit 4 [Drosophila
>gi|1036831648|ref|XP_017138246.1| PREDICTED: uncharacterized protein LOC108153010 isoform X2
>gi|1036831629|ref|XP_017138245.1| PREDICTED: uncharacterized protein LOC108153010 isoform X1
>gi|1036831667|ref|XP_017138247.1| PREDICTED: single-pass membrane and coiled-coil domain-conta
>gi|1036766868|ref|XP_017134598.1| PREDICTED: uncharacterized protein LOC108150806 [Drosophila
>gi|1036758975|ref|XP_017156810.1| PREDICTED: oleosin-B6 [Drosophila miranda]
>gi|1036850074|ref|XP_017139269.1| PREDICTED: aspartic and glutamic acid-rich protein [Drosophi
>gi|1036850093|ref|XP_017139270.1| PREDICTED: uncharacterized protein LOC108153667 [Drosophila
>gi|1036824174|ref|XP_017137819.1| PREDICTED: myeloid zinc finger 1 isoform X1 [Drosophila mira
>gi|1036824195|ref|XP_017137820.1| PREDICTED: myeloid zinc finger 1 isoform X2 [Drosophila mira
>gi|1036824214|ref|XP_017137821.1| PREDICTED: myeloid zinc finger 1 isoform X3 [Drosophila mira
>gi|1036824237|ref|XP_017137822.1| PREDICTED: myeloid zinc finger 1 isoform X4 [Drosophila mira
>gi|1036824256|ref|XP_017137823.1| PREDICTED: zinc finger protein 768 isoform X5 [Drosophila mi
>gi|1036824275|ref|XP_017137824.1| PREDICTED: uncharacterized protein LOC108152763 [Drosophila
>gi|1036763048|ref|XP_017157049.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm2 [Drosopl
>gi|1036846165|ref|XP_017139042.1| PREDICTED: gonadotropin-releasing hormone receptor [Drosophi
>gi|1036831227|ref|XP_017138221.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
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>gi|1036831208|ref|XP_017138220.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1036831172|ref|XP_017138218.1| PREDICTED: solute carrier family 12 member 2 isoform X1 [Dro
>gi|1036831191|ref|XP_017138219.1| PREDICTED: solute carrier family 12 member 2 isoform X1 [Dro
>gi|1036817463|ref|XP_017137450.1| PREDICTED: RNA polymerase II-associated protein 1 [Drosophil
>gi|1036817519|ref|XP_017137454.1| PREDICTED: general odorant-binding protein 69a [Drosophila m
>gi|1036817481|ref|XP_017137452.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosoph
>gi|1036817501|ref|XP_017137453.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosoph
>gi|1036826387|ref|XP_017137941.1| PREDICTED: calcium-binding mitochondrial carrier protein SCa
>gi|1036826406|ref|XP_017137942.1| PREDICTED: calcium-binding mitochondrial carrier protein SCa
>gi|1036826424|ref|XP_017137943.1| PREDICTED: probable calcium-binding mitochondrial carrier Cl
>gi|1036826442|ref|XP_017137944.1| PREDICTED: probable calcium-binding mitochondrial carrier Cl
>gi|1036836023|ref|XP_017138485.1| PREDICTED: ADP-ribosylation factor GTPase-activating protein
>gi|1036836004|ref|XP_017138484.1| PREDICTED: ADP-ribosylation factor GTPase-activating protein
>gi|1036847661|ref|XP_017139127.1| PREDICTED: paired box protein Pax-6 isoform X1 [Drosophila m
>gi|1036847680|ref|XP_017139128.1| PREDICTED: paired box protein Pax-1 isoform X2 [Drosophila m
>gi|1036846398|ref|XP_017139055.1| PREDICTED: paired box protein Pax-6 [Drosophila miranda]
>gi|1036768499|ref|XP_017134685.1| PREDICTED: cuticle protein 38-like [Drosophila miranda]
>gi|1036768575|ref|XP_017134689.1| PREDICTED: uncharacterized protein LOC108150890 [Drosophila
>gi|1036766481|ref|XP_017134577.1| PREDICTED: uncharacterized protein LOC108150787 [Drosophila
>gi|1036828127|ref|XP_017138043.1| PREDICTED: protein odr-4 homolog [Drosophila miranda]
>gi|1036828091|ref|XP_017138040.1| PREDICTED: uncharacterized protein LOC108152905 [Drosophila
>gi|1036828110|ref|XP_017138041.1| PREDICTED: uncharacterized protein LOC108152905 [Drosophila
>gi|1036847085|ref|XP_017139095.1| PREDICTED: uncharacterized protein LOC108153540 isoform X1
>gi|1036847104|ref|XP_017139096.1| PREDICTED: uncharacterized protein LOC108153540 isoform X2
>gi|1036848931|ref|XP_017139199.1| PREDICTED: clavesin-1 [Drosophila miranda]
>gi|1036780551|ref|XP_017135379.1| PREDICTED: 1,5-anhydro-D-fructose reductase isoform X2 [Dros
>gi|1036780532|ref|XP_017135378.1| PREDICTED: 1,5-anhydro-D-fructose reductase isoform X1 [Dros
>gi|1036780570|ref|XP_017135380.1| PREDICTED: aldose reductase isoform X3 [Drosophila miranda]
>gi|1036780589|ref|XP_017135381.1| PREDICTED: kxDL motif-containing protein CG10681 isoform X1
>gi|1036780607|ref|XP_017135383.1| PREDICTED: kxDL motif-containing protein CG10681 isoform X2
>gi|1036780514|ref|XP_017135377.1| PREDICTED: protein TSSC1 [Drosophila miranda]
>gi|1036780625|ref|XP_017135384.1| PREDICTED: ubiquitin-conjugating enzyme E2 C [Drosophila mi
>gi|1036780498|ref|XP_017135376.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1036780479|ref|XP_017135375.1| PREDICTED: citron Rho-interacting kinase isoform X2 [Drosoph
>gi|1036780461|ref|XP_017135374.1| PREDICTED: citron Rho-interacting kinase isoform X1 [Drosoph
>gi|1036781203|ref|XP_017135419.1| PREDICTED: protein LSM14 homolog B-A isoform X1 [Drosophila
>gi|1036781222|ref|XP_017135420.1| PREDICTED: protein LSM14 homolog B-A isoform X2 [Drosophila
>gi|1036781184|ref|XP_017135418.1| PREDICTED: histone acetyltransferase KAT2A [Drosophila mira
>gi|1036781278|ref|XP_017135423.1| PREDICTED: uncharacterized protein LOC108151372 [Drosophila
>gi|1036781259|ref|XP_017135422.1| PREDICTED: eukaryotic translation initiation factor 2 subun
>gi|1036781240|ref|XP_017135421.1| PREDICTED: phosphoacetylglucosamine mutase [Drosophila mira
>gi|1036785067|ref|XP_017135642.1| PREDICTED: melanotransferrin [Drosophila miranda]
>gi|1036785159|ref|XP_017135647.1| PREDICTED: uncharacterized protein LOC108151514 [Drosophila
>gi|1036785141|ref|XP_017135646.1| PREDICTED: probable phosphomannomutase [Drosophila miranda]
>gi|1036785084|ref|XP_017135643.1| PREDICTED: kelch domain-containing protein 4 [Drosophila mi
>gi|1036785103|ref|XP_017135644.1| PREDICTED: uncharacterized protein LOC108151510 [Drosophila
>gi|1036785122|ref|XP_017135645.1| PREDICTED: ankyrin repeat domain-containing protein SOWAHA-1
>gi|1036843319|ref|XP_017138896.1| PREDICTED: homeobox protein araucan [Drosophila miranda]
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>gi|1036837311|ref|XP_017138561.1| PREDICTED: homeobox protein caupolican [Drosophila miranda]
>gi|1036832127|ref|XP_017138271.1| PREDICTED: homeobox protein caupolican [Drosophila miranda]
>gi|1036834640|ref|XP_017138403.1| PREDICTED: protein ABHD18 [Drosophila miranda]
>gi|1036812315|ref|XP_017137165.1| PREDICTED: tyrosine-protein phosphatase 69D [Drosophila mira
>gi|1036796922|ref|XP_017136311.1| PREDICTED: uncharacterized protein LOC108151911 [Drosophila
>gi|1036796937|ref|XP_017136312.1| PREDICTED: probable small nuclear ribonucleoprotein Sm D1 [
>gi|1036796871|ref|XP_017136308.1| PREDICTED: kinesin light chain [Drosophila miranda]
>gi|1036796888|ref|XP_017136309.1| PREDICTED: kinesin light chain [Drosophila miranda]
>gi|1036796855|ref|XP_017136307.1| PREDICTED: ankyrin repeat domain-containing protein 11 [Dros
>gi|1036796905|ref|XP_017136310.1| PREDICTED: hsp70 nucleotide exchange factor fes1 [Drosophila
>gi|1036781387|ref|XP_017135430.1| PREDICTED: vacuolar protein sorting-associated protein 13D
>gi|1036781425|ref|XP_017135432.1| PREDICTED: huntingtin-interacting protein 1 isoform X2 [Dros
>gi|1036781406|ref|XP_017135431.1| PREDICTED: huntingtin-interacting protein 1 isoform X1 [Dros
>gi|1036758994|ref|XP_017156811.1| PREDICTED: ATP-binding cassette sub-family A member 17-like
>gi|1036759010|ref|XP_017156812.1| PREDICTED: uncharacterized protein LOC108165288 [Drosophila
>gi|1036759029|ref|XP_017156813.1| PREDICTED: protein SDA1 homolog [Drosophila miranda]
>gi|1036828835|ref|XP_017138082.1| PREDICTED: calcium channel flower isoform X1 [Drosophila mi
>gi|1036828853|ref|XP_017138083.1| PREDICTED: calcium channel flower isoform X2 [Drosophila mi
>gi|1036799054|ref|XP_017136432.1| PREDICTED: zinc finger protein 140 [Drosophila miranda]
>gi|1036799035|ref|XP_017136431.1| PREDICTED: mRNA-decapping enzyme subunit 2 [Drosophila mira
>gi|1036825673|ref|XP_017137899.1| PREDICTED: kelch-like protein diablo [Drosophila miranda]
>gi|1036834122|ref|XP_017138375.1| PREDICTED: uncharacterized protein C15orf41 homolog [Drosop]
>gi|1036834138|ref|XP_017138376.1| PREDICTED: zinc finger protein 706-like [Drosophila miranda]
>gi|1036834157|ref|XP_017138377.1| PREDICTED: zinc finger protein 706-like [Drosophila miranda]
>gi|1036847586|ref|XP_017139122.1| PREDICTED: uncharacterized protein LOC108153561 isoform X1
>gi|1036847605|ref|XP_017139123.1| PREDICTED: uncharacterized protein LOC108153561 isoform X1
>gi|1036847622|ref|XP_017139124.1| PREDICTED: uncharacterized protein LOC108153561 isoform X2
>gi|1036840742|ref|XP_017138758.1| PREDICTED: uncharacterized protein LOC108153334 isoform X11
>gi|1036840704|ref|XP_017138756.1| PREDICTED: uncharacterized protein LOC108153334 isoform X9
>gi|1036840614|ref|XP_017138750.1| PREDICTED: uncharacterized protein LOC108153334 isoform X4
>gi|1036840668|ref|XP_017138754.1| PREDICTED: uncharacterized protein LOC108153334 isoform X7
>gi|1036840723|ref|XP_017138757.1| PREDICTED: uncharacterized protein LOC108153334 isoform X10
>gi|1036840649|ref|XP_017138753.1| PREDICTED: uncharacterized protein LOC108153334 isoform X6
>gi|1036840481|ref|XP_017138743.1| PREDICTED: uncharacterized protein LOC108153334 isoform X2
>gi|1036840779|ref|XP_017138760.1| PREDICTED: gamma-aminobutyric acid receptor subunit alpha-3
>gi|1036840687|ref|XP_017138755.1| PREDICTED: uncharacterized protein LOC108153334 isoform X8
>gi|1036840760|ref|XP_017138759.1| PREDICTED: gamma-aminobutyric acid receptor subunit alpha-3
>gi|1036840633|ref|XP_017138752.1| PREDICTED: uncharacterized protein LOC108153334 isoform X5
>gi|1036840595|ref|XP_017138749.1| PREDICTED: uncharacterized protein LOC108153334 isoform X3
>gi|1036840444|ref|XP_017138741.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840462|ref|XP_017138742.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840500|ref|XP_017138744.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840517|ref|XP_017138745.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840536|ref|XP_017138746.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840557|ref|XP_017138747.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840576|ref|XP_017138748.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036844916|ref|XP_017138978.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X6
>gi|1036844879|ref|XP_017138976.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X4
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>gi|1036844897|ref|XP_017138977.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X5
>gi|1036844858|ref|XP_017138975.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X3
>gi|1036844837|ref|XP_017138974.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X2
>gi|1036844784|ref|XP_017138971.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X1
>gi|1036844803|ref|XP_017138972.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X1
>gi|1036844819|ref|XP_017138973.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X1
>gi|1036845067|ref|XP_017138986.1| PREDICTED: protein kinase DC2 isoform X1 [Drosophila miranda
>gi|1036845082|ref|XP_017138987.1| PREDICTED: protein kinase DC2 isoform X1 [Drosophila miranda
>gi|1036845099|ref|XP_017138988.1| PREDICTED: protein kinase DC2 isoform X2 [Drosophila miranda
>gi|1036845390|ref|XP_017139006.1| PREDICTED: group XIIA secretory phospholipase A2 [Drosophila
>gi|1036835833|ref|XP_017138473.1| PREDICTED: E3 ubiquitin-protein ligase NRDP1 [Drosophila mi
>gi|1036835852|ref|XP_017138474.1| PREDICTED: uncharacterized protein LOC108153177 [Drosophila
>gi|1036799217|ref|XP_017136442.1| PREDICTED: putative U5 small nuclear ribonucleoprotein 200
>gi|1036797300|ref|XP_017136333.1| PREDICTED: chromosomal serine/threonine-protein kinase JIL-
>gi|1036797319|ref|XP_017136334.1| PREDICTED: chromosomal serine/threonine-protein kinase JIL-
>gi|1036834679|ref|XP_017138405.1| PREDICTED: iodotyrosine deiodinase 1-like [Drosophila miran
>gi|1036834661|ref|XP_017138404.1| PREDICTED: uncharacterized protein LOC108153126 [Drosophila
>gi|1036834714|ref|XP_017138407.1| PREDICTED: uncharacterized protein LOC108153129 [Drosophila
>gi|1036834696|ref|XP_017138406.1| PREDICTED: CCAAT/enhancer-binding protein gamma [Drosophila
>gi|1036841595|ref|XP_017138809.1| PREDICTED: nedd8-activating enzyme E1 regulatory subunit [D
>gi|1036759048|ref|XP_017156814.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1036813031|ref|XP_017137198.1| PREDICTED: uncharacterized protein LOC108152406 [Drosophila
>gi|1036813052|ref|XP_017137199.1| PREDICTED: uncharacterized protein LOC108152406 [Drosophila
>gi|1036813072|ref|XP_017137200.1| PREDICTED: uncharacterized protein LOC108152406 [Drosophila
>gi|1036812791|ref|XP_017137187.1| PREDICTED: adenosine deaminase CECR1 [Drosophila miranda]
>gi|1036812810|ref|XP_017137188.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812829|ref|XP_017137189.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812848|ref|XP_017137190.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812867|ref|XP_017137191.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812884|ref|XP_017137193.1| PREDICTED: adenosine deaminase CECR1 [Drosophila miranda]
>gi|1036812901|ref|XP_017137194.1| PREDICTED: basic-leucine zipper transcription factor A [Dros
>gi|1036812918|ref|XP_017137195.1| PREDICTED: basic-leucine zipper transcription factor A [Dros
>gi|1036813012|ref|XP_017137197.1| PREDICTED: spermine oxidase-like [Drosophila miranda]
>gi|1036812937|ref|XP_017137196.1| PREDICTED: peroxisomal N(1)-acetyl-spermine/spermidine oxida
>gi|1036850379|ref|XP_017139285.1| PREDICTED: uncharacterized protein LOC108153680 [Drosophila
>gi|1036823821|ref|XP_017137799.1| PREDICTED: nuclear transcription factor Y subunit beta [Dros
>gi|1036847217|ref|XP_017139102.1| PREDICTED: 15 kDa selenoprotein [Drosophila miranda]
>gi|1036783551|ref|XP_017135555.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036783570|ref|XP_017135556.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036783591|ref|XP_017135557.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036783513|ref|XP_017135553.1| PREDICTED: NEDD4 family-interacting protein 1 [Drosophila mi
>gi|1036783532|ref|XP_017135554.1| PREDICTED: NEDD4 family-interacting protein 1 [Drosophila mi
>gi|1036783435|ref|XP_017135548.1| PREDICTED: zinc finger protein on ecdysone puffs isoform X1
>gi|1036783454|ref|XP_017135549.1| PREDICTED: zinc finger protein on ecdysone puffs isoform X1
>gi|1036783473|ref|XP_017135551.1| PREDICTED: zinc finger protein on ecdysone puffs isoform X2
>gi|1036783494|ref|XP_017135552.1| PREDICTED: uncharacterized protein LOC108151448 [Drosophila
>gi|1036849083|ref|XP_017139208.1| PREDICTED: prostaglandin D2 receptor [Drosophila miranda]
>gi|1036820663|ref|XP_017137622.1| PREDICTED: integral membrane protein GPR155 [Drosophila mira
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>gi|1036766046|ref|XP_017134554.1| PREDICTED: RNA-binding motif protein, X-linked 2 isoform X1
>gi|1036766064|ref|XP_017134556.1| PREDICTED: RNA-binding motif protein, X-linked 2 isoform X2
>gi|1036841110|ref|XP_017138781.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841091|ref|XP_017138780.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840981|ref|XP_017138773.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841055|ref|XP_017138778.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840943|ref|XP_017138771.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841072|ref|XP_017138779.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841036|ref|XP_017138777.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840909|ref|XP_017138768.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841017|ref|XP_017138775.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840926|ref|XP_017138769.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840962|ref|XP_017138772.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840998|ref|XP_017138774.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840888|ref|XP_017138767.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841128|ref|XP_017138782.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841204|ref|XP_017138786.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841185|ref|XP_017138785.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841166|ref|XP_017138784.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841147|ref|XP_017138783.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036813396|ref|XP_017137219.1| PREDICTED: uncharacterized protein LOC108152417 [Drosophila
>gi|1036813356|ref|XP_017137217.1| PREDICTED: uncharacterized protein LOC108152415 isoform X1
>gi|1036813375|ref|XP_017137218.1| PREDICTED: uncharacterized protein LOC108152415 isoform X2
>gi|1036813314|ref|XP_017137215.1| PREDICTED: folliculin [Drosophila miranda]
>gi|1036813430|ref|XP_017137221.1| PREDICTED: 60S ribosomal protein L18 [Drosophila miranda]
>gi|1036813413|ref|XP_017137220.1| PREDICTED: 39S ribosomal protein L50, mitochondrial [Drosopl
>gi|1036813337|ref|XP_017137216.1| PREDICTED: nuclear receptor coactivator 5 [Drosophila miran
>gi|1036770311|ref|XP_017134787.1| PREDICTED: uncharacterized protein LOC108150964 [Drosophila
>gi|1036770197|ref|XP_017134780.1| PREDICTED: cell division cycle protein 27 homolog [Drosophil
>gi|1036770273|ref|XP_017134785.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036770159|ref|XP_017134778.1| PREDICTED: protein KIAA0100 [Drosophila miranda]
>gi|1036770178|ref|XP_017134779.1| PREDICTED: trafficking protein particle complex subunit 11
>gi|1036770237|ref|XP_017134782.1| PREDICTED: trypsin alpha [Drosophila miranda]
>gi|1036770216|ref|XP_017134781.1| PREDICTED: trypsin beta-like [Drosophila miranda]
>gi|1036758083|ref|XP_017156756.1| PREDICTED: transmembrane protease serine 9 [Drosophila mira
>gi|1036770255|ref|XP_017134783.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1036770292|ref|XP_017134786.1| PREDICTED: stress-associated endoplasmic reticulum protein 2
>gi|1036770349|ref|XP_017134790.1| PREDICTED: UPF0235 protein C15orf40 homolog [Drosophila mir
>gi|1036770330|ref|XP_017134788.1| PREDICTED: heat shock protein 83 [Drosophila miranda]
>gi|1036794352|ref|XP_017136167.1| PREDICTED: uncharacterized protein LOC108151819 [Drosophila
>gi|1036759067|ref|XP_017156815.1| PREDICTED: uncharacterized protein LOC108165292 [Drosophila
>gi|1036794387|ref|XP_017136169.1| PREDICTED: cell wall protein IFF6 [Drosophila miranda]
>gi|1036794371|ref|XP_017136168.1| PREDICTED: uncharacterized protein LOC108151820 [Drosophila
>gi|1036794317|ref|XP_017136165.1| PREDICTED: daple-like protein isoform X2 [Drosophila miranda
>gi|1036794302|ref|XP_017136164.1| PREDICTED: girdin isoform X1 [Drosophila miranda]
>gi|1036794333|ref|XP_017136166.1| PREDICTED: girdin isoform X3 [Drosophila miranda]
>gi|1036829922|ref|XP_017138144.1| PREDICTED: titin homolog [Drosophila miranda]
>gi|1036829960|ref|XP_017138146.1| PREDICTED: uncharacterized protein LOC108152957 [Drosophila
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>gi|1036829978|ref|XP_017138147.1| PREDICTED: uncharacterized protein LOC108152958 [Drosophila
>gi|1036829941|ref|XP_017138145.1| PREDICTED: tRNA (guanine(37)-N1)-methyltransferase [Drosoph
>gi|1036843652|ref|XP_017138917.1| PREDICTED: short/branched chain specific acyl-CoA dehydroger
>gi|1036843670|ref|XP_017138918.1| PREDICTED: ninjurin-1 [Drosophila miranda]
>gi|1036782545|ref|XP_017135493.1| PREDICTED: long-chain-fatty-acid--CoA ligase 5 isoform X1 [I
>gi|1036782564|ref|XP_017135494.1| PREDICTED: long-chain-fatty-acid--CoA ligase 5 isoform X1 [I
>gi|1036782582|ref|XP_017135495.1| PREDICTED: long-chain-fatty-acid--CoA ligase 5 isoform X2 [I
>gi|1036782526|ref|XP_017135492.1| PREDICTED: UDP-glucose:glycoprotein glucosyltransferase [Dro
>gi|1036782600|ref|XP_017135497.1| PREDICTED: cell cycle checkpoint control protein RAD9A [Dros
>gi|1036782657|ref|XP_017135500.1| PREDICTED: biogenesis of lysosome-related organelles complex
>gi|1036782674|ref|XP_017135501.1| PREDICTED: glutaredoxin-C4 [Drosophila miranda]
>gi|1036782638|ref|XP_017135499.1| PREDICTED: dysbindin protein homolog [Drosophila miranda]
>gi|1036782619|ref|XP_017135498.1| PREDICTED: ADP-ribosylation factor-like protein 6-interactin
>gi|1036759086|ref|XP_017156816.1| PREDICTED: transmembrane protease serine 9 [Drosophila miran
>gi|1036820924|ref|XP_017137637.1| PREDICTED: protein I'm not dead yet isoform X1 [Drosophila n
>gi|1036820943|ref|XP_017137638.1| PREDICTED: protein I'm not dead yet isoform X2 [Drosophila n
>gi|1036847497|ref|XP_017139119.1| PREDICTED: uncharacterized protein LOC108153557 isoform X1
>gi|1036847514|ref|XP_017139120.1| PREDICTED: uncharacterized protein LOC108153557 isoform X2
>gi|1036832533|ref|XP_017138292.1| PREDICTED: catalase isoform X1 [Drosophila miranda]
>gi|1036832552|ref|XP_017138293.1| PREDICTED: catalase isoform X2 [Drosophila miranda]
>gi|1036844356|ref|XP_017138953.1| PREDICTED: uncharacterized protein LOC108153447 [Drosophila
>gi|1036844374|ref|XP_017138954.1| PREDICTED: uncharacterized protein LOC108153447 [Drosophila
>gi|1036808939|ref|XP_017136984.1| PREDICTED: uncharacterized protein LOC108152274 [Drosophila
>gi|1036808919|ref|XP_017136983.1| PREDICTED: uncharacterized protein LOC108152273 [Drosophila
>gi|1036826370|ref|XP_017137940.1| PREDICTED: nuclear hormone receptor FTZ-F1 [Drosophila miran
>gi|1036820721|ref|XP_017137625.1| PREDICTED: ubiquitin-like-conjugating enzyme ATG3 [Drosophil
>gi|1036820700|ref|XP_017137624.1| PREDICTED: nuclear fragile X mental retardation-interacting
>gi|1036820741|ref|XP_017137626.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036763115|ref|XP_017157050.1| PREDICTED: protein terminus [Drosophila miranda]
>gi|1036764894|ref|XP_017157142.1| PREDICTED: M protein, serotype 2.1-like [Drosophila miranda]
>gi|1036847760|ref|XP_017139132.1| PREDICTED: uncharacterized protein LOC108153567 [Drosophila
>gi|1036849122|ref|XP_017139210.1| PREDICTED: uncharacterized protein LOC108153620 isoform X1
>gi|1036849139|ref|XP_017139211.1| PREDICTED: uncharacterized protein LOC108153620 isoform X1
>gi|1036849155|ref|XP_017139212.1| PREDICTED: uncharacterized protein LOC108153620 isoform X2
>gi|1036851689|ref|XP_017139363.1| PREDICTED: uncharacterized protein LOC108153745 [Drosophila
>gi|1036847869|ref|XP_017139138.1| PREDICTED: uncharacterized protein LOC108153571 [Drosophila
>gi|1036759105|ref|XP_017156817.1| PREDICTED: putative odorant receptor 69a [Drosophila miranda
>gi|1036759126|ref|XP_017156819.1| PREDICTED: putative odorant receptor 69a [Drosophila miranda
>gi|1036765869|ref|XP_017134544.1| PREDICTED: malate dehydrogenase, mitochondrial [Drosophila n
>gi|1036764947|ref|XP_017157143.1| PREDICTED: malate dehydrogenase, mitochondrial [Drosophila n
>gi|1036841227|ref|XP_017138787.1| PREDICTED: cullin homolog 1 [Drosophila miranda]
>gi|1036818364|ref|XP_017137502.1| PREDICTED: 10 kDa heat shock protein, mitochondrial [Drosopl
>gi|1036818346|ref|XP_017137501.1| PREDICTED: 39S ribosomal protein L20, mitochondrial [Drosopl
>gi|1036818328|ref|XP_017137500.1| PREDICTED: protein-S-isoprenylcysteine O-methyltransferase
>gi|1036818309|ref|XP_017137499.1| PREDICTED: adenosine kinase isoform X2 [Drosophila miranda]
>gi|1036818290|ref|XP_017137498.1| PREDICTED: adenosine kinase isoform X1 [Drosophila miranda]
>gi|1036829655|ref|XP_017138128.1| PREDICTED: 40S ribosomal protein S12 [Drosophila miranda]
>gi|1036829594|ref|XP_017138125.1| PREDICTED: zinc finger MYND domain-containing protein 10 hor
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>gi|1036829636|ref|XP_017138127.1| PREDICTED: matrix-remodeling-associated protein 7 [Drosophil
>gi|1036829615|ref|XP_017138126.1| PREDICTED: uncharacterized protein LOC108152942 [Drosophila
>gi|1036843889|ref|XP_017138928.1| PREDICTED: cyclin-H [Drosophila miranda]
>gi|1036764131|ref|XP_017157102.1| PREDICTED: protein embryonic gonad [Drosophila miranda]
>gi|1036825202|ref|XP_017137876.1| PREDICTED: ras-interacting protein RIP3-like [Drosophila mi
>gi|1036825183|ref|XP_017137875.1| PREDICTED: ras-interacting protein RIP3-like [Drosophila mi
>gi|1036759145|ref|XP_017156820.1| PREDICTED: catalase C-like [Drosophila miranda]
>gi|1036844555|ref|XP_017138965.1| PREDICTED: beta-1-syntrophin isoform X1 [Drosophila miranda]
>gi|1036844573|ref|XP_017138966.1| PREDICTED: beta-1-syntrophin isoform X1 [Drosophila miranda]
>gi|1036844727|ref|XP_017138967.1| PREDICTED: beta-1-syntrophin isoform X1 [Drosophila miranda]
>gi|1036844746|ref|XP_017138969.1| PREDICTED: beta-1-syntrophin isoform X2 [Drosophila miranda]
>gi|1036844765|ref|XP_017138970.1| PREDICTED: brain acid soluble protein 1 [Drosophila miranda]
>gi|1036770366|ref|XP_017134791.1| PREDICTED: uncharacterized protein LOC108150967 [Drosophila
>gi|1036767446|ref|XP_017134630.1| PREDICTED: uncharacterized protein LOC108150834 [Drosophila
>gi|1036849381|ref|XP_017139225.1| PREDICTED: uncharacterized protein LOC108153633 [Drosophila
>gi|1036849365|ref|XP_017139224.1| PREDICTED: translation initiation factor IF-2 [Drosophila mi
>gi|1036767741|ref|XP_017134643.1| PREDICTED: uncharacterized protein LOC108150851 [Drosophila
>gi|1036768687|ref|XP_017134695.1| PREDICTED: mucin-7 [Drosophila miranda]
>gi|1036823459|ref|XP_017137780.1| PREDICTED: uncharacterized protein LOC108152743 [Drosophila
>gi|1036823436|ref|XP_017137779.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036823478|ref|XP_017137781.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu
>gi|1036823417|ref|XP_017137777.1| PREDICTED: zinc finger protein OZF [Drosophila miranda]
>gi|1036836973|ref|XP_017138540.1| PREDICTED: uncharacterized protein LOC108153223 isoform X1
>gi|1036836992|ref|XP_017138542.1| PREDICTED: uncharacterized protein LOC108153223 isoform X2
>gi|1036832570|ref|XP_017138294.1| PREDICTED: ninein [Drosophila miranda]
>gi|1036759164|ref|XP_017156821.1| PREDICTED: acanthoscurrin-2 [Drosophila miranda]
>gi|1036759183|ref|XP_017156822.1| PREDICTED: translation initiation factor IF-2-like, partial
>gi|1036852527|ref|XP_017139416.1| PREDICTED: uncharacterized protein LOC108153783 [Drosophila
>gi|1036828033|ref|XP_017138037.1| PREDICTED: alsin homolog [Drosophila miranda]
>gi|1036828073|ref|XP_017138039.1| PREDICTED: uncharacterized protein LOC108152904 isoform X2
>gi|1036828054|ref|XP_017138038.1| PREDICTED: uncharacterized protein LOC108152904 isoform X1
>gi|1036764702|ref|XP_017157131.1| PREDICTED: endocuticle structural glycoprotein ABD-5 [Drosop
>gi|1036841558|ref|XP_017138807.1| PREDICTED: ankyrin repeat and BTB/POZ domain-containing prot
>gi|1036841539|ref|XP_017138806.1| PREDICTED: ankyrin repeat and BTB/POZ domain-containing prot
>gi|1036847741|ref|XP_017139131.1| PREDICTED: copper transport protein ATOX1 [Drosophila mirand
>gi|1036851857|ref|XP_017139374.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036851675|ref|XP_017139362.1| PREDICTED: probable fumarate hydratase, mitochondrial [Dros
>gi|1036811343|ref|XP_017137112.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036811365|ref|XP_017137113.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036811282|ref|XP_017137109.1| PREDICTED: fibroleukin [Drosophila miranda]
>gi|1036811305|ref|XP_017137110.1| PREDICTED: fibroleukin [Drosophila miranda]
>gi|1036811384|ref|XP_017137114.1| PREDICTED: ORM1-like protein [Drosophila miranda]
>gi|1036811324|ref|XP_017137111.1| PREDICTED: asparagine--tRNA ligase, cytoplasmic-like [Drosop
>gi|1036811912|ref|XP_017137141.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036811931|ref|XP_017137142.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036847271|ref|XP_017139106.1| PREDICTED: uncharacterized protein LOC108153549 [Drosophila
>gi|1036847290|ref|XP_017139107.1| PREDICTED: neuropeptide Y receptor type 6 [Drosophila mirand
>gi|1036847886|ref|XP_017139139.1| PREDICTED: uncharacterized protein LOC108153572 [Drosophila
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>gi|1036847904|ref|XP_017139140.1| PREDICTED: uncharacterized protein LOC108153572 [Drosophila
>gi|1036847941|ref|XP_017139143.1| PREDICTED: leucine-rich repeat-containing protein 9 [Drosop
>gi|1036847922|ref|XP_017139142.1| PREDICTED: glutamate receptor 2 [Drosophila miranda]
>gi|1036767964|ref|XP_017134655.1| PREDICTED: uncharacterized protein LOC108150864 [Drosophila
>gi|1036851890|ref|XP_017139377.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036851906|ref|XP_017139378.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036851925|ref|XP_017139379.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036834015|ref|XP_017138369.1| PREDICTED: ADP-dependent glucokinase [Drosophila miranda]
>gi|1036823840|ref|XP_017137800.1| PREDICTED: heat shock 70 kDa protein 4 isoform X1 [Drosophil
>gi|1036823859|ref|XP_017137801.1| PREDICTED: heat shock 70 kDa protein 4 isoform X1 [Drosophil
>gi|1036823878|ref|XP_017137802.1| PREDICTED: heat shock 70 kDa protein 4 isoform X2 [Drosophil
>gi|1036818969|ref|XP_017137537.1| PREDICTED: neuralized-like protein 4 [Drosophila miranda]
>gi|1036838446|ref|XP_017138630.1| PREDICTED: RNA exonuclease 4 [Drosophila miranda]
>gi|1036838463|ref|XP_017138631.1| PREDICTED: M-phase-specific PLK1-interacting protein [Drosop
>gi|1036809572|ref|XP_017137015.1| PREDICTED: formin-like protein CG32138 isoform X5 [Drosophil
>gi|1036809502|ref|XP_017137011.1| PREDICTED: formin-like protein CG32138 isoform X1 [Drosophil
>gi|1036809589|ref|XP_017137016.1| PREDICTED: formin-like protein CG32138 isoform X6 [Drosophil
>gi|1036809521|ref|XP_017137012.1| PREDICTED: formin-like protein CG32138 isoform X2 [Drosophil
>gi|1036809538|ref|XP_017137013.1| PREDICTED: formin-like protein CG32138 isoform X3 [Drosophil
>gi|1036809555|ref|XP_017137014.1| PREDICTED: formin-like protein CG32138 isoform X4 [Drosophil
>gi|1036809464|ref|XP_017137009.1| PREDICTED: peroxisome biogenesis factor 1 [Drosophila mirand
>gi|1036809446|ref|XP_017137008.1| PREDICTED: fibroblast growth factor receptor homolog 2 [Dros
>gi|1036809483|ref|XP_017137010.1| PREDICTED: fat-body protein 1 [Drosophila miranda]
>gi|1036846586|ref|XP_017139066.1| PREDICTED: fat-body protein 1 [Drosophila miranda]
>gi|1036835572|ref|XP_017138458.1| PREDICTED: transcription factor Sox-14 [Drosophila miranda]
>gi|1036759200|ref|XP_017156823.1| PREDICTED: BTB/POZ domain-containing protein 2-like [Drosop
>gi|1036835554|ref|XP_017138457.1| PREDICTED: RNA-binding protein fusilli [Drosophila miranda]
>gi|1036841614|ref|XP_017138810.1| PREDICTED: SOX domain-containing protein dichaete [Drosophil
>gi|1036852250|ref|XP_017139400.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036837329|ref|XP_017138562.1| PREDICTED: succinyl-CoA ligase [ADP/GDP-forming] subunit alp
>gi|1036811950|ref|XP_017137143.1| PREDICTED: protein encore isoform X1 [Drosophila miranda]
>gi|1036811988|ref|XP_017137145.1| PREDICTED: protein encore isoform X3 [Drosophila miranda]
>gi|1036811969|ref|XP_017137144.1| PREDICTED: protein encore isoform X2 [Drosophila miranda]
>gi|1036812007|ref|XP_017137146.1| PREDICTED: protein encore isoform X4 [Drosophila miranda]
>gi|1036846199|ref|XP_017139043.1| PREDICTED: LIM/homeobox protein Awh isoform X1 [Drosophila m
>gi|1036846216|ref|XP_017139044.1| PREDICTED: LIM/homeobox protein Awh isoform X2 [Drosophila m
>gi|1036846235|ref|XP_017139045.1| PREDICTED: LIM/homeobox protein Awh isoform X3 [Drosophila m
>gi|1036846256|ref|XP_017139046.1| PREDICTED: LIM/homeobox protein Awh isoform X4 [Drosophila m
>gi|1036763260|ref|XP_017157056.1| PREDICTED: uncharacterized protein LOC108165511 [Drosophila
>gi|1036833374|ref|XP_017138342.1| PREDICTED: uncharacterized protein LOC108153072 [Drosophila
>gi|1036759219|ref|XP_017156824.1| PREDICTED: gustatory and odorant receptor 63a [Drosophila mi
>gi|1036833340|ref|XP_017138339.1| PREDICTED: hsp90 co-chaperone Cdc37 [Drosophila miranda]
>gi|1036833356|ref|XP_017138341.1| PREDICTED: dnaJ homolog subfamily B member 13 [Drosophila mi
>gi|1036819433|ref|XP_017137560.1| PREDICTED: patj homolog isoform X1 [Drosophila miranda]
>gi|1036819473|ref|XP_017137563.1| PREDICTED: patj homolog isoform X3 [Drosophila miranda]
>gi|1036819454|ref|XP_017137561.1| PREDICTED: patj homolog isoform X2 [Drosophila miranda]
>gi|1036819492|ref|XP_017137564.1| PREDICTED: protein JTB [Drosophila miranda]
>gi|1036832072|ref|XP_017138268.1| PREDICTED: bromodomain-containing protein DDB_G0280777 isofo
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>gi|1036832089|ref|XP_017138269.1| PREDICTED: basic-leucine zipper transcription factor A isof
>gi|1036848164|ref|XP_017139154.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036769686|ref|XP_017134754.1| PREDICTED: uncharacterized protein LOC108150935 [Drosophila
>gi|1036767187|ref|XP_017134617.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1036851618|ref|XP_017139359.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1036850597|ref|XP_017139298.1| PREDICTED: uncharacterized protein LOC108153687 isoform X1
>gi|1036850616|ref|XP_017139299.1| PREDICTED: uncharacterized protein LOC108153687 isoform X2
>gi|1036805293|ref|XP_017136784.1| PREDICTED: CUE domain-containing protein 1 isoform X1 [Dros
>gi|1036805312|ref|XP_017136785.1| PREDICTED: CUE domain-containing protein 1 isoform X2 [Dros
>gi|1036805331|ref|XP_017136786.1| PREDICTED: CUE domain-containing protein 1 isoform X3 [Dros
>gi|1036805348|ref|XP_017136787.1| PREDICTED: ras-like protein 3 [Drosophila miranda]
>gi|1036805274|ref|XP_017136783.1| PREDICTED: HBS1-like protein [Drosophila miranda]
>gi|1036834916|ref|XP_017138419.1| PREDICTED: transmembrane protein 134 [Drosophila miranda]
>gi|1036823038|ref|XP_017137755.1| PREDICTED: uncharacterized protein LOC108152728 [Drosophila
>gi|1036823019|ref|XP_017137754.1| PREDICTED: FACT complex subunit spt16 isoform X2 [Drosophila
>gi|1036823000|ref|XP_017137752.1| PREDICTED: FACT complex subunit spt16 isoform X1 [Drosophila
>gi|1036840088|ref|XP_017138722.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840107|ref|XP_017138723.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840126|ref|XP_017138724.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840145|ref|XP_017138725.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840164|ref|XP_017138726.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X2 [Dros
>gi|1036840183|ref|XP_017138727.1| PREDICTED: adenine phosphoribosyltransferase [Drosophila mi
>gi|1036840204|ref|XP_017138728.1| PREDICTED: adenine phosphoribosyltransferase [Drosophila mi
>gi|1036836729|ref|XP_017138526.1| PREDICTED: lipoma HMGIC fusion partner-like 3 protein [Dros
>gi|1036846935|ref|XP_017139086.1| PREDICTED: lysosome membrane protein 2 isoform X1 [Drosophil
>gi|1036846954|ref|XP_017139087.1| PREDICTED: lysosome membrane protein 2 isoform X1 [Drosophil
>gi|1036846973|ref|XP_017139088.1| PREDICTED: lysosome membrane protein 2 isoform X2 [Drosophil
>gi|1036851944|ref|XP_017139380.1| PREDICTED: alpha-protein kinase 1 [Drosophila miranda]
>gi|1036848126|ref|XP_017139152.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dr
>gi|1036771948|ref|XP_017134879.1| PREDICTED: microfibrillar-associated protein 1 [Drosophila
>gi|1036759238|ref|XP_017156825.1| PREDICTED: uncharacterized protein LOC108165301 [Drosophila
>gi|1036771966|ref|XP_017134881.1| PREDICTED: microtubule-associated protein futsch-like [Dros
>gi|1036771929|ref|XP_017134878.1| PREDICTED: ankyrin repeat domain-containing protein 26 [Dro
>gi|1036771910|ref|XP_017134877.1| PREDICTED: uncharacterized protein LOC108151029 [Drosophila
>gi|1036788535|ref|XP_017135837.1| PREDICTED: RING finger protein 10 [Drosophila miranda]
>gi|1036788606|ref|XP_017135841.1| PREDICTED: choline-phosphate cytidylyltransferase A [Drosop
>gi|1036788570|ref|XP_017135839.1| PREDICTED: choline-phosphate cytidylyltransferase A [Drosop
>gi|1036788588|ref|XP_017135840.1| PREDICTED: choline-phosphate cytidylyltransferase A [Drosop
>gi|1036788519|ref|XP_017135836.1| PREDICTED: DNA polymerase eta [Drosophila miranda]
>gi|1036788550|ref|XP_017135838.1| PREDICTED: solute carrier family 22 member 3 [Drosophila mi
>gi|1036807066|ref|XP_017136880.1| PREDICTED: poly(rC)-binding protein 3 isoform X16 [Drosophil
>gi|1036807048|ref|XP_017136879.1| PREDICTED: poly(rC)-binding protein 3 isoform X15 [Drosophil
>gi|1036807030|ref|XP_017136878.1| PREDICTED: poly(rC)-binding protein 3 isoform X14 [Drosophil
>gi|1036807016|ref|XP_017136876.1| PREDICTED: poly(rC)-binding protein 3 isoform X13 [Drosophil
>gi|1036806997|ref|XP_017136875.1| PREDICTED: poly(rC)-binding protein 3 isoform X12 [Drosophil
>gi|1036806981|ref|XP_017136874.1| PREDICTED: poly(rC)-binding protein 3 isoform X11 [Drosophil
>gi|1036806964|ref|XP_017136873.1| PREDICTED: poly(rC)-binding protein 3 isoform X10 [Drosophil
>gi|1036806927|ref|XP_017136871.1| PREDICTED: poly(rC)-binding protein 3 isoform X8 [Drosophila
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>gi|1036806816|ref|XP_017136865.1| PREDICTED: poly(rC)-binding protein 3 isoform X2 [Drosophila
>gi|1036806946|ref|XP_017136872.1| PREDICTED: poly(rC)-binding protein 3 isoform X9 [Drosophila
>gi|1036806909|ref|XP_017136870.1| PREDICTED: poly(rC)-binding protein 3 isoform X7 [Drosophila
>gi|1036806890|ref|XP_017136869.1| PREDICTED: poly(rC)-binding protein 3 isoform X6 [Drosophila
>gi|1036806872|ref|XP_017136868.1| PREDICTED: poly(rC)-binding protein 3 isoform X5 [Drosophila
>gi|1036806853|ref|XP_017136867.1| PREDICTED: poly(rC)-binding protein 3 isoform X4 [Drosophila
>gi|1036806834|ref|XP_017136866.1| PREDICTED: poly(rC)-binding protein 3 isoform X3 [Drosophila
>gi|1036806742|ref|XP_017136861.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036806762|ref|XP_017136862.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036806781|ref|XP_017136863.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036806799|ref|XP_017136864.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036768667|ref|XP_017134694.1| PREDICTED: neuropeptide-like protein 29 [Drosophila miranda]
>gi|1036832974|ref|XP_017138318.1| PREDICTED: delta-1-pyrroline-5-carboxylate dehydrogenase, mi
>gi|1036825981|ref|XP_017137917.1| PREDICTED: nucleolar and coiled-body phosphoprotein 1-like
>gi|1036825939|ref|XP_017137915.1| PREDICTED: nucleolin 2-like isoform X1 [Drosophila miranda]
>gi|1036825958|ref|XP_017137916.1| PREDICTED: nucleolin 1-like isoform X2 [Drosophila miranda]
>gi|1036838182|ref|XP_017138614.1| PREDICTED: uncharacterized protein LOC108153263 [Drosophila
>gi|1036826144|ref|XP_017137927.1| PREDICTED: small integral membrane protein 8 [Drosophila mi
>gi|1036826127|ref|XP_017137926.1| PREDICTED: eukaryotic translation initiation factor 2A [Dros
>gi|1036846605|ref|XP_017139067.1| PREDICTED: uncharacterized protein LOC108153520 [Drosophila
>gi|1036799073|ref|XP_017136433.1| PREDICTED: serine/arginine repetitive matrix protein 1 [Dros
>gi|1036799107|ref|XP_017136435.1| PREDICTED: 40S ribosomal protein S4 [Drosophila miranda]
>gi|1036799090|ref|XP_017136434.1| PREDICTED: syntaxin-12 [Drosophila miranda]
>gi|1036799124|ref|XP_017136436.1| PREDICTED: uncharacterized protein LOC108151983 [Drosophila
>gi|1036768442|ref|XP_017134682.1| PREDICTED: uncharacterized protein LOC108150884 [Drosophila
>gi|1036767923|ref|XP_017134653.1| PREDICTED: uncharacterized protein LOC108150862 [Drosophila
>gi|1036759257|ref|XP_017156826.1| PREDICTED: uncharacterized protein LOC108165303 [Drosophila
>gi|1036848278|ref|XP_017139161.1| PREDICTED: serine/threonine-protein phosphatase PP1-like [Dr
>gi|1036835868|ref|XP_017138475.1| PREDICTED: leucine-rich repeat-containing protein 15 [Drosop
>gi|1036852125|ref|XP_017139392.1| PREDICTED: DC-STAMP domain-containing protein 1 [Drosophila
>gi|1036814002|ref|XP_017137255.1| PREDICTED: uncharacterized protein LOC108152434 [Drosophila
>gi|1036813984|ref|XP_017137254.1| PREDICTED: chondroadherin-like protein [Drosophila miranda]
>gi|1036759273|ref|XP_017156827.1| PREDICTED: hydrogenase nickel incorporation protein HypB-li
>gi|1036766387|ref|XP_017134570.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036766138|ref|XP_017134560.1| PREDICTED: collagen alpha-1(VIII) chain-like [Drosophila mi
>gi|1036850436|ref|XP_017139289.1| PREDICTED: cysteine and histidine-rich protein 1-B-like [Dro
>gi|1036850453|ref|XP_017139290.1| PREDICTED: neuropeptide-like 2 [Drosophila miranda]
>gi|1036847960|ref|XP_017139144.1| PREDICTED: uncharacterized protein LOC108153576 [Drosophila
>gi|1036846074|ref|XP_017139037.1| PREDICTED: uncharacterized protein LOC108153497 [Drosophila
>gi|1036759293|ref|XP_017156828.1| PREDICTED: uncharacterized protein LOC108165305, partial [D
>gi|1036846055|ref|XP_017139036.1| PREDICTED: nucleolin [Drosophila miranda]
>gi|1036852417|ref|XP_017139410.1| PREDICTED: uncharacterized protein LOC108153779 [Drosophila
>gi|1036821869|ref|XP_017137689.1| PREDICTED: uncharacterized protein LOC108152694 [Drosophila
>gi|1036821890|ref|XP_017137690.1| PREDICTED: transmembrane protein 19 isoform X1 [Drosophila m
>gi|1036821911|ref|XP_017137691.1| PREDICTED: transmembrane protein 19 isoform X2 [Drosophila m
>gi|1036821850|ref|XP_017137688.1| PREDICTED: POC1 centriolar protein homolog isoform X2 [Droso
>gi|1036821813|ref|XP_017137685.1| PREDICTED: POC1 centriolar protein homolog isoform X1 [Droso
>gi|1036821831|ref|XP_017137686.1| PREDICTED: POC1 centriolar protein homolog isoform X1 [Droso
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>gi|1036846624|ref|XP_017139068.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036849657|ref|XP_017139242.1| PREDICTED: tetratricopeptide repeat protein 36 homolog [Dros
>gi|1036849617|ref|XP_017139240.1| PREDICTED: uncharacterized protein LOC108153647 isoform X1
>gi|1036849636|ref|XP_017139241.1| PREDICTED: uncharacterized protein LOC108153647 isoform X2
>gi|1036842712|ref|XP_017138861.1| PREDICTED: whirlin isoform X3 [Drosophila miranda]
>gi|1036842833|ref|XP_017138868.1| PREDICTED: whirlin isoform X10 [Drosophila miranda]
>gi|1036842780|ref|XP_017138865.1| PREDICTED: whirlin isoform X7 [Drosophila miranda]
>gi|1036842728|ref|XP_017138862.1| PREDICTED: whirlin isoform X4 [Drosophila miranda]
>gi|1036842815|ref|XP_017138867.1| PREDICTED: whirlin isoform X9 [Drosophila miranda]
>gi|1036842693|ref|XP_017138859.1| PREDICTED: whirlin isoform X2 [Drosophila miranda]
>gi|1036842797|ref|XP_017138866.1| PREDICTED: whirlin isoform X8 [Drosophila miranda]
>gi|1036842747|ref|XP_017138863.1| PREDICTED: whirlin isoform X5 [Drosophila miranda]
>gi|1036842764|ref|XP_017138864.1| PREDICTED: whirlin isoform X6 [Drosophila miranda]
>gi|1036842677|ref|XP_017138858.1| PREDICTED: whirlin isoform X1 [Drosophila miranda]
>gi|1036842851|ref|XP_017138869.1| PREDICTED: whirlin isoform X11 [Drosophila miranda]
>gi|1036842864|ref|XP_017138870.1| PREDICTED: uncharacterized protein LOC108153394 [Drosophila
>gi|1036777927|ref|XP_017135223.1| PREDICTED: ral guanine nucleotide dissociation stimulator-l
>gi|1036777921|ref|XP_017135222.1| PREDICTED: G patch domain-containing protein 1 homolog [Dros
>gi|1036777911|ref|XP_017135220.1| PREDICTED: dynactin subunit 1 [Drosophila miranda]
>gi|1036777915|ref|XP_017135221.1| PREDICTED: dynactin subunit 1 [Drosophila miranda]
>gi|1036777939|ref|XP_017135224.1| PREDICTED: bicaudal D-related protein homolog isoform X1 [D
>gi|1036777964|ref|XP_017135225.1| PREDICTED: bicaudal D-related protein homolog isoform X1 [D
>gi|1036777976|ref|XP_017135226.1| PREDICTED: bicaudal D-related protein homolog isoform X2 [D
>gi|1036777990|ref|XP_017135227.1| PREDICTED: bicaudal D-related protein homolog isoform X2 [D
>gi|1036838482|ref|XP_017138632.1| PREDICTED: gastrula zinc finger protein XlCGF57.1 [Drosophil
>gi|1036785178|ref|XP_017135650.1| PREDICTED: ataxin-2 homolog [Drosophila miranda]
>gi|1036785322|ref|XP_017135656.1| PREDICTED: uncharacterized protein LOC108151519 [Drosophila
>gi|1036785197|ref|XP_017135651.1| PREDICTED: heat shock 70 kDa protein cognate 1 [Drosophila
>gi|1036785286|ref|XP_017135654.1| PREDICTED: uncharacterized protein LOC108151517 isoform X3
>gi|1036785269|ref|XP_017135653.1| PREDICTED: uncharacterized protein LOC108151517 isoform X2
>gi|1036785233|ref|XP_017135652.1| PREDICTED: uncharacterized protein LOC108151517 isoform X1
>gi|1036759312|ref|XP_017156830.1| PREDICTED: cathepsin L-like [Drosophila miranda]
>gi|1036785304|ref|XP_017135655.1| PREDICTED: pre-intermoult gene 1 protein-like [Drosophila m
>gi|1036759331|ref|XP_017156831.1| PREDICTED: histone deacetylase HDT3-like [Drosophila miranda
>gi|1036839962|ref|XP_017138714.1| PREDICTED: digestive cysteine proteinase 1 [Drosophila mira
>gi|1036813746|ref|XP_017137240.1| PREDICTED: formin-binding protein 1-like isoform X3 [Drosopl
>gi|1036813710|ref|XP_017137238.1| PREDICTED: formin-binding protein 1-like isoform X1 [Drosopl
>gi|1036813729|ref|XP_017137239.1| PREDICTED: formin-binding protein 1-like isoform X2 [Drosopl
>gi|1036813799|ref|XP_017137243.1| PREDICTED: formin-binding protein 1-like isoform X6 [Drosopl
>gi|1036813818|ref|XP_017137244.1| PREDICTED: formin-binding protein 1-like isoform X7 [Drosopl
>gi|1036813763|ref|XP_017137241.1| PREDICTED: formin-binding protein 1-like isoform X4 [Drosopl
>gi|1036813781|ref|XP_017137242.1| PREDICTED: formin-binding protein 1-like isoform X5 [Drosopl
>gi|1036813837|ref|XP_017137245.1| PREDICTED: formin-binding protein 1-like isoform X8 [Drosopl
>gi|1036813857|ref|XP_017137246.1| PREDICTED: formin-binding protein 1-like isoform X9 [Drosopl
>gi|1036786248|ref|XP_017135709.1| PREDICTED: uncharacterized protein LOC108151553 [Drosophila
>gi|1036786195|ref|XP_017135706.1| PREDICTED: uncharacterized protein LOC108151551 isoform X1
>gi|1036786212|ref|XP_017135707.1| PREDICTED: uncharacterized protein LOC108151551 isoform X2
>gi|1036786029|ref|XP_017135696.1| PREDICTED: semaphorin-5A isoform X2 [Drosophila miranda]
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>gi|1036785988|ref|XP_017135694.1| PREDICTED: semaphorin-5A isoform X1 [Drosophila miranda]
>gi|1036786008|ref|XP_017135695.1| PREDICTED: semaphorin-5A isoform X1 [Drosophila miranda]
>gi|1036786231|ref|XP_017135708.1| PREDICTED: uncharacterized protein LOC108151552 [Drosophila
>gi|1036786048|ref|XP_017135697.1| PREDICTED: uncharacterized protein LOC108151547 isoform X1
>gi|1036786065|ref|XP_017135698.1| PREDICTED: uncharacterized protein LOC108151547 isoform X2
>gi|1036786084|ref|XP_017135699.1| PREDICTED: uncharacterized protein LOC108151548 isoform X1
>gi|1036786103|ref|XP_017135700.1| PREDICTED: uncharacterized protein LOC108151548 isoform X2
>gi|1036786159|ref|XP_017135704.1| PREDICTED: uncharacterized protein LOC108151550 isoform X1
>gi|1036786177|ref|XP_017135705.1| PREDICTED: uncharacterized protein LOC108151550 isoform X2
>gi|1036786141|ref|XP_017135703.1| PREDICTED: nuclease-sensitive element-binding protein 1 iso:
>gi|1036786122|ref|XP_017135702.1| PREDICTED: nuclease-sensitive element-binding protein 1 iso:
>gi|1036759350|ref|XP_017156832.1| PREDICTED: uncharacterized protein LOC108165308 [Drosophila
>gi|1036790950|ref|XP_017135974.1| PREDICTED: sterol regulatory element-binding protein 1 [Dros
>gi|1036790969|ref|XP_017135975.1| PREDICTED: sterol regulatory element-binding protein 1 [Dros
>gi|1036791116|ref|XP_017135984.1| PREDICTED: uncharacterized protein LOC108151719 isoform X3
>gi|1036791040|ref|XP_017135979.1| PREDICTED: uncharacterized protein LOC108151718 isoform X1
>gi|1036791059|ref|XP_017135980.1| PREDICTED: uncharacterized protein LOC108151718 isoform X2
>gi|1036791097|ref|XP_017135983.1| PREDICTED: uncharacterized protein LOC108151719 isoform X2
>gi|1036791078|ref|XP_017135982.1| PREDICTED: uncharacterized protein LOC108151719 isoform X1
>gi|1036791135|ref|XP_017135985.1| PREDICTED: uncharacterized protein LOC108151720 [Drosophila
>gi|1036791154|ref|XP_017135986.1| PREDICTED: coiled-coil domain-containing protein 58 [Drosoph
>gi|1036791173|ref|XP_017135987.1| PREDICTED: DNA repair protein SWI5 homolog [Drosophila mirar
>gi|1036790984|ref|XP_017135976.1| PREDICTED: importin subunit alpha-7 [Drosophila miranda]
>gi|1036791003|ref|XP_017135977.1| PREDICTED: probable cytochrome P450 305a1 isoform X1 [Droso]
>gi|1036791021|ref|XP_017135978.1| PREDICTED: probable cytochrome P450 305a1 isoform X2 [Droso]
>gi|1036834413|ref|XP_017138389.1| PREDICTED: protein cycle isoform X2 [Drosophila miranda]
>gi|1036834394|ref|XP_017138387.1| PREDICTED: protein cycle isoform X1 [Drosophila miranda]
>gi|1036834432|ref|XP_017138390.1| PREDICTED: acidic fibroblast growth factor intracellular-bir
>gi|1036834470|ref|XP_017138392.1| PREDICTED: deformed epidermal autoregulatory factor 1 isofor
>gi|1036834489|ref|XP_017138393.1| PREDICTED: deformed epidermal autoregulatory factor 1 isofor
>gi|1036826297|ref|XP_017137936.1| PREDICTED: serine/threonine-protein kinase tricorner [Drosop
>gi|1036826315|ref|XP_017137937.1| PREDICTED: uncharacterized protein LOC108152840 [Drosophila
>gi|1036803956|ref|XP_017136707.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036803974|ref|XP_017136708.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036833484|ref|XP_017138347.1| PREDICTED: bifunctional 3'-phosphoadenosine 5'-phosphosulfat
>gi|1036833448|ref|XP_017138345.1| PREDICTED: bifunctional 3'-phosphoadenosine 5'-phosphosulfat
>gi|1036833467|ref|XP_017138346.1| PREDICTED: bifunctional 3'-phosphoadenosine 5'-phosphosulfat
>gi|1036831476|ref|XP_017138236.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1036831495|ref|XP_017138237.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1036831512|ref|XP_017138238.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1036838201|ref|XP_017138616.1| PREDICTED: uncharacterized protein LOC108153264 [Drosophila
>gi|1036803770|ref|XP_017136698.1| PREDICTED: immediate early response 3-interacting protein 1
>gi|1036803751|ref|XP_017136697.1| PREDICTED: biogenesis of lysosome-related organelles comple>
>gi|1036803716|ref|XP_017136695.1| PREDICTED: uncharacterized protein YER152C [Drosophila mirar
>gi|1036803608|ref|XP_017136689.1| PREDICTED: proton-coupled amino acid transporter 1 isoform >
>gi|1036803627|ref|XP_017136690.1| PREDICTED: proton-coupled amino acid transporter 1 isoform >
>gi|1036803645|ref|XP_017136691.1| PREDICTED: proton-coupled amino acid transporter 1 isoform >
>gi|1036803663|ref|XP_017136692.1| PREDICTED: proton-coupled amino acid transporter 1 isoform >
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>gi|1036803571|ref|XP_017136687.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803590|ref|XP_017136688.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803555|ref|XP_017136686.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803536|ref|XP_017136685.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803517|ref|XP_017136684.1| PREDICTED: proton-coupled amino acid transporter 4 isoform
>gi|1036803698|ref|XP_017136694.1| PREDICTED: proton-coupled amino acid transporter 1 [Drosophi
>gi|1036803680|ref|XP_017136693.1| PREDICTED: proton-coupled amino acid transporter 3 [Drosophi
>gi|1036803732|ref|XP_017136696.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1036823614|ref|XP_017137789.1| PREDICTED: uncharacterized protein LOC108152746 isoform X4
>gi|1036823499|ref|XP_017137783.1| PREDICTED: uncharacterized protein LOC108152745 [Drosophila
>gi|1036823518|ref|XP_017137784.1| PREDICTED: histidine-rich glycoprotein isoform X1 [Drosophil
>gi|1036823537|ref|XP_017137785.1| PREDICTED: histidine-rich glycoprotein isoform X2 [Drosophil
>gi|1036823559|ref|XP_017137786.1| PREDICTED: histidine-rich glycoprotein isoform X3 [Drosophil
>gi|1036823577|ref|XP_017137787.1| PREDICTED: histidine-rich glycoprotein isoform X3 [Drosophil
>gi|1036823595|ref|XP_017137788.1| PREDICTED: histidine-rich glycoprotein isoform X3 [Drosophil
>gi|1036825772|ref|XP_017137905.1| PREDICTED: diphosphoinositol polyphosphate phosphohydrolase
>gi|1036825790|ref|XP_017137906.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036764021|ref|XP_017157100.1| PREDICTED: larval cuticle protein LCP-17-like [Drosophila mi
>gi|1036834545|ref|XP_017138397.1| PREDICTED: mitochondrial inner membrane protein OXA1L [Droso
>gi|1036834566|ref|XP_017138398.1| PREDICTED: MIP18 family protein CG7949 [Drosophila miranda]
>gi|1036844391|ref|XP_017138956.1| PREDICTED: histidine-rich protein PFHRP-II isoform X1 [Dros
>gi|1036844409|ref|XP_017138957.1| PREDICTED: sarcoplasmic reticulum histidine-rich calcium-bi
>gi|1036844427|ref|XP_017138958.1| PREDICTED: biogenesis of lysosome-related organelles comple
>gi|1036826109|ref|XP_017137925.1| PREDICTED: ATP-dependent RNA helicase DDX42 [Drosophila mira
>gi|1036804160|ref|XP_017136719.1| PREDICTED: zinc finger MIZ domain-containing protein 2 [Dros
>gi|1036759367|ref|XP_017156833.1| PREDICTED: keratin, type II cytoskeletal 2 epidermal [Drosop
>gi|1036783689|ref|XP_017135562.1| PREDICTED: transmembrane protein 184B isoform X1 [Drosophila
>gi|1036783707|ref|XP_017135564.1| PREDICTED: transmembrane protein 184B isoform X2 [Drosophila
>gi|1036783762|ref|XP_017135567.1| PREDICTED: 40S ribosomal protein S6-like [Drosophila miranda
>gi|1036783726|ref|XP_017135565.1| PREDICTED: rho GTPase-activating protein 17 [Drosophila mira
>gi|1036783610|ref|XP_017135558.1| PREDICTED: uncharacterized protein LOC108151451 [Drosophila
>gi|1036783629|ref|XP_017135559.1| PREDICTED: uncharacterized protein LOC108151451 [Drosophila
>gi|1036783745|ref|XP_017135566.1| PREDICTED: peroxisome biogenesis factor 10 [Drosophila mirar
>gi|1036783650|ref|XP_017135560.1| PREDICTED: molybdenum cofactor biosynthesis protein 1 isofor
>gi|1036783671|ref|XP_017135561.1| PREDICTED: molybdenum cofactor biosynthesis protein 1 isofor
>gi|1036783779|ref|XP_017135568.1| PREDICTED: probable prefoldin subunit 2 [Drosophila miranda]
>gi|1036827959|ref|XP_017138034.1| PREDICTED: unc-112-related protein [Drosophila miranda]
>gi|1036830330|ref|XP_017138169.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036830348|ref|XP_017138171.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036830367|ref|XP_017138172.1| PREDICTED: putative uncharacterized protein DDB_G0272516 iso
>gi|1036830385|ref|XP_017138173.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036830404|ref|XP_017138174.1| PREDICTED: adenylate cyclase, terminal-differentiation speci
>gi|1036830423|ref|XP_017138175.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X5
>gi|1036830440|ref|XP_017138176.1| PREDICTED: G-box-binding factor isoform X6 [Drosophila miran
>gi|1036830459|ref|XP_017138177.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X7
>gi|1036830478|ref|XP_017138178.1| PREDICTED: uncharacterized protein LOC108152969 isoform X8
>gi|1036830495|ref|XP_017138179.1| PREDICTED: uncharacterized protein LOC108152969 isoform X9
>gi|1036830514|ref|XP_017138180.1| PREDICTED: uncharacterized protein LOC108152969 isoform X10
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>gi|1036842584|ref|XP_017138853.1| PREDICTED: uncharacterized protein LOC108153389 [Drosophila
>gi|1036842530|ref|XP_017138851.1| PREDICTED: glutamate decarboxylase [Drosophila miranda]
>gi|1036842565|ref|XP_017138852.1| PREDICTED: glutamate decarboxylase [Drosophila miranda]
>gi|1036848824|ref|XP_017139192.1| PREDICTED: fumarylacetoacetase [Drosophila miranda]
>gi|1036830702|ref|XP_017138191.1| PREDICTED: bone morphogenetic protein receptor type-2 [Dros
>gi|1036830721|ref|XP_017138192.1| PREDICTED: bone morphogenetic protein receptor type-2 [Dros
>gi|1036830740|ref|XP_017138194.1| PREDICTED: cytochrome P450 302a1, mitochondrial [Drosophila
>gi|1036851838|ref|XP_017139373.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036851819|ref|XP_017139371.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036846735|ref|XP_017139075.1| PREDICTED: uncharacterized protein LOC108153526 isoform X2
>gi|1036846717|ref|XP_017139074.1| PREDICTED: uncharacterized protein LOC108153526 isoform X1
>gi|1036846754|ref|XP_017139076.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846771|ref|XP_017139077.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846790|ref|XP_017139078.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846808|ref|XP_017139079.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846827|ref|XP_017139080.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036832162|ref|XP_017138274.1| PREDICTED: microspherule protein 1 [Drosophila miranda]
>gi|1036834451|ref|XP_017138391.1| PREDICTED: BCL2/adenovirus E1B 19 kDa protein-interacting pr
>gi|1036794954|ref|XP_017136201.1| PREDICTED: gustatory receptor for sugar taste 64f isoform X1
>gi|1036794970|ref|XP_017136202.1| PREDICTED: gustatory receptor for sugar taste 64f isoform X2
>gi|1036795027|ref|XP_017136205.1| PREDICTED: gustatory receptor for sugar taste 64e isoform X1
>gi|1036795046|ref|XP_017136206.1| PREDICTED: gustatory receptor for sugar taste 64e isoform X2
>gi|1036795119|ref|XP_017136211.1| PREDICTED: gustatory receptor for sugar taste 64d isoform X4
>gi|1036795081|ref|XP_017136208.1| PREDICTED: gustatory receptor for sugar taste 64c isoform X2
>gi|1036795100|ref|XP_017136209.1| PREDICTED: gustatory receptor for sugar taste 64c isoform X3
>gi|1036795065|ref|XP_017136207.1| PREDICTED: gustatory receptor for sugar taste 64c isoform X1
>gi|1036795213|ref|XP_017136216.1| PREDICTED: gustatory receptor for sugar taste 64c-like isofo
>gi|1036795194|ref|XP_017136215.1| PREDICTED: gustatory receptor for sugar taste 64c-like isofo
>gi|1036795230|ref|XP_017136217.1| PREDICTED: gustatory receptor for sugar taste 64b [Drosophil
>gi|1036795008|ref|XP_017136204.1| PREDICTED: gustatory receptor for sugar taste 64a isoform X2
>gi|1036794988|ref|XP_017136203.1| PREDICTED: gustatory receptor for sugar taste 64a isoform X1
>gi|1036795139|ref|XP_017136212.1| PREDICTED: gustatory receptor for sugar taste 61a isoform X1
>gi|1036795175|ref|XP_017136214.1| PREDICTED: gustatory receptor for sugar taste 61a isoform X3
>gi|1036795156|ref|XP_017136213.1| PREDICTED: gustatory receptor for sugar taste 61a isoform X2
>gi|1036794824|ref|XP_017136193.1| PREDICTED: centrosomal protein of 290 kDa isoform X1 [Drosop
>gi|1036794843|ref|XP_017136194.1| PREDICTED: centrosomal protein of 290 kDa isoform X1 [Drosop
>gi|1036794860|ref|XP_017136195.1| PREDICTED: centrosomal protein of 290 kDa isoform X2 [Drosop
>gi|1036795267|ref|XP_017136219.1| PREDICTED: enoyl-CoA delta isomerase 2, mitochondrial-like
>gi|1036795248|ref|XP_017136218.1| PREDICTED: enoyl-CoA delta isomerase 2, mitochondrial-like
>gi|1036794938|ref|XP_017136200.1| PREDICTED: uncharacterized protein LOC108151835 [Drosophila
>gi|1036794879|ref|XP_017136197.1| PREDICTED: uncharacterized protein LOC108151834 isoform X1
>gi|1036794898|ref|XP_017136198.1| PREDICTED: uncharacterized protein LOC108151834 isoform X1
>gi|1036794918|ref|XP_017136199.1| PREDICTED: uncharacterized protein LOC108151834 isoform X2
>gi|1036781482|ref|XP_017135435.1| PREDICTED: nuclear receptor coactivator 6 [Drosophila mirand
>gi|1036781444|ref|XP_017135433.1| PREDICTED: monocarboxylate transporter 1 [Drosophila miranda
>gi|1036781463|ref|XP_017135434.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036805988|ref|XP_017136824.1| PREDICTED: uncharacterized protein LOC108152187 [Drosophila
>gi|1036805970|ref|XP_017136823.1| PREDICTED: gastrula zinc finger protein XlCGF53.1 [Drosophil
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>gi|1036806098|ref|XP_017136825.1| PREDICTED: LOW QUALITY PROTEIN: glucose transporter type 1
>gi|1036846991|ref|XP_017139089.1| PREDICTED: protein FAM69C [Drosophila miranda]
>gi|1036838144|ref|XP_017138612.1| PREDICTED: uncharacterized protein LOC108153260 [Drosophila
>gi|1036838125|ref|XP_017138611.1| PREDICTED: myb-like protein AA [Drosophila miranda]
>gi|1036788823|ref|XP_017135855.1| PREDICTED: phosphatase and actin regulator 4 isoform X11 [Dr
>gi|1036788750|ref|XP_017135850.1| PREDICTED: phosphatase and actin regulator 4 isoform X7 [Dro
>gi|1036788716|ref|XP_017135848.1| PREDICTED: neurogenic protein mastermind isoform X5 [Drosopl
>gi|1036788697|ref|XP_017135847.1| PREDICTED: neurogenic protein mastermind isoform X4 [Drosopl
>gi|1036788678|ref|XP_017135846.1| PREDICTED: neurogenic protein mastermind isoform X3 [Drosopl
>gi|1036788659|ref|XP_017135845.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosopl
>gi|1036788624|ref|XP_017135842.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosopl
>gi|1036788641|ref|XP_017135844.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosopl
>gi|1036788767|ref|XP_017135851.1| PREDICTED: phosphatase and actin regulator 4 isoform X8 [Dro
>gi|1036788842|ref|XP_017135856.1| PREDICTED: phosphatase and actin regulator 2 isoform X12 [Di
>gi|1036788786|ref|XP_017135852.1| PREDICTED: phosphatase and actin regulator 2 isoform X9 [Dro
>gi|1036788804|ref|XP_017135853.1| PREDICTED: neurogenic protein mastermind isoform X10 [Drosop
>gi|1036788736|ref|XP_017135849.1| PREDICTED: neurogenic protein mastermind isoform X6 [Drosopl
>gi|1036788861|ref|XP_017135857.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036788912|ref|XP_017135859.1| PREDICTED: uncharacterized protein LOC108151651 [Drosophila
>gi|1036788879|ref|XP_017135858.1| PREDICTED: protein nutcracker [Drosophila miranda]
>gi|1036809606|ref|XP_017137017.1| PREDICTED: integrator complex subunit 10 [Drosophila miranda
>gi|1036809657|ref|XP_017137020.1| PREDICTED: mpv17-like protein 2 [Drosophila miranda]
>gi|1036809676|ref|XP_017137023.1| PREDICTED: mpv17-like protein 2 [Drosophila miranda]
>gi|1036809623|ref|XP_017137018.1| PREDICTED: GPI mannosyltransferase 3 isoform X1 [Drosophila
>gi|1036809640|ref|XP_017137019.1| PREDICTED: GPI mannosyltransferase 3 isoform X2 [Drosophila
>gi|1036806136|ref|XP_017136827.1| PREDICTED: rab11 family-interacting protein 4B isoform X2 [I
>gi|1036806169|ref|XP_017136830.1| PREDICTED: rab11 family-interacting protein 4B isoform X4 [I
>gi|1036806187|ref|XP_017136831.1| PREDICTED: rab11 family-interacting protein 4B isoform X5 [I
>gi|1036806205|ref|XP_017136832.1| PREDICTED: rab11 family-interacting protein 4B isoform X6 [I
>gi|1036806224|ref|XP_017136833.1| PREDICTED: rab11 family-interacting protein 4B isoform X7 [I
>gi|1036806279|ref|XP_017136836.1| PREDICTED: peptidyl-prolyl cis-trans isomerase [Drosophila r
>gi|1036806117|ref|XP_017136826.1| PREDICTED: bromodomain-containing protein DDB_G0280777 isofc
>gi|1036806151|ref|XP_017136829.1| PREDICTED: bromodomain-containing protein DDB_G0280777 isofc
>gi|1036806243|ref|XP_017136834.1| PREDICTED: rab11 family-interacting protein 4B isoform X8 [I
>gi|1036806262|ref|XP_017136835.1| PREDICTED: MAGE-like protein 2 [Drosophila miranda]
>gi|1036806297|ref|XP_017136837.1| PREDICTED: uncharacterized protein LOC108152194 [Drosophila
>gi|1036764444|ref|XP_017157118.1| PREDICTED: alpha/beta-gliadin A-V [Drosophila miranda]
>gi|1036806412|ref|XP_017136844.1| PREDICTED: frizzled isoform X1 [Drosophila miranda]
>gi|1036806433|ref|XP_017136845.1| PREDICTED: frizzled isoform X2 [Drosophila miranda]
>gi|1036806452|ref|XP_017136846.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036851543|ref|XP_017139355.1| PREDICTED: uncharacterized protein LOC108153736 [Drosophila
>gi|1036839083|ref|XP_017138670.1| PREDICTED: uncharacterized protein LOC108153294 [Drosophila
>gi|1036759386|ref|XP_017156834.1| PREDICTED: digestive cysteine proteinase 1-like, partial [Di
>gi|1036787666|ref|XP_017135789.1| PREDICTED: uncharacterized protein LOC108151609 isoform X1
>gi|1036787681|ref|XP_017135790.1| PREDICTED: uncharacterized protein LOC108151609 isoform X2
>gi|1036787696|ref|XP_017135791.1| PREDICTED: uncharacterized protein LOC108151610 [Drosophila
>gi|1036787592|ref|XP_017135783.1| PREDICTED: uncharacterized protein LOC108151608 isoform X1
>gi|1036787607|ref|XP_017135785.1| PREDICTED: PDZ domain-containing protein 2 isoform X2 [Drosc
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>gi|1036787622|ref|XP_017135786.1| PREDICTED: PDZ domain-containing protein 2 isoform X3 [Dros
>gi|1036787637|ref|XP_017135787.1| PREDICTED: PDZ domain-containing protein 2 isoform X4 [Dros
>gi|1036787651|ref|XP_017135788.1| PREDICTED: PDZ domain-containing protein 2 isoform X5 [Dros
>gi|1036759403|ref|XP_017156835.1| PREDICTED: uncharacterized protein LOC108165311 [Drosophila
>gi|1036759420|ref|XP_017156836.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036834033|ref|XP_017138370.1| PREDICTED: phosphate carrier protein, mitochondrial [Drosopl
>gi|1036834049|ref|XP_017138371.1| PREDICTED: phosphate carrier protein, mitochondrial [Drosopl
>gi|1036824733|ref|XP_017137852.1| PREDICTED: protein phosphatase 1 regulatory subunit 3C isofo
>gi|1036824714|ref|XP_017137851.1| PREDICTED: protein phosphatase 1 regulatory subunit 3C isofo
>gi|1036836428|ref|XP_017138509.1| PREDICTED: COMM domain-containing protein 4 [Drosophila mira
>gi|1036836466|ref|XP_017138511.1| PREDICTED: leucokinin [Drosophila miranda]
>gi|1036836447|ref|XP_017138510.1| PREDICTED: tctex1 domain-containing protein 2 [Drosophila mi
>gi|1036848088|ref|XP_017139150.1| PREDICTED: homeobox protein Nkx-6.1 [Drosophila miranda]
>gi|1036841523|ref|XP_017138804.1| PREDICTED: ecdysone 20-monooxygenase isoform X3 [Drosophila
>gi|1036841469|ref|XP_017138801.1| PREDICTED: ecdysone 20-monooxygenase isoform X1 [Drosophila
>gi|1036841486|ref|XP_017138802.1| PREDICTED: ecdysone 20-monooxygenase isoform X1 [Drosophila
>gi|1036841505|ref|XP_017138803.1| PREDICTED: ecdysone 20-monooxygenase isoform X2 [Drosophila
>gi|1036817367|ref|XP_017137445.1| PREDICTED: uncharacterized protein LOC108152545 isoform X1
>gi|1036817386|ref|XP_017137446.1| PREDICTED: uncharacterized protein LOC108152545 isoform X2
>gi|1036817407|ref|XP_017137447.1| PREDICTED: uncharacterized protein LOC108152545 isoform X2
>gi|1036817348|ref|XP_017137444.1| PREDICTED: ATP-dependent DNA helicase Q5 [Drosophila miranda
>gi|1036801968|ref|XP_017136600.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036802022|ref|XP_017136603.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036801984|ref|XP_017136601.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036802004|ref|XP_017136602.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036802040|ref|XP_017136605.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036791429|ref|XP_017136001.1| PREDICTED: protein alan shepard isoform X9 [Drosophila miran
>gi|1036791502|ref|XP_017136006.1| PREDICTED: protein alan shepard isoform X13 [Drosophila mira
>gi|1036791447|ref|XP_017136002.1| PREDICTED: protein alan shepard isoform X10 [Drosophila mira
>gi|1036791393|ref|XP_017135999.1| PREDICTED: protein alan shepard isoform X7 [Drosophila miran
>gi|1036791301|ref|XP_017135994.1| PREDICTED: protein alan shepard isoform X2 [Drosophila miran
>gi|1036791412|ref|XP_017136000.1| PREDICTED: protein alan shepard isoform X8 [Drosophila miran
>gi|1036791483|ref|XP_017136005.1| PREDICTED: protein alan shepard isoform X12 [Drosophila mira
>gi|1036791375|ref|XP_017135998.1| PREDICTED: protein alan shepard isoform X6 [Drosophila miran
>gi|1036791356|ref|XP_017135997.1| PREDICTED: protein alan shepard isoform X5 [Drosophila miran
>gi|1036791320|ref|XP_017135995.1| PREDICTED: protein alan shepard isoform X3 [Drosophila miran
>gi|1036791338|ref|XP_017135996.1| PREDICTED: protein alan shepard isoform X4 [Drosophila miran
>gi|1036791266|ref|XP_017135993.1| PREDICTED: protein alan shepard isoform X1 [Drosophila miran
>gi|1036791465|ref|XP_017136004.1| PREDICTED: protein alan shepard isoform X11 [Drosophila mira
>gi|1036850180|ref|XP_017139275.1| PREDICTED: uncharacterized protein LOC108153672 [Drosophila
>gi|1036852180|ref|XP_017139395.1| PREDICTED: glutamate dehydrogenase, mitochondrial-like [Dros
>gi|1036767244|ref|XP_017134620.1| PREDICTED: uncharacterized protein LOC108150824 [Drosophila
>gi|1036767263|ref|XP_017134621.1| PREDICTED: uncharacterized protein LOC108150824 [Drosophila
>gi|1036766739|ref|XP_017134591.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036821581|ref|XP_017137672.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036759439|ref|XP_017156837.1| PREDICTED: uncharacterized protein LOC108165314 [Drosophila
>gi|1036821463|ref|XP_017137666.1| PREDICTED: uncharacterized protein LOC108152680 isoform X1
>gi|1036821484|ref|XP_017137667.1| PREDICTED: uncharacterized protein LOC108152680 isoform X2
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>gi|1036821505|ref|XP_017137668.1| PREDICTED: uncharacterized protein LOC108152680 isoform X2
>gi|1036821525|ref|XP_017137669.1| PREDICTED: uncharacterized protein LOC108152681 isoform X1
>gi|1036821544|ref|XP_017137670.1| PREDICTED: uncharacterized protein LOC108152681 isoform X2
>gi|1036821444|ref|XP_017137665.1| PREDICTED: ATP-binding cassette sub-family A member 3 [Drosc
>gi|1036821562|ref|XP_017137671.1| PREDICTED: uncharacterized protein LOC108152682 [Drosophila
>gi|1036799479|ref|XP_017136456.1| PREDICTED: junctional adhesion molecule A isoform X2 [Drosop
>gi|1036799460|ref|XP_017136455.1| PREDICTED: fibroblast growth factor receptor-like 1 isoform
>gi|1036799536|ref|XP_017136460.1| PREDICTED: N-alpha-acetyltransferase daf-31 [Drosophila mira
>gi|1036799554|ref|XP_017136461.1| PREDICTED: small integral membrane protein 13 [Drosophila mi
>gi|1036799517|ref|XP_017136458.1| PREDICTED: protein TSSC4 [Drosophila miranda]
>gi|1036799498|ref|XP_017136457.1| PREDICTED: caspase Dronc [Drosophila miranda]
>gi|1036836151|ref|XP_017138493.1| PREDICTED: uncharacterized protein LOC108153189 isoform X1
>gi|1036836169|ref|XP_017138494.1| PREDICTED: uncharacterized protein LOC108153189 isoform X2
>gi|1036836186|ref|XP_017138495.1| PREDICTED: uncharacterized protein LOC108153189 isoform X3
>gi|1036836132|ref|XP_017138491.1| PREDICTED: SCY1-like protein 2 isoform X2 [Drosophila mirand
>gi|1036836113|ref|XP_017138490.1| PREDICTED: SCY1-like protein 2 isoform X1 [Drosophila mirand
>gi|1036794728|ref|XP_017136188.1| PREDICTED: nuclear distribution protein nudE homolog isoform
>gi|1036794710|ref|XP_017136187.1| PREDICTED: nuclear distribution protein nudE homolog isoform
>gi|1036794748|ref|XP_017136189.1| PREDICTED: type 1 phosphatidylinositol 4,5-bisphosphate 4-ph
>gi|1036794768|ref|XP_017136190.1| PREDICTED: type 1 phosphatidylinositol 4,5-bisphosphate 4-ph
>gi|1036794786|ref|XP_017136191.1| PREDICTED: type 1 phosphatidylinositol 4,5-bisphosphate 4-ph
>gi|1036794805|ref|XP_017136192.1| PREDICTED: uncharacterized protein LOC108151831 [Drosophila
>gi|1036794693|ref|XP_017136186.1| PREDICTED: calpain-B [Drosophila miranda]
>gi|1036814019|ref|XP_017137256.1| PREDICTED: transcription initiation factor TFIID subunit 2
>gi|1036814055|ref|XP_017137258.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1036814037|ref|XP_017137257.1| PREDICTED: ataxin-2 homolog isoform X1 [Drosophila miranda]
>gi|1036818985|ref|XP_017137538.1| PREDICTED: glutamic acid-rich protein [Drosophila miranda]
>gi|1036842659|ref|XP_017138857.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036842621|ref|XP_017138855.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036842640|ref|XP_017138856.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036842603|ref|XP_017138854.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036797036|ref|XP_017136317.1| PREDICTED: spermine oxidase [Drosophila miranda]
>gi|1036797175|ref|XP_017136326.1| PREDICTED: transmembrane protein 60 [Drosophila miranda]
>gi|1036797193|ref|XP_017136327.1| PREDICTED: transmembrane protein 60 [Drosophila miranda]
>gi|1036797017|ref|XP_017136316.1| PREDICTED: proline-rich protein PRCC [Drosophila miranda]
>gi|1036797100|ref|XP_017136321.1| PREDICTED: protein brother isoform X4 [Drosophila miranda]
>gi|1036797052|ref|XP_017136318.1| PREDICTED: protein brother isoform X1 [Drosophila miranda]
>gi|1036797085|ref|XP_017136320.1| PREDICTED: protein brother isoform X3 [Drosophila miranda]
>gi|1036797068|ref|XP_017136319.1| PREDICTED: protein brother isoform X2 [Drosophila miranda]
>gi|1036797118|ref|XP_017136322.1| PREDICTED: protein brother isoform X5 [Drosophila miranda]
>gi|1036797156|ref|XP_017136325.1| PREDICTED: transmembrane protein 35 [Drosophila miranda]
>gi|1036797137|ref|XP_017136324.1| PREDICTED: vacuolar protein sorting-associated protein VTA1
>gi|1036785913|ref|XP_017135692.1| PREDICTED: peroxisomal membrane protein 2 [Drosophila mirand
>gi|1036785929|ref|XP_017135693.1| PREDICTED: peroxisomal membrane protein 2 [Drosophila mirand
>gi|1036785816|ref|XP_017135686.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785778|ref|XP_017135684.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785833|ref|XP_017135687.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785797|ref|XP_017135685.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
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>gi|1036785852|ref|XP_017135688.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036785875|ref|XP_017135690.1| PREDICTED: uncharacterized protein C9orf78 [Drosophila mirar
>gi|1036785894|ref|XP_017135691.1| PREDICTED: 60S ribosomal protein L23a [Drosophila miranda]
>gi|1036764803|ref|XP_017157137.1| PREDICTED: ras-related protein Rab-34 [Drosophila miranda]
>gi|1036836651|ref|XP_017138522.1| PREDICTED: WD repeat-containing protein 78 [Drosophila mirar
>gi|1036836670|ref|XP_017138523.1| PREDICTED: WD repeat-containing protein 78 [Drosophila mirar
>gi|1036836616|ref|XP_017138521.1| PREDICTED: uncharacterized protein LOC108153210 [Drosophila
>gi|1036836689|ref|XP_017138524.1| PREDICTED: uncharacterized protein LOC108153212 [Drosophila
>gi|1036836710|ref|XP_017138525.1| PREDICTED: glycine-rich cell wall structural protein [Drosop
>gi|1036792113|ref|XP_017136041.1| PREDICTED: protein still life, isoform SIF type 1 isoform XS
>gi|1036792035|ref|XP_017136037.1| PREDICTED: protein still life, isoform SIF type 1 isoform X5
>gi|1036792094|ref|XP_017136040.1| PREDICTED: protein still life, isoform SIF type 1 isoform X8
>gi|1036792187|ref|XP_017136045.1| PREDICTED: protein still life, isoform SIF type 1 isoform X1
>gi|1036792168|ref|XP_017136044.1| PREDICTED: protein still life, isoform SIF type 1 isoform X1
>gi|1036792054|ref|XP_017136038.1| PREDICTED: protein still life, isoform SIF type 1 isoform X6
>gi|1036792150|ref|XP_017136043.1| PREDICTED: protein still life, isoform SIF type 1 isoform X1
>gi|1036792132|ref|XP_017136042.1| PREDICTED: protein still life, isoform SIF type 1 isoform X1
>gi|1036792073|ref|XP_017136039.1| PREDICTED: protein still life, isoform SIF type 1 isoform X7
>gi|1036791927|ref|XP_017136030.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791946|ref|XP_017136031.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791964|ref|XP_017136032.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791983|ref|XP_017136034.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036792001|ref|XP_017136035.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036792018|ref|XP_017136036.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791908|ref|XP_017136029.1| PREDICTED: protein still life, isoforms C/SIF type 2 isoforn
>gi|1036791890|ref|XP_017136028.1| PREDICTED: protein still life, isoforms C/SIF type 2 isoforn
>gi|1036791871|ref|XP_017136027.1| PREDICTED: protein still life, isoforms C/SIF type 2 isoforn
>gi|1036792202|ref|XP_017136046.1| PREDICTED: LOW QUALITY PROTEIN: separin [Drosophila miranda]
>gi|1036846308|ref|XP_017139049.1| PREDICTED: protein lin-28 homolog isoform X2 [Drosophila mir
>gi|1036846290|ref|XP_017139048.1| PREDICTED: protein lin-28 homolog isoform X1 [Drosophila mir
>gi|1036768887|ref|XP_017134707.1| PREDICTED: uncharacterized protein LOC108150904 isoform X2
>gi|1036768830|ref|XP_017134704.1| PREDICTED: uncharacterized protein LOC108150904 isoform X1
>gi|1036768849|ref|XP_017134705.1| PREDICTED: uncharacterized protein LOC108150904 isoform X1
>gi|1036768868|ref|XP_017134706.1| PREDICTED: uncharacterized protein LOC108150904 isoform X1
>gi|1036769248|ref|XP_017134729.1| PREDICTED: chromatin modification-related protein MEAF6 isoi
>gi|1036769267|ref|XP_017134730.1| PREDICTED: chromatin modification-related protein MEAF6 isoi
>gi|1036769286|ref|XP_017134731.1| PREDICTED: chromatin modification-related protein MEAF6 isoi
>gi|1036769122|ref|XP_017134721.1| PREDICTED: transmembrane 9 superfamily member 3 [Drosophila
>gi|1036769194|ref|XP_017134726.1| PREDICTED: collagen alpha-2(IX) chain [Drosophila miranda]
>gi|1036769159|ref|XP_017134723.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila r
>gi|1036769140|ref|XP_017134722.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila r
>gi|1036759457|ref|XP_017156838.1| PREDICTED: cytochrome P450 307a1 [Drosophila miranda]
>gi|1036768794|ref|XP_017134702.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1036768811|ref|XP_017134703.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1036769086|ref|XP_017134719.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036769105|ref|XP_017134720.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036769305|ref|XP_017134733.1| PREDICTED: bladder cancer-associated protein [Drosophila mir
>gi|1036769322|ref|XP_017134734.1| PREDICTED: bladder cancer-associated protein [Drosophila mir
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>gi|1036769213|ref|XP_017134727.1| PREDICTED: CAAX prenyl protease 2 [Drosophila miranda]
>gi|1036769176|ref|XP_017134725.1| PREDICTED: succinyl-CoA ligase [GDP-forming] subunit beta, m
>gi|1036769231|ref|XP_017134728.1| PREDICTED: uncharacterized protein LOC108150914 [Drosophila
>gi|1036768904|ref|XP_017134708.1| PREDICTED: collagen alpha-5(IV) chain isoform X1 [Drosophila
>gi|1036768922|ref|XP_017134709.1| PREDICTED: collagen alpha-5(IV) chain isoform X1 [Drosophila
>gi|1036768941|ref|XP_017134710.1| PREDICTED: collagen alpha-5(IV) chain isoform X2 [Drosophila
>gi|1036768960|ref|XP_017134711.1| PREDICTED: collagen alpha-1(X) chain isoform X3 [Drosophila
>gi|1036769001|ref|XP_017134713.1| PREDICTED: collagen alpha-1(X) chain isoform X5 [Drosophila
>gi|1036768982|ref|XP_017134712.1| PREDICTED: collagen alpha-1(I) chain isoform X4 [Drosophila
>gi|1036769018|ref|XP_017134714.1| PREDICTED: collagen alpha-1(I) chain isoform X6 [Drosophila
>gi|1036769035|ref|XP_017134716.1| PREDICTED: collagen alpha-1(I) chain isoform X6 [Drosophila
>gi|1036769051|ref|XP_017134717.1| PREDICTED: collagen alpha-1(III) chain isoform X7 [Drosophil
>gi|1036769067|ref|XP_017134718.1| PREDICTED: collagen alpha-1(V) chain isoform X8 [Drosophila
>gi|1036766665|ref|XP_017134586.1| PREDICTED: prisilkin-39 isoform X2 [Drosophila miranda]
>gi|1036766646|ref|XP_017134585.1| PREDICTED: prisilkin-39 isoform X1 [Drosophila miranda]
>gi|1036852398|ref|XP_017139409.1| PREDICTED: neuropeptide Y receptor type 1 [Drosophila miran
>gi|1036768612|ref|XP_017134690.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosop
>gi|1036767126|ref|XP_017134614.1| PREDICTED: uncharacterized protein LOC108150818 [Drosophila
>gi|1036759475|ref|XP_017156839.1| PREDICTED: uncharacterized protein LOC108165316 [Drosophila
>gi|1036784396|ref|XP_017135603.1| PREDICTED: probable prefoldin subunit 4 [Drosophila miranda]
>gi|1036784358|ref|XP_017135601.1| PREDICTED: probable Golgi SNAP receptor complex member 2 iso
>gi|1036784341|ref|XP_017135600.1| PREDICTED: probable Golgi SNAP receptor complex member 2 iso
>gi|1036784377|ref|XP_017135602.1| PREDICTED: ubiquitin-conjugating enzyme E2 variant 2 [Droso
>gi|1036784322|ref|XP_017135599.1| PREDICTED: cytochrome c1, heme protein, mitochondrial [Dros
>gi|1036784268|ref|XP_017135595.1| PREDICTED: kinesin-like protein KIF3A [Drosophila miranda]
>gi|1036784287|ref|XP_017135597.1| PREDICTED: putative phospholipase B-like lamina ancestor [D
>gi|1036784306|ref|XP_017135598.1| PREDICTED: putative phospholipase B-like lamina ancestor [D
>gi|1036843783|ref|XP_017138925.1| PREDICTED: protein alan shepard [Drosophila miranda]
>gi|1036843802|ref|XP_017138926.1| PREDICTED: protein alan shepard [Drosophila miranda]
>gi|1036759494|ref|XP_017156841.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036765309|ref|XP_017157163.1| PREDICTED: lachesin [Drosophila miranda]
>gi|1036766794|ref|XP_017134594.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036767206|ref|XP_017134618.1| PREDICTED: uncharacterized protein LOC108150823 [Drosophila
>gi|1036767225|ref|XP_017134619.1| PREDICTED: uncharacterized protein LOC108150823 [Drosophila
>gi|1036843397|ref|XP_017138902.1| PREDICTED: uncharacterized protein LOC108153413 isoform X3
>gi|1036843338|ref|XP_017138897.1| PREDICTED: uncharacterized protein LOC108153413 isoform X1
>gi|1036843357|ref|XP_017138898.1| PREDICTED: uncharacterized protein LOC108153413 isoform X1
>gi|1036843377|ref|XP_017138901.1| PREDICTED: uncharacterized protein LOC108153413 isoform X2
>gi|1036843452|ref|XP_017138905.1| PREDICTED: uncharacterized protein LOC108153415 [Drosophila
>gi|1036843414|ref|XP_017138903.1| PREDICTED: uncharacterized protein LOC108153413 isoform X4
>gi|1036843433|ref|XP_017138904.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036849873|ref|XP_017139256.1| PREDICTED: nucleolin [Drosophila miranda]
>gi|1036849892|ref|XP_017139257.1| PREDICTED: nucleolin [Drosophila miranda]
>gi|1036830552|ref|XP_017138183.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036830644|ref|XP_017138188.1| PREDICTED: phospholipid scramblase 1-like isoform X2 [Drosop
>gi|1036830624|ref|XP_017138187.1| PREDICTED: phospholipid scramblase 1-like isoform X1 [Drosop
>gi|1036830570|ref|XP_017138184.1| PREDICTED: phospholipid scramblase 2-like isoform X1 [Drosop
>gi|1036830586|ref|XP_017138185.1| PREDICTED: phospholipid scramblase 2-like isoform X2 [Drosop
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>gi|1036830605|ref|XP_017138186.1| PREDICTED: phospholipid scramblase 1-like isoform X3 [Droso
>gi|1036825593|ref|XP_017137894.1| PREDICTED: odorant receptor 67c [Drosophila miranda]
>gi|1036825554|ref|XP_017137891.1| PREDICTED: odorant receptor 67d isoform X1 [Drosophila mira
>gi|1036825572|ref|XP_017137892.1| PREDICTED: odorant receptor 67d isoform X2 [Drosophila mira
>gi|1036825633|ref|XP_017137896.1| PREDICTED: uncharacterized protein LOC108152818 isoform X2
>gi|1036825652|ref|XP_017137897.1| PREDICTED: uncharacterized protein LOC108152818 isoform X3
>gi|1036825614|ref|XP_017137895.1| PREDICTED: uncharacterized protein LOC108152818 isoform X1
>gi|1036851326|ref|XP_017139341.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036759513|ref|XP_017156842.1| PREDICTED: uncharacterized protein LOC108165319 [Drosophila
>gi|1036759532|ref|XP_017156843.1| PREDICTED: uncharacterized protein LOC108165320 [Drosophila
>gi|1036776402|ref|XP_017135128.1| PREDICTED: probable ATP-dependent RNA helicase DDX20 [Droso
>gi|1036776855|ref|XP_017135156.1| PREDICTED: uncharacterized protein LOC108151197 [Drosophila
>gi|1036776838|ref|XP_017135155.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1036776687|ref|XP_017135145.1| PREDICTED: AF4/FMR2 family member 4 isoform X14 [Drosophila
>gi|1036776721|ref|XP_017135147.1| PREDICTED: homeotic protein female sterile isoform X16 [Dros
>gi|1036776637|ref|XP_017135142.1| PREDICTED: AF4/FMR2 family member 4 isoform X11 [Drosophila
>gi|1036776668|ref|XP_017135144.1| PREDICTED: AF4/FMR2 family member 4 isoform X13 [Drosophila
>gi|1036776421|ref|XP_017135129.1| PREDICTED: ecdysone-induced protein 74EF isoform X1 [Drosop
>gi|1036776551|ref|XP_017135136.1| PREDICTED: ecdysone-induced protein 74EF isoform X6 [Drosop
>gi|1036776532|ref|XP_017135135.1| PREDICTED: ecdysone-induced protein 74EF isoform X5 [Drosop
>gi|1036776588|ref|XP_017135138.1| PREDICTED: ecdysone-induced protein 74EF isoform X8 [Drosop
>gi|1036776651|ref|XP_017135143.1| PREDICTED: AF4/FMR2 family member 4 isoform X12 [Drosophila
>gi|1036776440|ref|XP_017135130.1| PREDICTED: ecdysone-induced protein 74EF isoform X2 [Drosop
>gi|1036776570|ref|XP_017135137.1| PREDICTED: ecdysone-induced protein 74EF isoform X7 [Drosop
>gi|1036776607|ref|XP_017135139.1| PREDICTED: ecdysone-induced protein 74EF isoform X9 [Drosop
>gi|1036776623|ref|XP_017135140.1| PREDICTED: ecdysone-induced protein 74EF isoform X10 [Droso
>gi|1036776804|ref|XP_017135153.1| PREDICTED: ecdysone-induced protein 74EF isoform X19 [Droso
>gi|1036776821|ref|XP_017135154.1| PREDICTED: ecdysone-induced protein 74EF isoform X20 [Droso
>gi|1036776459|ref|XP_017135131.1| PREDICTED: ecdysone-induced protein 74EF isoform X3 [Drosop
>gi|1036776738|ref|XP_017135148.1| PREDICTED: homeotic protein female sterile isoform X17 [Dros
>gi|1036776755|ref|XP_017135149.1| PREDICTED: homeotic protein female sterile isoform X17 [Dros
>gi|1036776771|ref|XP_017135150.1| PREDICTED: homeotic protein female sterile isoform X17 [Dros
>gi|1036776787|ref|XP_017135151.1| PREDICTED: homeotic protein female sterile isoform X18 [Dros
>gi|1036776478|ref|XP_017135132.1| PREDICTED: ecdysone-induced protein 74EF isoform X4 [Drosop
>gi|1036776497|ref|XP_017135133.1| PREDICTED: ecdysone-induced protein 74EF isoform X4 [Drosop
>gi|1036776513|ref|XP_017135134.1| PREDICTED: ecdysone-induced protein 74EF isoform X4 [Drosop
>gi|1036776703|ref|XP_017135146.1| PREDICTED: ecdysone-induced protein 74EF isoform X15 [Droso
>gi|1036776872|ref|XP_017135158.1| PREDICTED: uncharacterized protein LOC108151198 [Drosophila
>gi|1036759551|ref|XP_017156844.1| PREDICTED: cytosolic non-specific dipeptidase-like [Drosophi
>gi|1036852379|ref|XP_017139407.1| PREDICTED: UNC93-like protein isoform X2 [Drosophila miranda
>gi|1036852360|ref|XP_017139406.1| PREDICTED: UNC93-like protein isoform X1 [Drosophila miranda
>gi|1036766939|ref|XP_017134603.1| PREDICTED: uncharacterized protein LOC108150809 isoform X2
>gi|1036766920|ref|XP_017134602.1| PREDICTED: uncharacterized protein LOC108150809 isoform X1
>gi|1036768649|ref|XP_017134693.1| PREDICTED: uncharacterized protein LOC108150894 [Drosophila
>gi|1036767484|ref|XP_017134632.1| PREDICTED: uncharacterized protein LOC108150836 [Drosophila
>gi|1036765364|ref|XP_017157166.1| PREDICTED: uncharacterized protein LOC108165603 isoform X1
>gi|1036765378|ref|XP_017157167.1| PREDICTED: uncharacterized protein LOC108165603 isoform X2
>gi|1036765393|ref|XP_017157168.1| PREDICTED: uncharacterized protein LOC108165603 isoform X3
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>gi|1036759570|ref|XP_017156845.1| PREDICTED: uncharacterized protein LOC108165322, partial [Di
>gi|1036782761|ref|XP_017135506.1| PREDICTED: chromosome transmission fidelity protein 8 homolo
>gi|1036782691|ref|XP_017135502.1| PREDICTED: DNA excision repair protein haywire [Drosophila m
>gi|1036782709|ref|XP_017135503.1| PREDICTED: histone-lysine N-methyltransferase E(z) [Drosophi
>gi|1036782779|ref|XP_017135508.1| PREDICTED: nuclear envelope phosphatase-regulatory subunit 1
>gi|1036782728|ref|XP_017135504.1| PREDICTED: protein FAM49B isoform X1 [Drosophila miranda]
>gi|1036782745|ref|XP_017135505.1| PREDICTED: protein FAM49B isoform X2 [Drosophila miranda]
>gi|1036804980|ref|XP_017136766.1| PREDICTED: ankyrin repeat and MYND domain-containing protein
>gi|1036804999|ref|XP_017136767.1| PREDICTED: 1,2-dihydroxy-3-keto-5-methylthiopentene dioxygen
>gi|1036804942|ref|XP_017136764.1| PREDICTED: ras guanine nucleotide exchange factor P isoform
>gi|1036804961|ref|XP_017136765.1| PREDICTED: ras guanine nucleotide exchange factor P isoform
>gi|1036805034|ref|XP_017136769.1| PREDICTED: U4/U6 small nuclear ribonucleoprotein Prp4 [Droso
>gi|1036805013|ref|XP_017136768.1| PREDICTED: solute carrier organic anion transporter family m
>gi|1036805088|ref|XP_017136772.1| PREDICTED: uncharacterized protein LOC108152164 [Drosophila
>gi|1036759588|ref|XP_017156846.1| PREDICTED: uncharacterized protein LOC108165323 [Drosophila
>gi|1036805071|ref|XP_017136771.1| PREDICTED: uncharacterized protein LOC108152163 [Drosophila
>gi|1036805053|ref|XP_017136770.1| PREDICTED: cytochrome b-c1 complex subunit 8 [Drosophila mir
>gi|1036848950|ref|XP_017139200.1| PREDICTED: glycine receptor subunit alphaZ1 [Drosophila mira
>gi|1036847401|ref|XP_017139113.1| PREDICTED: MICOS complex subunit MIC13 homolog QIL1 [Drosoph
>gi|1036795780|ref|XP_017136248.1| PREDICTED: uncharacterized protein LOC108151867 isoform X2
>gi|1036795727|ref|XP_017136245.1| PREDICTED: uncharacterized protein LOC108151867 isoform X1
>gi|1036795745|ref|XP_017136246.1| PREDICTED: uncharacterized protein LOC108151867 isoform X1
>gi|1036795763|ref|XP_017136247.1| PREDICTED: uncharacterized protein LOC108151867 isoform X1
>gi|1036795799|ref|XP_017136249.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1036795817|ref|XP_017136250.1| PREDICTED: uncharacterized protein LOC108151870 [Drosophila
>gi|1036795836|ref|XP_017136251.1| PREDICTED: UDP-sugar transporter UST74c [Drosophila miranda]
>gi|1036795874|ref|XP_017136254.1| PREDICTED: ubiquinone biosynthesis protein COQ4 homolog, mit
>gi|1036795893|ref|XP_017136255.1| PREDICTED: ubiquinone biosynthesis protein COQ4 homolog, mit
>gi|1036795912|ref|XP_017136256.1| PREDICTED: ubiquinone biosynthesis protein COQ4 homolog, mit
>gi|1036795855|ref|XP_017136252.1| PREDICTED: RAB6-interacting golgin [Drosophila miranda]
>gi|1036769992|ref|XP_017134770.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC7 [Dro
>gi|1036770030|ref|XP_017134772.1| PREDICTED: uncharacterized protein LOC108150950 isoform X1
>gi|1036770047|ref|XP_017134773.1| PREDICTED: uncharacterized protein LOC108150950 isoform X2
>gi|1036769895|ref|XP_017134765.1| PREDICTED: exocyst complex component 1 [Drosophila miranda]
>gi|1036769857|ref|XP_017134762.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036769876|ref|XP_017134764.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036769916|ref|XP_017134766.1| PREDICTED: LOW QUALITY PROTEIN: proline-rich extensin-like p
>gi|1036759606|ref|XP_017156847.1| PREDICTED: uncharacterized protein LOC108165324 [Drosophila
>gi|1036769741|ref|XP_017134756.1| PREDICTED: cadherin-87A [Drosophila miranda]
>gi|1036769760|ref|XP_017134757.1| PREDICTED: cadherin-87A [Drosophila miranda]
>gi|1036769954|ref|XP_017134768.1| PREDICTED: basic helix-loop-helix neural transcription facto
>gi|1036769799|ref|XP_017134759.1| PREDICTED: uncharacterized protein LOC108150939 isoform X2
>gi|1036769779|ref|XP_017134758.1| PREDICTED: uncharacterized protein LOC108150939 isoform X1
>gi|1036769818|ref|XP_017134760.1| PREDICTED: uncharacterized protein LOC108150939 isoform X3
>gi|1036769837|ref|XP_017134761.1| PREDICTED: uncharacterized protein LOC108150939 isoform X4
>gi|1036770011|ref|XP_017134771.1| PREDICTED: allatostatins MIP [Drosophila miranda]
>gi|1036759625|ref|XP_017156848.1| PREDICTED: polycystic kidney disease 2-like 2 protein [Droso
>gi|1036759642|ref|XP_017156849.1| PREDICTED: uncharacterized protein LOC108165327 [Drosophila
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>gi|1036769935|ref|XP_017134767.1| PREDICTED: transmembrane protein 209 [Drosophila miranda]
>gi|1036769973|ref|XP_017134769.1| PREDICTED: odorant receptor 74a [Drosophila miranda]
>gi|1036759658|ref|XP_017156850.1| PREDICTED: uncharacterized protein LOC108165328 [Drosophila
>gi|1036766761|ref|XP_017134592.1| PREDICTED: NADH-quinone oxidoreductase subunit E [Drosophila
>gi|1036766443|ref|XP_017134575.1| PREDICTED: uncharacterized protein LOC108150785 [Drosophila
>gi|1036832514|ref|XP_017138291.1| PREDICTED: UPF0687 protein C20orf27 homolog [Drosophila mira
>gi|1036792425|ref|XP_017136059.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792504|ref|XP_017136064.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792388|ref|XP_017136057.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792406|ref|XP_017136058.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792484|ref|XP_017136063.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792582|ref|XP_017136068.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792627|ref|XP_017136071.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792465|ref|XP_017136062.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792616|ref|XP_017136070.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792563|ref|XP_017136067.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792662|ref|XP_017136073.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792601|ref|XP_017136069.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792444|ref|XP_017136060.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792523|ref|XP_017136065.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792542|ref|XP_017136066.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792682|ref|XP_017136074.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792646|ref|XP_017136072.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036818174|ref|XP_017137491.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila m
>gi|1036818193|ref|XP_017137492.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila m
>gi|1036818212|ref|XP_017137493.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila m
>gi|1036818231|ref|XP_017137494.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila m
>gi|1036773858|ref|XP_017134991.1| PREDICTED: zinc finger CCCH domain-containing protein 3 [Dro
>gi|1036773747|ref|XP_017134984.1| PREDICTED: pseudouridylate synthase 7 homolog [Drosophila mi
>gi|1036773982|ref|XP_017134998.1| PREDICTED: uncharacterized protein LOC108151109 [Drosophila
>gi|1036773802|ref|XP_017134987.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036773821|ref|XP_017134988.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036773839|ref|XP_017134989.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036773963|ref|XP_017134997.1| PREDICTED: pyrimidodiazepine synthase [Drosophila miranda]
>gi|1036773875|ref|XP_017134992.1| PREDICTED: pyrimidodiazepine synthase [Drosophila miranda]
>gi|1036773946|ref|XP_017134996.1| PREDICTED: pyrimidodiazepine synthase isoform X2 [Drosophila
>gi|1036773928|ref|XP_017134995.1| PREDICTED: pyrimidodiazepine synthase isoform X1 [Drosophila
>gi|1036773894|ref|XP_017134993.1| PREDICTED: pyrimidodiazepine synthase isoform X1 [Drosophila
>gi|1036773909|ref|XP_017134994.1| PREDICTED: pyrimidodiazepine synthase isoform X2 [Drosophila
>gi|1036773766|ref|XP_017134985.1| PREDICTED: zinc transporter foi [Drosophila miranda]
>gi|1036773785|ref|XP_017134986.1| PREDICTED: zinc transporter foi [Drosophila miranda]
>gi|1036790045|ref|XP_017135925.1| PREDICTED: zinc finger protein 345 [Drosophila miranda]
>gi|1036789994|ref|XP_017135922.1| PREDICTED: E3 ubiquitin-protein ligase SHPRH isoform X1 [Dro
>gi|1036790011|ref|XP_017135923.1| PREDICTED: E3 ubiquitin-protein ligase SHPRH isoform X2 [Dro
>gi|1036790027|ref|XP_017135924.1| PREDICTED: zinc finger protein 91 [Drosophila miranda]
>gi|1036801261|ref|XP_017136558.1| PREDICTED: serine/threonine-protein phosphatase 6 regulatory
>gi|1036801280|ref|XP_017136559.1| PREDICTED: serine/threonine-protein phosphatase 6 regulatory
>gi|1036801301|ref|XP_017136560.1| PREDICTED: serine/threonine-protein phosphatase 6 regulatory
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>gi|1036764770|ref|XP_017157135.1| PREDICTED: Krueppel-like factor 7 [Drosophila miranda]
>gi|1036764787|ref|XP_017157136.1| PREDICTED: pupal cuticle protein 20 [Drosophila miranda]
>gi|1036768630|ref|XP_017134692.1| PREDICTED: uncharacterized protein LOC108150893 [Drosophila
>gi|1036768004|ref|XP_017134658.1| PREDICTED: uncharacterized protein LOC108150866 [Drosophila
>gi|1036765832|ref|XP_017134542.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036765852|ref|XP_017134543.1| PREDICTED: endocuticle structural glycoprotein ABD-5 [Drosop
>gi|1036767559|ref|XP_017134636.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036764187|ref|XP_017157105.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036764205|ref|XP_017157106.1| PREDICTED: larval cuticle protein 5-like [Drosophila miranda
>gi|1036766577|ref|XP_017134583.1| PREDICTED: larval cuticle protein 8 [Drosophila miranda]
>gi|1036767795|ref|XP_017134645.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036766175|ref|XP_017134562.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036767353|ref|XP_017134625.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036766902|ref|XP_017134601.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036766777|ref|XP_017134593.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036766721|ref|XP_017134590.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036764353|ref|XP_017157115.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036764373|ref|XP_017157116.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036849855|ref|XP_017139255.1| PREDICTED: protein lethal(3)malignant blood neoplasm 1 [Dros
>gi|1036849193|ref|XP_017139214.1| PREDICTED: DNA-binding protein D-ETS-3 isoform X1 [Drosophil
>gi|1036849211|ref|XP_017139215.1| PREDICTED: DNA-binding protein D-ETS-3 isoform X2 [Drosophil
>gi|1036759675|ref|XP_017156852.1| PREDICTED: DNA-binding protein D-ETS-3-like [Drosophila mir
>gi|1036842968|ref|XP_017138877.1| PREDICTED: transmembrane emp24 domain-containing protein B
>gi|1036797790|ref|XP_017136360.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036797810|ref|XP_017136361.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036797677|ref|XP_017136354.1| PREDICTED: protein SCAI isoform X4 [Drosophila miranda]
>gi|1036797644|ref|XP_017136352.1| PREDICTED: protein SCAI isoform X2 [Drosophila miranda]
>gi|1036797661|ref|XP_017136353.1| PREDICTED: protein SCAI isoform X3 [Drosophila miranda]
>gi|1036797613|ref|XP_017136350.1| PREDICTED: protein SCAI isoform X1 [Drosophila miranda]
>gi|1036797628|ref|XP_017136351.1| PREDICTED: protein SCAI isoform X1 [Drosophila miranda]
>gi|1036797576|ref|XP_017136348.1| PREDICTED: uncharacterized protein LOC108151934 [Drosophila
>gi|1036797595|ref|XP_017136349.1| PREDICTED: uncharacterized protein LOC108151934 [Drosophila
>gi|1036797826|ref|XP_017136362.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm5 [Drosop
>gi|1036797710|ref|XP_017136357.1| PREDICTED: T-cell activation inhibitor, mitochondrial isofo
>gi|1036797694|ref|XP_017136356.1| PREDICTED: T-cell activation inhibitor, mitochondrial isofo
>gi|1036797752|ref|XP_017136358.1| PREDICTED: T-cell activation inhibitor, mitochondrial isofo
>gi|1036797771|ref|XP_017136359.1| PREDICTED: uncharacterized protein LOC108151939 [Drosophila
>gi|1036815454|ref|XP_017137337.1| PREDICTED: organic cation transporter protein isoform X1 [D
>gi|1036815473|ref|XP_017137338.1| PREDICTED: organic cation transporter protein isoform X2 [D
>gi|1036815503|ref|XP_017137339.1| PREDICTED: organic cation transporter protein isoform X3 [D
>gi|1036815567|ref|XP_017137342.1| PREDICTED: brachyurin-like [Drosophila miranda]
>gi|1036759695|ref|XP_017156853.1| PREDICTED: LOW QUALITY PROTEIN: brachyurin-like [Drosophila
>gi|1036815525|ref|XP_017137340.1| PREDICTED: transmembrane protease serine 12 [Drosophila mir
>gi|1036815548|ref|XP_017137341.1| PREDICTED: protein tantalus [Drosophila miranda]
>gi|1036815646|ref|XP_017137347.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036815628|ref|XP_017137346.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036815605|ref|XP_017137345.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036815588|ref|XP_017137344.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
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>gi|1036759714|ref|XP_017156854.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036850286|ref|XP_017139280.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036759732|ref|XP_017156855.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036846510|ref|XP_017139062.1| PREDICTED: tensin-1 isoform X2 [Drosophila miranda]
>gi|1036846472|ref|XP_017139060.1| PREDICTED: protein sprint isoform X1 [Drosophila miranda]
>gi|1036846491|ref|XP_017139061.1| PREDICTED: protein sprint isoform X1 [Drosophila miranda]
>gi|1036846529|ref|XP_017139063.1| PREDICTED: serine protease 3-like [Drosophila miranda]
>gi|1036846548|ref|XP_017139064.1| PREDICTED: serine protease 3-like [Drosophila miranda]
>gi|1036846567|ref|XP_017139065.1| PREDICTED: serine protease 3-like [Drosophila miranda]
>gi|1036828928|ref|XP_017138088.1| PREDICTED: tensin-4 isoform X1 [Drosophila miranda]
>gi|1036828947|ref|XP_017138089.1| PREDICTED: uncharacterized protein LOC108152920 isoform X2
>gi|1036833857|ref|XP_017138359.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mir
>gi|1036833874|ref|XP_017138360.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mir
>gi|1036836748|ref|XP_017138527.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036836765|ref|XP_017138529.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036836838|ref|XP_017138533.1| PREDICTED: peptidoglycan-recognition protein LD isoform X4
>gi|1036836819|ref|XP_017138532.1| PREDICTED: peptidoglycan-recognition protein LD isoform X3
>gi|1036836801|ref|XP_017138531.1| PREDICTED: peptidoglycan-recognition protein LD isoform X2
>gi|1036836783|ref|XP_017138530.1| PREDICTED: peptidoglycan-recognition protein LD isoform X1
>gi|1036836857|ref|XP_017138534.1| PREDICTED: peptidoglycan-recognition protein LD isoform X5
>gi|1036836878|ref|XP_017138535.1| PREDICTED: transmembrane protein 11 homolog, mitochondrial
>gi|1036836897|ref|XP_017138536.1| PREDICTED: transmembrane protein 11 homolog, mitochondrial
>gi|1036837237|ref|XP_017138556.1| PREDICTED: membrane-associated tyrosine- and threonine-spec
>gi|1036837256|ref|XP_017138557.1| PREDICTED: membrane-associated tyrosine- and threonine-spec
>gi|1036803499|ref|XP_017136683.1| PREDICTED: nucleolar protein 12 [Drosophila miranda]
>gi|1036803442|ref|XP_017136680.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 36 isoform
>gi|1036803460|ref|XP_017136681.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 36 isoform
>gi|1036817291|ref|XP_017137441.1| PREDICTED: bifunctional coenzyme A synthase [Drosophila mira
>gi|1036817329|ref|XP_017137443.1| PREDICTED: thioredoxin-like protein 1 [Drosophila miranda]
>gi|1036817310|ref|XP_017137442.1| PREDICTED: proliferation-associated protein 2G4 [Drosophila
>gi|1036805557|ref|XP_017136799.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 64E [Drosop
>gi|1036805576|ref|XP_017136800.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 64E [Drosop
>gi|1036805595|ref|XP_017136801.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 64E [Drosop
>gi|1036844482|ref|XP_017138961.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036844499|ref|XP_017138962.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036829057|ref|XP_017138096.1| PREDICTED: R3H and coiled-coil domain-containing protein 1
>gi|1036829074|ref|XP_017138097.1| PREDICTED: R3H and coiled-coil domain-containing protein 1
>gi|1036826962|ref|XP_017137975.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036826981|ref|XP_017137976.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036827000|ref|XP_017137977.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036826923|ref|XP_017137973.1| PREDICTED: phospholipid scramblase 1 isoform X1 [Drosophila
>gi|1036826941|ref|XP_017137974.1| PREDICTED: phospholipid scramblase 2 isoform X2 [Drosophila
>gi|1036839630|ref|XP_017138696.1| PREDICTED: ankyrin repeat domain-containing protein 54 [Dros
>gi|1036827038|ref|XP_017137979.1| PREDICTED: 85/88 kDa calcium-independent phospholipase A2 is
>gi|1036827019|ref|XP_017137978.1| PREDICTED: 85/88 kDa calcium-independent phospholipase A2 is
>gi|1036808148|ref|XP_017136940.1| PREDICTED: protein phosphatase inhibitor 2 isoform X1 [Drosc
>gi|1036808167|ref|XP_017136941.1| PREDICTED: protein phosphatase inhibitor 2 isoform X2 [Drosc
>gi|1036808075|ref|XP_017136936.1| PREDICTED: mediator of RNA polymerase II transcription subun
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>gi|1036808020|ref|XP_017136933.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808039|ref|XP_017136934.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808001|ref|XP_017136932.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808111|ref|XP_017136938.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808056|ref|XP_017136935.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808092|ref|XP_017136937.1| PREDICTED: uncharacterized protein LOC108152243 isoform X6
>gi|1036808271|ref|XP_017136947.1| PREDICTED: probable insulin-like peptide 5 [Drosophila miran
>gi|1036808130|ref|XP_017136939.1| PREDICTED: acid sphingomyelinase-like phosphodiesterase 3b
>gi|1036808220|ref|XP_017136944.1| PREDICTED: probable insulin-like peptide 4 isoform X1 [Droso
>gi|1036808236|ref|XP_017136945.1| PREDICTED: probable insulin-like peptide 4 isoform X2 [Droso
>gi|1036808253|ref|XP_017136946.1| PREDICTED: probable insulin-like peptide 4 isoform X3 [Droso
>gi|1036808290|ref|XP_017136948.1| PREDICTED: probable insulin-like peptide 3 [Drosophila miran
>gi|1036808202|ref|XP_017136943.1| PREDICTED: probable insulin-like peptide 2 [Drosophila miran
>gi|1036808184|ref|XP_017136942.1| PREDICTED: probable insulin-like peptide 1 [Drosophila miran
>gi|1036807925|ref|XP_017136927.1| PREDICTED: synaptopodin 2-like protein isoform X1 [Drosophil
>gi|1036807944|ref|XP_017136929.1| PREDICTED: synaptopodin 2-like protein isoform X2 [Drosophil
>gi|1036807963|ref|XP_017136930.1| PREDICTED: synaptopodin 2-like protein isoform X3 [Drosophil
>gi|1036807981|ref|XP_017136931.1| PREDICTED: synaptopodin 2-like protein isoform X4 [Drosophil
>gi|1036822438|ref|XP_017137720.1| PREDICTED: anion exchange protein 3 isoform X2 [Drosophila r
>gi|1036822402|ref|XP_017137718.1| PREDICTED: anion exchange protein 2 isoform X1 [Drosophila r
>gi|1036822419|ref|XP_017137719.1| PREDICTED: anion exchange protein 2 isoform X1 [Drosophila r
>gi|1036820285|ref|XP_017137608.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1036820302|ref|XP_017137609.1| PREDICTED: homeobox protein prospero isoform X2 [Drosophila
>gi|1036820318|ref|XP_017137610.1| PREDICTED: homeobox protein prospero isoform X3 [Drosophila
>gi|1036820335|ref|XP_017137611.1| PREDICTED: homeobox protein prospero isoform X4 [Drosophila
>gi|1036820388|ref|XP_017137614.1| PREDICTED: homeobox protein prospero isoform X7 [Drosophila
>gi|1036820405|ref|XP_017137615.1| PREDICTED: homeobox protein prospero isoform X8 [Drosophila
>gi|1036820424|ref|XP_017137616.1| PREDICTED: homeobox protein prospero isoform X9 [Drosophila
>gi|1036820353|ref|XP_017137612.1| PREDICTED: homeobox protein prospero isoform X5 [Drosophila
>gi|1036820463|ref|XP_017137619.1| PREDICTED: homeobox protein prospero isoform X11 [Drosophila
>gi|1036820369|ref|XP_017137613.1| PREDICTED: homeobox protein prospero isoform X6 [Drosophila
>gi|1036820443|ref|XP_017137617.1| PREDICTED: homeobox protein prospero isoform X10 [Drosophila
>gi|1036820482|ref|XP_017137620.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036820499|ref|XP_017137621.1| PREDICTED: ATP synthase subunit b, mitochondrial [Drosophila
>gi|1036837200|ref|XP_017138554.1| PREDICTED: N-alpha-acetyltransferase 60 isoform X2 [Drosophi
>gi|1036837164|ref|XP_017138552.1| PREDICTED: N-alpha-acetyltransferase 60 isoform X1 [Drosophi
>gi|1036837182|ref|XP_017138553.1| PREDICTED: N-alpha-acetyltransferase 60 isoform X1 [Drosophi
>gi|1036759752|ref|XP_017156856.1| PREDICTED: integrator complex subunit 7 [Drosophila miranda]
>gi|1036817195|ref|XP_017137435.1| PREDICTED: nibrin [Drosophila miranda]
>gi|1036817252|ref|XP_017137438.1| PREDICTED: uncharacterized protein LOC108152539 [Drosophila
>gi|1036817270|ref|XP_017137439.1| PREDICTED: adrenodoxin [Drosophila miranda]
>gi|1036817214|ref|XP_017137436.1| PREDICTED: uncharacterized protein LOC108152538 isoform X1
>gi|1036817233|ref|XP_017137437.1| PREDICTED: uncharacterized protein LOC108152538 isoform X2
>gi|1036793701|ref|XP_017136128.1| PREDICTED: claspin [Drosophila miranda]
>gi|1036793757|ref|XP_017136131.1| PREDICTED: peroxisomal membrane protein PMP34 [Drosophila mi
>gi|1036793739|ref|XP_017136130.1| PREDICTED: THUMP domain-containing protein 1 homolog [Drosop
>gi|1036793720|ref|XP_017136129.1| PREDICTED: lysine-specific demethylase 8 [Drosophila miranda
>gi|1036793773|ref|XP_017136133.1| PREDICTED: B-cell receptor-associated protein 31 [Drosophila
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>gi|1036793795|ref|XP_017136134.1| PREDICTED: B-cell receptor-associated protein 31 [Drosophila
>gi|1036793812|ref|XP_017136135.1| PREDICTED: protein DPCD [Drosophila miranda]
>gi|1036793868|ref|XP_017136136.1| PREDICTED: uncharacterized protein LOC108151797 [Drosophila
>gi|1036842984|ref|XP_017138878.1| PREDICTED: UPF0415 protein C7orf25 homolog [Drosophila mira
>gi|1036834602|ref|XP_017138400.1| PREDICTED: uncharacterized protein LOC108153123 [Drosophila
>gi|1036834585|ref|XP_017138399.1| PREDICTED: dimethyladenosine transferase 1, mitochondrial [
>gi|1036834621|ref|XP_017138402.1| PREDICTED: cytochrome c oxidase assembly factor 3, mitochond
>gi|1036816225|ref|XP_017137379.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816244|ref|XP_017137381.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816263|ref|XP_017137382.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816282|ref|XP_017137383.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816152|ref|XP_017137375.1| PREDICTED: dynein intermediate chain 3, ciliary [Drosophila
>gi|1036816171|ref|XP_017137376.1| PREDICTED: diphthamide biosynthesis protein 1 [Drosophila m
>gi|1036816188|ref|XP_017137377.1| PREDICTED: diphthamide biosynthesis protein 1 [Drosophila m
>gi|1036816206|ref|XP_017137378.1| PREDICTED: diphthamide biosynthesis protein 1 [Drosophila m
>gi|1036801688|ref|XP_017136583.1| PREDICTED: uncharacterized protein LOC108152071 [Drosophila
>gi|1036801706|ref|XP_017136584.1| PREDICTED: uncharacterized protein LOC108152071 [Drosophila
>gi|1036801724|ref|XP_017136585.1| PREDICTED: biogenesis of lysosome-related organelles comple
>gi|1036801653|ref|XP_017136581.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801634|ref|XP_017136580.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801672|ref|XP_017136582.1| PREDICTED: general transcription factor IIE subunit 1 [Droso
>gi|1036839157|ref|XP_017138675.1| PREDICTED: F-box/LRR-repeat protein 16 isoform X1 [Drosophil
>gi|1036839176|ref|XP_017138676.1| PREDICTED: F-box/LRR-repeat protein 16 isoform X2 [Drosophil
>gi|1036839194|ref|XP_017138677.1| PREDICTED: uncharacterized protein LOC108153298 [Drosophila
>gi|1036816592|ref|XP_017137401.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC8 [Dro
>gi|1036816573|ref|XP_017137400.1| PREDICTED: serine/threonine-protein kinase RIO1 [Drosophila
>gi|1036816534|ref|XP_017137397.1| PREDICTED: ATP-binding cassette sub-family G member 1 isofor
>gi|1036816553|ref|XP_017137399.1| PREDICTED: ATP-binding cassette sub-family G member 1 isofor
>gi|1036833084|ref|XP_017138324.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833102|ref|XP_017138325.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833121|ref|XP_017138326.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833138|ref|XP_017138328.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833156|ref|XP_017138329.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833229|ref|XP_017138333.1| PREDICTED: acyl-protein thioesterase 1 isoform X3 [Drosophil
>gi|1036833174|ref|XP_017138330.1| PREDICTED: acyl-protein thioesterase 1 isoform X2 [Drosophil
>gi|1036833193|ref|XP_017138331.1| PREDICTED: acyl-protein thioesterase 1 isoform X2 [Drosophil
>gi|1036833212|ref|XP_017138332.1| PREDICTED: acyl-protein thioesterase 1 isoform X2 [Drosophil
>gi|1036838610|ref|XP_017138643.1| PREDICTED: PCI domain-containing protein 2 homolog [Drosophi
>gi|1036848678|ref|XP_017139184.1| PREDICTED: uncharacterized protein LOC108153603 isoform X2
>gi|1036848659|ref|XP_017139183.1| PREDICTED: uncharacterized protein LOC108153603 isoform X1
>gi|1036848640|ref|XP_017139182.1| PREDICTED: alcohol dehydrogenase [NADP(+)] [Drosophila mira
>gi|1036842273|ref|XP_017138839.1| PREDICTED: aldose reductase isoform X1 [Drosophila miranda]
>gi|1036842328|ref|XP_017138840.1| PREDICTED: aldose reductase isoform X2 [Drosophila miranda]
>gi|1036849429|ref|XP_017139229.1| PREDICTED: putative protein TPRXL [Drosophila miranda]
>gi|1036849413|ref|XP_017139227.1| PREDICTED: uncharacterized serine-rich protein C215.13-like
>gi|1036830884|ref|XP_017138202.1| PREDICTED: uncharacterized protein LOC108152986 isoform X1
>gi|1036830922|ref|XP_017138204.1| PREDICTED: uncharacterized protein LOC108152986 isoform X3
>gi|1036830903|ref|XP_017138203.1| PREDICTED: uncharacterized protein LOC108152986 isoform X2
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>gi|1036830943|ref|XP_017138205.1| PREDICTED: uncharacterized protein LOC108152986 isoform X4
>gi|1036823934|ref|XP_017137805.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036823953|ref|XP_017137806.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036823971|ref|XP_017137807.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036824063|ref|XP_017137813.1| PREDICTED: uncharacterized protein LOC108152760 [Drosophila
>gi|1036824046|ref|XP_017137812.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036823897|ref|XP_017137803.1| PREDICTED: protein O-mannosyltransferase 1 [Drosophila miran
>gi|1036823916|ref|XP_017137804.1| PREDICTED: protein O-mannosyltransferase 1 [Drosophila miran
>gi|1036823990|ref|XP_017137809.1| PREDICTED: uncharacterized protein LOC108152757 [Drosophila
>gi|1036824029|ref|XP_017137811.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036824010|ref|XP_017137810.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036799366|ref|XP_017136450.1| PREDICTED: MOB kinase activator-like 2 isoform X4 [Drosophil
>gi|1036799328|ref|XP_017136448.1| PREDICTED: MOB kinase activator-like 2 isoform X2 [Drosophil
>gi|1036799309|ref|XP_017136447.1| PREDICTED: MOB kinase activator-like 2 isoform X1 [Drosophil
>gi|1036799422|ref|XP_017136453.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1036799347|ref|XP_017136449.1| PREDICTED: MOB kinase activator-like 2 isoform X3 [Drosophil
>gi|1036799385|ref|XP_017136451.1| PREDICTED: putative alpha-L-fucosidase [Drosophila miranda]
>gi|1036799404|ref|XP_017136452.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1036759771|ref|XP_017156857.1| PREDICTED: uncharacterized protein LOC108165334 [Drosophila
>gi|1036799441|ref|XP_017136454.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1036806637|ref|XP_017136857.1| PREDICTED: protein charybde isoform X2 [Drosophila miranda]
>gi|1036806619|ref|XP_017136856.1| PREDICTED: protein charybde isoform X1 [Drosophila miranda]
>gi|1036847028|ref|XP_017139091.1| PREDICTED: uncharacterized protein LOC108153537 isoform X2
>gi|1036847009|ref|XP_017139090.1| PREDICTED: uncharacterized protein LOC108153537 isoform X1
>gi|1036759788|ref|XP_017156858.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036850578|ref|XP_017139297.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036806526|ref|XP_017136850.1| PREDICTED: uncharacterized protein LOC108152202 [Drosophila
>gi|1036806582|ref|XP_017136853.1| PREDICTED: androgen-induced gene 1 protein [Drosophila miran
>gi|1036806489|ref|XP_017136848.1| PREDICTED: methylenetetrahydrofolate reductase 1 [Drosophila
>gi|1036806601|ref|XP_017136855.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036806545|ref|XP_017136851.1| PREDICTED: eukaryotic translation initiation factor 4H [Dros
>gi|1036806471|ref|XP_017136847.1| PREDICTED: YLP motif-containing protein 1 [Drosophila miranc
>gi|1036806563|ref|XP_017136852.1| PREDICTED: uncharacterized protein LOC108152205 [Drosophila
>gi|1036806507|ref|XP_017136849.1| PREDICTED: glutaminyl-peptide cyclotransferase-like protein
>gi|1036759807|ref|XP_017156859.1| PREDICTED: CAAX prenyl protease 1 homolog [Drosophila miranc
>gi|1036840850|ref|XP_017138764.1| PREDICTED: AF4/FMR2 family member 4 [Drosophila miranda]
>gi|1036848257|ref|XP_017139160.1| PREDICTED: gamma-tubulin complex component 2 homolog isofor
>gi|1036848238|ref|XP_017139159.1| PREDICTED: gamma-tubulin complex component 2 homolog isofor
>gi|1036833947|ref|XP_017138364.1| PREDICTED: protein scylla [Drosophila miranda]
>gi|1036814350|ref|XP_017137274.1| PREDICTED: uncharacterized protein LOC108152442 isoform X1
>gi|1036814369|ref|XP_017137277.1| PREDICTED: uncharacterized protein LOC108152442 isoform X2
>gi|1036814213|ref|XP_017137267.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814232|ref|XP_017137268.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814253|ref|XP_017137269.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814272|ref|XP_017137270.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814291|ref|XP_017137271.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814329|ref|XP_017137273.1| PREDICTED: uncharacterized protein LOC108152440 isoform X3
>gi|1036814310|ref|XP_017137272.1| PREDICTED: uncharacterized protein LOC108152440 isoform X2
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>gi|1036810546|ref|XP_017137070.1| PREDICTED: protein FAM91A1 [Drosophila miranda]
>gi|1036810491|ref|XP_017137067.1| PREDICTED: ubiquitin-protein ligase E3A isoform X1 [Drosophi
>gi|1036810509|ref|XP_017137068.1| PREDICTED: ubiquitin-protein ligase E3A isoform X2 [Drosophi
>gi|1036810528|ref|XP_017137069.1| PREDICTED: ubiquitin-protein ligase E3A isoform X3 [Drosophi
>gi|1036825126|ref|XP_017137872.1| PREDICTED: procollagen-lysine,2-oxoglutarate 5-dioxygenase
>gi|1036825144|ref|XP_017137873.1| PREDICTED: immunoglobulin domain-containing protein oig-1 [
>gi|1036825164|ref|XP_017137874.1| PREDICTED: immunoglobulin domain-containing protein oig-1 [
>gi|1036824867|ref|XP_017137858.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucuro
>gi|1036824904|ref|XP_017137859.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucuro
>gi|1036767759|ref|XP_017134644.1| PREDICTED: protein FAM188B2 [Drosophila miranda]
>gi|1036793284|ref|XP_017136107.1| PREDICTED: uncharacterized aarF domain-containing protein ki
>gi|1036793339|ref|XP_017136109.1| PREDICTED: uncharacterized protein LOC108151775 [Drosophila
>gi|1036793265|ref|XP_017136106.1| PREDICTED: protein wntless [Drosophila miranda]
>gi|1036793305|ref|XP_017136108.1| PREDICTED: putative Dol-P-Glc:Glc(2)Man(9)GlcNAc(2)-PP-Dol a
>gi|1036793246|ref|XP_017136105.1| PREDICTED: brain-specific angiogenesis inhibitor 1-associate
>gi|1036793210|ref|XP_017136104.1| PREDICTED: putative uncharacterized protein DDB_G0277255 iso
>gi|1036793358|ref|XP_017136110.1| PREDICTED: uncharacterized protein LOC108151776 [Drosophila
>gi|1036823224|ref|XP_017137766.1| PREDICTED: sodium-dependent phosphate transporter 1-B [Droso
>gi|1036823243|ref|XP_017137767.1| PREDICTED: sodium-dependent phosphate transporter 1-B [Droso
>gi|1036843637|ref|XP_017138916.1| PREDICTED: uncharacterized protein LOC108153424 [Drosophila
>gi|1036813914|ref|XP_017137250.1| PREDICTED: forkhead box protein K2 isoform X2 [Drosophila mi
>gi|1036813876|ref|XP_017137247.1| PREDICTED: forkhead box protein K2 isoform X1 [Drosophila mi
>gi|1036813894|ref|XP_017137248.1| PREDICTED: forkhead box protein K2 isoform X1 [Drosophila mi
>gi|1036813932|ref|XP_017137251.1| PREDICTED: 39S ribosomal protein L2, mitochondrial [Drosophi
>gi|1036813950|ref|XP_017137252.1| PREDICTED: ubiquitin fusion degradation protein 1 homolog [
>gi|1036829033|ref|XP_017138095.1| PREDICTED: superoxide dismutase [Cu-Zn] isoform X2 [Drosophi
>gi|1036829015|ref|XP_017138094.1| PREDICTED: superoxide dismutase [Cu-Zn] isoform X1 [Drosophi
>gi|1036829003|ref|XP_017138092.1| PREDICTED: transmembrane protein 161B [Drosophila miranda]
>gi|1036851458|ref|XP_017139349.1| PREDICTED: protein new-glue 2-like [Drosophila miranda]
>gi|1036851472|ref|XP_017139350.1| PREDICTED: uncharacterized protein LOC108153732 [Drosophila
>gi|1036769705|ref|XP_017134755.1| PREDICTED: uncharacterized protein LOC108150936 [Drosophila
>gi|1036759825|ref|XP_017156860.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036819553|ref|XP_017137567.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036819513|ref|XP_017137565.1| PREDICTED: nucleobindin-2 [Drosophila miranda]
>gi|1036819534|ref|XP_017137566.1| PREDICTED: probable ATP-dependent RNA helicase DDX52 [Drosop
>gi|1036845261|ref|XP_017138998.1| PREDICTED: uncharacterized protein LOC108153475 [Drosophila
>gi|1036845244|ref|XP_017138997.1| PREDICTED: arylsulfatase B [Drosophila miranda]
>gi|1036849909|ref|XP_017139258.1| PREDICTED: battenin [Drosophila miranda]
>gi|1036849928|ref|XP_017139259.1| PREDICTED: battenin [Drosophila miranda]
>gi|1036849947|ref|XP_017139261.1| PREDICTED: battenin [Drosophila miranda]
>gi|1036765695|ref|XP_017134534.1| PREDICTED: glutamate receptor ionotropic, kainate 4 [Drosoph
>gi|1036765727|ref|XP_017134535.1| PREDICTED: glutamate receptor ionotropic, kainate 4 [Drosoph
>gi|1036759844|ref|XP_017156861.1| PREDICTED: glutamate receptor ionotropic, kainate 5 [Drosoph
>gi|1036759863|ref|XP_017156862.1| PREDICTED: gamma-aminobutyric acid receptor alpha-like [Dros
>gi|1036778515|ref|XP_017135258.1| PREDICTED: EF-hand domain-containing family member B-like [
>gi|1036778554|ref|XP_017135260.1| PREDICTED: glycerol-3-phosphate phosphatase [Drosophila mira
>gi|1036778535|ref|XP_017135259.1| PREDICTED: glycerol-3-phosphate phosphatase [Drosophila mira
>gi|1036778665|ref|XP_017135267.1| PREDICTED: mediator of RNA polymerase II transcription subun
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>gi|1036778646|ref|XP_017135266.1| PREDICTED: dihydrolipoyl dehydrogenase, mitochondrial [Dros
>gi|1036778573|ref|XP_017135261.1| PREDICTED: tryptophan--tRNA ligase [Drosophila miranda]
>gi|1036778592|ref|XP_017135262.1| PREDICTED: cationic amino acid transporter 2 [Drosophila mi
>gi|1036778610|ref|XP_017135263.1| PREDICTED: arylsulfatase B isoform X1 [Drosophila miranda]
>gi|1036778629|ref|XP_017135265.1| PREDICTED: arylsulfatase B isoform X2 [Drosophila miranda]
>gi|1036778684|ref|XP_017135268.1| PREDICTED: uncharacterized protein LOC108151261 [Drosophila
>gi|1036766561|ref|XP_017134581.1| PREDICTED: uncharacterized protein LOC108150790 [Drosophila
>gi|1036759882|ref|XP_017156863.1| PREDICTED: arylsulfatase B [Drosophila miranda]
>gi|1036844991|ref|XP_017138982.1| PREDICTED: CD151 antigen [Drosophila miranda]
>gi|1036845010|ref|XP_017138983.1| PREDICTED: CD151 antigen [Drosophila miranda]
>gi|1036841752|ref|XP_017138819.1| PREDICTED: LOW QUALITY PROTEIN: solute carrier family 26 mem
>gi|1036821183|ref|XP_017137650.1| PREDICTED: WD repeat-containing protein 92 [Drosophila miran
>gi|1036821199|ref|XP_017137651.1| PREDICTED: WD repeat-containing protein 92 [Drosophila miran
>gi|1036821163|ref|XP_017137649.1| PREDICTED: LOW QUALITY PROTEIN: tetratricopeptide repeat pro
>gi|1036849999|ref|XP_017139264.1| PREDICTED: uncharacterized protein LOC108153663 [Drosophila
>gi|1036850019|ref|XP_017139266.1| PREDICTED: uncharacterized protein LOC108153663 [Drosophila
>gi|1036759901|ref|XP_017156864.1| PREDICTED: dihydrolipoyl dehydrogenase-like [Drosophila mira
>gi|1036801175|ref|XP_017136553.1| PREDICTED: ecdysone-induced protein 75B isoform X1 [Drosophi
>gi|1036801225|ref|XP_017136556.1| PREDICTED: ecdysone-induced protein 75B, isoforms C/D isofor
>gi|1036801208|ref|XP_017136555.1| PREDICTED: ecdysone-induced protein 75B, isoforms C/D isofor
>gi|1036801192|ref|XP_017136554.1| PREDICTED: ecdysone-induced protein 75B isoform X2 [Drosophi
>gi|1036801242|ref|XP_017136557.1| PREDICTED: ecdysone-induced protein 75B, isoforms C/D isofor
>gi|1036759920|ref|XP_017156865.1| PREDICTED: uncharacterized protein LOC108165343 [Drosophila
>gi|1036849767|ref|XP_017139249.1| PREDICTED: uncharacterized protein LOC108153654 [Drosophila
>gi|1036849785|ref|XP_017139250.1| PREDICTED: uncharacterized protein LOC108153655 isoform X1
>gi|1036849803|ref|XP_017139252.1| PREDICTED: uncharacterized protein LOC108153655 isoform X2
>gi|1036765659|ref|XP_017157176.1| PREDICTED: uncharacterized protein LOC108165611 [Drosophila
>gi|1036829193|ref|XP_017138104.1| PREDICTED: uncharacterized protein LOC108152929 [Drosophila
>gi|1036829212|ref|XP_017138105.1| PREDICTED: uncharacterized protein LOC108152929 [Drosophila
>gi|1036829231|ref|XP_017138106.1| PREDICTED: probable 28S ribosomal protein S26, mitochondrial
>gi|1036829172|ref|XP_017138103.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC4 [Dro
>gi|1036787005|ref|XP_017135751.1| PREDICTED: ATR-interacting protein mus304 [Drosophila miranc
>gi|1036787043|ref|XP_017135753.1| PREDICTED: uncharacterized protein LOC108151585 [Drosophila
>gi|1036787024|ref|XP_017135752.1| PREDICTED: probable cytochrome P450 312a1 [Drosophila miranc
>gi|1036787061|ref|XP_017135754.1| PREDICTED: uncharacterized protein LOC108151586 isoform X1
>gi|1036787080|ref|XP_017135755.1| PREDICTED: uncharacterized protein LOC108151586 isoform X2
>gi|1036787116|ref|XP_017135757.1| PREDICTED: uncharacterized protein LOC108151589 [Drosophila
>gi|1036787135|ref|XP_017135758.1| PREDICTED: uncharacterized protein LOC108151590 [Drosophila
>gi|1036787153|ref|XP_017135759.1| PREDICTED: uncharacterized protein LOC108151591 [Drosophila
>gi|1036787099|ref|XP_017135756.1| PREDICTED: uncharacterized protein LOC108151587 [Drosophila
>gi|1036831774|ref|XP_017138251.1| PREDICTED: cell death protein hid [Drosophila miranda]
>gi|1036850056|ref|XP_017139268.1| PREDICTED: larval serum protein 1 gamma chain [Drosophila mi
>gi|1036767164|ref|XP_017134616.1| PREDICTED: uncharacterized protein LOC108150820 [Drosophila
>gi|1036850670|ref|XP_017139303.1| PREDICTED: uncharacterized protein LOC108153690 [Drosophila
>gi|1036849494|ref|XP_017139232.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036763349|ref|XP_017157060.1| PREDICTED: cell death protein Grim [Drosophila miranda]
>gi|1036766326|ref|XP_017134566.1| PREDICTED: uncharacterized protein LOC108150780 isoform X1
>gi|1036766347|ref|XP_017134568.1| PREDICTED: uncharacterized protein LOC108150780 isoform X2
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>gi|1036849580|ref|XP_017139238.1| PREDICTED: uncharacterized protein LOC108153645 [Drosophila
>gi|1036764169|ref|XP_017157104.1| PREDICTED: coactosin-like protein [Drosophila miranda]
>gi|1036765778|ref|XP_017134539.1| PREDICTED: mitochondrial dicarboxylate carrier [Drosophila
>gi|1036763136|ref|XP_017157051.1| PREDICTED: cell death protein rpr [Drosophila miranda]
>gi|1036832217|ref|XP_017138277.1| PREDICTED: uncharacterized protein LOC108153032 [Drosophila
>gi|1036832198|ref|XP_017138276.1| PREDICTED: gamma-glutamylcyclotransferase [Drosophila miran
>gi|1036832181|ref|XP_017138275.1| PREDICTED: gamma-glutamylcyclotransferase [Drosophila miran
>gi|1036835683|ref|XP_017138464.1| PREDICTED: angiopoietin-related protein 1-like [Drosophila
>gi|1036835630|ref|XP_017138461.1| PREDICTED: somatostatin receptor type 2 [Drosophila miranda]
>gi|1036835701|ref|XP_017138465.1| PREDICTED: uncharacterized protein LOC108153171 [Drosophila
>gi|1036835647|ref|XP_017138462.1| PREDICTED: somatostatin receptor type 5 [Drosophila miranda]
>gi|1036835666|ref|XP_017138463.1| PREDICTED: somatostatin receptor type 5 [Drosophila miranda]
>gi|1036838106|ref|XP_017138610.1| PREDICTED: protein terminus [Drosophila miranda]
>gi|1036825411|ref|XP_017137889.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Drosop
>gi|1036825392|ref|XP_017137888.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1036825375|ref|XP_017137886.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1036825356|ref|XP_017137885.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1036825430|ref|XP_017137890.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1036825337|ref|XP_017137884.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1036825299|ref|XP_017137882.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 isoform X1
>gi|1036825318|ref|XP_017137883.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 isoform X2
>gi|1036842403|ref|XP_017138843.1| PREDICTED: uncharacterized protein LOC108153381 [Drosophila
>gi|1036808882|ref|XP_017136980.1| PREDICTED: LOW QUALITY PROTEIN: ubiquitin carboxyl-terminal
>gi|1036808901|ref|XP_017136982.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase nonstop-li
>gi|1036808863|ref|XP_017136979.1| PREDICTED: LOW QUALITY PROTEIN: protein aurora borealis [Dro
>gi|1036819061|ref|XP_017137541.1| PREDICTED: uncharacterized protein LOC108152597 [Drosophila
>gi|1036819080|ref|XP_017137543.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A [Dro
>gi|1036763295|ref|XP_017157057.1| PREDICTED: uncharacterized protein LOC108165512 [Drosophila
>gi|1036788386|ref|XP_017135827.1| PREDICTED: caprin homolog [Drosophila miranda]
>gi|1036788454|ref|XP_017135832.1| PREDICTED: gram-negative bacteria-binding protein 2 [Drosoph
>gi|1036788404|ref|XP_017135828.1| PREDICTED: gram-negative bacteria-binding protein 1 isoform
>gi|1036788420|ref|XP_017135829.1| PREDICTED: gram-negative bacteria-binding protein 2 isoform
>gi|1036788437|ref|XP_017135830.1| PREDICTED: gram-negative bacteria-binding protein 2-like [Dr
>gi|1036788471|ref|XP_017135833.1| PREDICTED: SAGA-associated factor 11 homolog [Drosophila mir
>gi|1036788503|ref|XP_017135835.1| PREDICTED: uncharacterized protein LOC108151640 [Drosophila
>gi|1036788487|ref|XP_017135834.1| PREDICTED: charged multivesicular body protein 1b [Drosophil
>gi|1036847180|ref|XP_017139100.1| PREDICTED: probable cytochrome P450 12c1, mitochondrial [Dro
>gi|1036759939|ref|XP_017156866.1| PREDICTED: mitochondrial dicarboxylate carrier, partial [Dro
>gi|1036787986|ref|XP_017135807.1| PREDICTED: uncharacterized protein LOC108151619 [Drosophila
>gi|1036787821|ref|XP_017135799.1| PREDICTED: uncharacterized protein LOC108151613 [Drosophila
>gi|1036787859|ref|XP_017135801.1| PREDICTED: uncharacterized protein LOC108151615 [Drosophila
>gi|1036787840|ref|XP_017135800.1| PREDICTED: equilibrative nucleoside transporter 4 [Drosophil
>gi|1036787915|ref|XP_017135804.1| PREDICTED: WW domain-binding protein 2 isoform X1 [Drosophil
>gi|1036787968|ref|XP_017135806.1| PREDICTED: WW domain-binding protein 2 isoform X3 [Drosophil
>gi|1036787932|ref|XP_017135805.1| PREDICTED: WW domain-binding protein 2 isoform X2 [Drosophil
>gi|1036787710|ref|XP_017135792.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X1
>gi|1036787727|ref|XP_017135793.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 1 isoform X2
>gi|1036787783|ref|XP_017135796.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X5
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>gi|1036787802|ref|XP_017135797.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X6
>gi|1036787764|ref|XP_017135795.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 1 isoform X4
>gi|1036787746|ref|XP_017135794.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X3
>gi|1036787877|ref|XP_017135802.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1036787896|ref|XP_017135803.1| PREDICTED: facilitated trehalose transporter Tret1 isoform X
>gi|1036810362|ref|XP_017137059.1| PREDICTED: titin [Drosophila miranda]
>gi|1036818506|ref|XP_017137510.1| PREDICTED: splicing factor 3A subunit 2 [Drosophila miranda]
>gi|1036818487|ref|XP_017137509.1| PREDICTED: mucin-19 isoform X4 [Drosophila miranda]
>gi|1036818469|ref|XP_017137507.1| PREDICTED: mucin-19 isoform X3 [Drosophila miranda]
>gi|1036818450|ref|XP_017137506.1| PREDICTED: mucin-19 isoform X2 [Drosophila miranda]
>gi|1036818432|ref|XP_017137505.1| PREDICTED: mucin-19 isoform X1 [Drosophila miranda]
>gi|1036809408|ref|XP_017137006.1| PREDICTED: serine/threonine-protein kinase unc-51 [Drosophil
>gi|1036809427|ref|XP_017137007.1| PREDICTED: leucine-rich repeat and transmembrane domain-cont
>gi|1036850037|ref|XP_017139267.1| PREDICTED: uncharacterized protein LOC108153664 [Drosophila
>gi|1036759958|ref|XP_017156867.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1036816362|ref|XP_017137387.1| PREDICTED: uncharacterized protein LOC108152508 [Drosophila
>gi|1036816381|ref|XP_017137388.1| PREDICTED: AP2-associated protein kinase 1 [Drosophila miran
>gi|1036759975|ref|XP_017156868.1| PREDICTED: uncharacterized protein LOC108165346 [Drosophila
>gi|1036816479|ref|XP_017137394.1| PREDICTED: uncharacterized protein C16orf52 homolog A [Droso
>gi|1036816322|ref|XP_017137385.1| PREDICTED: adenylate cyclase type 2 isoform X2 [Drosophila r
>gi|1036816303|ref|XP_017137384.1| PREDICTED: adenylate cyclase type 2 isoform X1 [Drosophila r
>gi|1036816341|ref|XP_017137386.1| PREDICTED: adenylate cyclase type 2 isoform X3 [Drosophila r
>gi|1036816498|ref|XP_017137395.1| PREDICTED: uncharacterized protein LOC108152513 [Drosophila
>gi|1036816400|ref|XP_017137390.1| PREDICTED: uncharacterized protein LOC108152510 isoform X1
>gi|1036816421|ref|XP_017137391.1| PREDICTED: uncharacterized protein LOC108152510 isoform X2
>gi|1036816440|ref|XP_017137392.1| PREDICTED: uncharacterized protein LOC108152510 isoform X3
>gi|1036816458|ref|XP_017137393.1| PREDICTED: uncharacterized protein LOC108152511 [Drosophila
>gi|1036816516|ref|XP_017137396.1| PREDICTED: uncharacterized protein LOC108152514 [Drosophila
>gi|1036804217|ref|XP_017136723.1| PREDICTED: rho GDP-dissociation inhibitor 1 [Drosophila mira
>gi|1036804236|ref|XP_017136724.1| PREDICTED: rho GDP-dissociation inhibitor 1 [Drosophila mira
>gi|1036804198|ref|XP_017136722.1| PREDICTED: metaxin-2 [Drosophila miranda]
>gi|1036804254|ref|XP_017136725.1| PREDICTED: larval/pupal cuticle protein H1C-like [Drosophila
>gi|1036804179|ref|XP_017136720.1| PREDICTED: Golgi reassembly-stacking protein 2 [Drosophila r
>gi|1036804273|ref|XP_017136726.1| PREDICTED: probable prefoldin subunit 6 [Drosophila miranda]
>gi|1036819395|ref|XP_017137558.1| PREDICTED: polycomb protein Su(z)12 isoform X1 [Drosophila r
>gi|1036819414|ref|XP_017137559.1| PREDICTED: polycomb protein Su(z)12 isoform X2 [Drosophila r
>gi|1036820778|ref|XP_017137628.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 2 is
>gi|1036820759|ref|XP_017137627.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 2 is
>gi|1036829903|ref|XP_017138142.1| PREDICTED: exosome complex component MTR3 [Drosophila miranc
>gi|1036829882|ref|XP_017138141.1| PREDICTED: U4/U6 small nuclear ribonucleoprotein Prp3 [Droso
>gi|1036770726|ref|XP_017134810.1| PREDICTED: chromodomain-helicase-DNA-binding protein Mi-2 hc
>gi|1036770747|ref|XP_017134811.1| PREDICTED: chromodomain-helicase-DNA-binding protein Mi-2 hc
>gi|1036770822|ref|XP_017134816.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770840|ref|XP_017134817.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770865|ref|XP_017134818.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770882|ref|XP_017134819.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770785|ref|XP_017134813.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 32 isoform
>gi|1036770766|ref|XP_017134812.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 32 isoform
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>gi|1036770804|ref|XP_017134814.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 32 isoform
>gi|1036792369|ref|XP_017136056.1| PREDICTED: uncharacterized protein LOC108151745 [Drosophila
>gi|1036792219|ref|XP_017136047.1| PREDICTED: protein suppressor of underreplication [Drosophil
>gi|1036792328|ref|XP_017136054.1| PREDICTED: uncharacterized protein LOC108151743 [Drosophila
>gi|1036792348|ref|XP_017136055.1| PREDICTED: uncharacterized protein LOC108151744 [Drosophila
>gi|1036792235|ref|XP_017136049.1| PREDICTED: succinyl-CoA:3-ketoacid coenzyme A transferase 1
>gi|1036792290|ref|XP_017136052.1| PREDICTED: uncharacterized protein LOC108151742 isoform X3
>gi|1036792273|ref|XP_017136051.1| PREDICTED: uncharacterized protein LOC108151742 isoform X2
>gi|1036792254|ref|XP_017136050.1| PREDICTED: uncharacterized protein LOC108151742 isoform X1
>gi|1036792309|ref|XP_017136053.1| PREDICTED: uncharacterized protein LOC108151742 isoform X4
>gi|1036852052|ref|XP_017139387.1| PREDICTED: proton-coupled amino acid transporter 4 [Drosophi
>gi|1036759995|ref|XP_017156869.1| PREDICTED: uncharacterized protein LOC108165347 [Drosophila
>gi|1036766366|ref|XP_017134569.1| PREDICTED: uncharacterized protein LOC108150781 [Drosophila
>gi|1036760013|ref|XP_017156870.1| PREDICTED: uncharacterized protein LOC108165348 [Drosophila
>gi|1036772467|ref|XP_017134910.1| PREDICTED: LOW QUALITY PROTEIN: titin-like [Drosophila miran
>gi|1036772444|ref|XP_017134909.1| PREDICTED: uncharacterized protein DDB_G0285291-like [Drosop
>gi|1036842460|ref|XP_017138846.1| PREDICTED: SEC14 domain and spectrin repeat-containing prote
>gi|1036767013|ref|XP_017134607.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036847199|ref|XP_017139101.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036849598|ref|XP_017139239.1| PREDICTED: uncharacterized protein LOC108153646 [Drosophila
>gi|1036848107|ref|XP_017139151.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036763227|ref|XP_017157055.1| PREDICTED: uncharacterized protein LOC108165509 [Drosophila
>gi|1036814604|ref|XP_017137290.1| PREDICTED: uncharacterized protein LOC108152447 [Drosophila
>gi|1036814407|ref|XP_017137279.1| PREDICTED: intraflagellar transport protein 172 homolog iso
>gi|1036814426|ref|XP_017137280.1| PREDICTED: intraflagellar transport protein 172 homolog iso
>gi|1036814622|ref|XP_017137291.1| PREDICTED: regulation of nuclear pre-mRNA domain-containing
>gi|1036814446|ref|XP_017137281.1| PREDICTED: adenylate cyclase type 8 isoform X1 [Drosophila
>gi|1036814465|ref|XP_017137282.1| PREDICTED: adenylate cyclase type 8 isoform X1 [Drosophila
>gi|1036814485|ref|XP_017137283.1| PREDICTED: adenylate cyclase type 8 isoform X2 [Drosophila
>gi|1036814504|ref|XP_017137284.1| PREDICTED: adenylate cyclase type 8 isoform X3 [Drosophila
>gi|1036814546|ref|XP_017137286.1| PREDICTED: adenylate cyclase type 8 isoform X5 [Drosophila
>gi|1036814583|ref|XP_017137288.1| PREDICTED: adenylate cyclase type 8 isoform X7 [Drosophila
>gi|1036814564|ref|XP_017137287.1| PREDICTED: adenylate cyclase type 8 isoform X6 [Drosophila
>gi|1036814527|ref|XP_017137285.1| PREDICTED: adenylate cyclase type 2 isoform X4 [Drosophila
>gi|1036786354|ref|XP_017135716.1| PREDICTED: adenylate cyclase type 2-like isoform X1 [Drosopl
>gi|1036786373|ref|XP_017135717.1| PREDICTED: adenylate cyclase type 8-like isoform X2 [Drosopl
>gi|1036786507|ref|XP_017135724.1| PREDICTED: cytochrome b reductase 1 isoform X3 [Drosophila
>gi|1036786430|ref|XP_017135720.1| PREDICTED: cytochrome b reductase 1 isoform X1 [Drosophila
>gi|1036786449|ref|XP_017135721.1| PREDICTED: cytochrome b reductase 1 isoform X2 [Drosophila
>gi|1036786470|ref|XP_017135722.1| PREDICTED: cytochrome b reductase 1 isoform X2 [Drosophila
>gi|1036786489|ref|XP_017135723.1| PREDICTED: cytochrome b reductase 1 isoform X2 [Drosophila
>gi|1036786526|ref|XP_017135725.1| PREDICTED: cytochrome b reductase 1 isoform X4 [Drosophila
>gi|1036786545|ref|XP_017135727.1| PREDICTED: uncharacterized protein LOC108151563 [Drosophila
>gi|1036786392|ref|XP_017135718.1| PREDICTED: xylosyltransferase oxt [Drosophila miranda]
>gi|1036786564|ref|XP_017135728.1| PREDICTED: 5'-3' exoribonuclease 2 homolog [Drosophila mira
>gi|1036786411|ref|XP_017135719.1| PREDICTED: protein ecdysoneless [Drosophila miranda]
>gi|1036835407|ref|XP_017138448.1| PREDICTED: uncharacterized protein LOC108153156 [Drosophila
>gi|1036835388|ref|XP_017138447.1| PREDICTED: peritrophin-48-like [Drosophila miranda]
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>gi|1036835332|ref|XP_017138443.1| PREDICTED: major royal jelly protein 1 [Drosophila miranda]
>gi|1036835350|ref|XP_017138444.1| PREDICTED: major royal jelly protein 1 [Drosophila miranda]
>gi|1036835313|ref|XP_017138442.1| PREDICTED: sex peptide receptor [Drosophila miranda]
>gi|1036835294|ref|XP_017138441.1| PREDICTED: uncharacterized protein LOC108153150 [Drosophila
>gi|1036835423|ref|XP_017138449.1| PREDICTED: rab-like protein 3 [Drosophila miranda]
>gi|1036835369|ref|XP_017138446.1| PREDICTED: choline-phosphate cytidylyltransferase A-like [D:
>gi|1036763770|ref|XP_017157085.1| PREDICTED: E3 ubiquitin-protein ligase MARCH6 [Drosophila mi
>gi|1036763985|ref|XP_017157098.1| PREDICTED: thioredoxin, mitochondrial [Drosophila miranda]
>gi|1036763967|ref|XP_017157097.1| PREDICTED: 39S ribosomal protein L23, mitochondrial [Drosoph
>gi|1036763788|ref|XP_017157086.1| PREDICTED: nucleolar MIF4G domain-containing protein 1 homol
>gi|1036763842|ref|XP_017157089.1| PREDICTED: uncharacterized protein C7orf50 homolog [Drosophi
>gi|1036763494|ref|XP_017157069.1| PREDICTED: uncharacterized protein LOC108165524 isoform X1
>gi|1036763512|ref|XP_017157070.1| PREDICTED: uncharacterized protein LOC108165524 isoform X2
>gi|1036763552|ref|XP_017157072.1| PREDICTED: uncharacterized protein LOC108165524 isoform X4
>gi|1036763533|ref|XP_017157071.1| PREDICTED: uncharacterized protein LOC108165524 isoform X3
>gi|1036763682|ref|XP_017157079.1| PREDICTED: uncharacterized protein LOC108165524 isoform X10
>gi|1036763860|ref|XP_017157090.1| PREDICTED: surface protein [Drosophila miranda]
>gi|1036763931|ref|XP_017157094.1| PREDICTED: uncharacterized protein LOC108165533 isoform X2
>gi|1036763949|ref|XP_017157095.1| PREDICTED: uncharacterized protein LOC108165533 isoform X3
>gi|1036763895|ref|XP_017157092.1| PREDICTED: uncharacterized protein LOC108165533 isoform X1
>gi|1036763913|ref|XP_017157093.1| PREDICTED: uncharacterized protein LOC108165533 isoform X1
>gi|1036763611|ref|XP_017157075.1| PREDICTED: uncharacterized protein LOC108165524 isoform X6
>gi|1036763586|ref|XP_017157073.1| PREDICTED: uncharacterized protein LOC108165524 isoform X5
>gi|1036763700|ref|XP_017157080.1| PREDICTED: microtubule-associated protein futsch isoform X1
>gi|1036763736|ref|XP_017157082.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036763718|ref|XP_017157081.1| PREDICTED: mucin-5AC isoform X12 [Drosophila miranda]
>gi|1036763630|ref|XP_017157076.1| PREDICTED: supervillin isoform X7 [Drosophila miranda]
>gi|1036763754|ref|XP_017157083.1| PREDICTED: AF4/FMR2 family member 4 isoform X14 [Drosophila
>gi|1036763877|ref|XP_017157091.1| PREDICTED: uncharacterized protein LOC108165531 [Drosophila
>gi|1036763806|ref|XP_017157087.1| PREDICTED: probable cytochrome P450 4d20 [Drosophila miranda
>gi|1036763646|ref|XP_017157077.1| PREDICTED: supervillin isoform X8 [Drosophila miranda]
>gi|1036763824|ref|XP_017157088.1| PREDICTED: uncharacterized protein LOC108165528 [Drosophila
>gi|1036763664|ref|XP_017157078.1| PREDICTED: supervillin isoform X9 [Drosophila miranda]
>gi|1036832665|ref|XP_017138300.1| PREDICTED: purine nucleoside phosphorylase isoform X3 [Dros
>gi|1036832646|ref|XP_017138299.1| PREDICTED: purine nucleoside phosphorylase isoform X2 [Dros
>gi|1036832627|ref|XP_017138298.1| PREDICTED: purine nucleoside phosphorylase isoform X1 [Dros
>gi|1036832701|ref|XP_017138302.1| PREDICTED: purine nucleoside phosphorylase isoform X5 [Dros
>gi|1036832684|ref|XP_017138301.1| PREDICTED: purine nucleoside phosphorylase isoform X4 [Dros
>gi|1036789463|ref|XP_017135890.1| PREDICTED: AF4/FMR2 family member 4 isoform X18 [Drosophila
>gi|1036789294|ref|XP_017135880.1| PREDICTED: uncharacterized protein LOC108151662 isoform X9
>gi|1036789201|ref|XP_017135875.1| PREDICTED: uncharacterized protein LOC108151662 isoform X4
>gi|1036789387|ref|XP_017135885.1| PREDICTED: AF4/FMR2 family member 4 isoform X14 [Drosophila
>gi|1036789349|ref|XP_017135883.1| PREDICTED: AF4/FMR2 family member 4 isoform X12 [Drosophila
>gi|1036789259|ref|XP_017135878.1| PREDICTED: uncharacterized protein LOC108151662 isoform X7
>gi|1036789165|ref|XP_017135873.1| PREDICTED: uncharacterized protein LOC108151662 isoform X2
>gi|1036789311|ref|XP_017135881.1| PREDICTED: uncharacterized protein LOC108151662 isoform X10
>gi|1036789276|ref|XP_017135879.1| PREDICTED: uncharacterized protein LOC108151662 isoform X8
>gi|1036789368|ref|XP_017135884.1| PREDICTED: AF4/FMR2 family member 4 isoform X13 [Drosophila
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>gi|1036789330|ref|XP_017135882.1| PREDICTED: AF4/FMR2 family member 4 isoform X11 [Drosophila
>gi|1036789183|ref|XP_017135874.1| PREDICTED: uncharacterized protein LOC108151662 isoform X3
>gi|1036789220|ref|XP_017135876.1| PREDICTED: uncharacterized protein LOC108151662 isoform X5
>gi|1036789144|ref|XP_017135872.1| PREDICTED: uncharacterized protein LOC108151662 isoform X1
>gi|1036789238|ref|XP_017135877.1| PREDICTED: uncharacterized protein LOC108151662 isoform X6
>gi|1036789406|ref|XP_017135886.1| PREDICTED: uncharacterized protein LOC108151662 isoform X15
>gi|1036789425|ref|XP_017135887.1| PREDICTED: uncharacterized protein LOC108151662 isoform X16
>gi|1036789482|ref|XP_017135891.1| PREDICTED: uncharacterized protein LOC108151662 isoform X19
>gi|1036789444|ref|XP_017135888.1| PREDICTED: AF4/FMR2 family member 4 isoform X17 [Drosophila
>gi|1036809813|ref|XP_017137030.1| PREDICTED: altered inheritance of mitochondria protein 3 iso
>gi|1036809788|ref|XP_017137029.1| PREDICTED: extensin isoform X1 [Drosophila miranda]
>gi|1036809943|ref|XP_017137036.1| PREDICTED: O-acetyl-ADP-ribose deacetylase 1 [Drosophila mir
>gi|1036809922|ref|XP_017137035.1| PREDICTED: O-acetyl-ADP-ribose deacetylase 1 [Drosophila mir
>gi|1036809903|ref|XP_017137033.1| PREDICTED: uncharacterized protein LOC108152312 [Drosophila
>gi|1036809884|ref|XP_017137032.1| PREDICTED: uncharacterized oxidoreductase YjmC [Drosophila m
>gi|1036809866|ref|XP_017137031.1| PREDICTED: uncharacterized protein LOC108152310 [Drosophila
>gi|1036809695|ref|XP_017137024.1| PREDICTED: uncharacterized protein LOC108152307 isoform X1
>gi|1036809751|ref|XP_017137027.1| PREDICTED: uncharacterized protein LOC108152307 isoform X4
>gi|1036809714|ref|XP_017137025.1| PREDICTED: uncharacterized protein LOC108152307 isoform X2
>gi|1036809769|ref|XP_017137028.1| PREDICTED: uncharacterized protein LOC108152307 isoform X5
>gi|1036809733|ref|XP_017137026.1| PREDICTED: GTPase-activating protein BEM2/IPL2 isoform X3 [D
>gi|1036809959|ref|XP_017137037.1| PREDICTED: multifunctional methyltransferase subunit TRM112-
>gi|1036836316|ref|XP_017138502.1| PREDICTED: integrin-linked protein kinase homolog pat-4 [Dro
>gi|1036841245|ref|XP_017138788.1| PREDICTED: ecdysone-induced protein 78C isoform X1 [Drosophi
>gi|1036841264|ref|XP_017138789.1| PREDICTED: ecdysone-induced protein 78C isoform X2 [Drosophi
>gi|1036841300|ref|XP_017138792.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036841282|ref|XP_017138790.1| PREDICTED: ecdysone-induced protein 78C isoform X3 [Drosophi
>gi|1036844209|ref|XP_017138947.1| PREDICTED: inositol monophosphatase 1 [Drosophila miranda]
>gi|1036849525|ref|XP_017139234.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036789773|ref|XP_017135909.1| PREDICTED: acetyl-coenzyme A synthetase [Drosophila miranda]
>gi|1036789813|ref|XP_017135912.1| PREDICTED: glycine cleavage system H protein, mitochondrial
>gi|1036789794|ref|XP_017135910.1| PREDICTED: uncharacterized protein LOC108151679 [Drosophila
>gi|1036789754|ref|XP_017135908.1| PREDICTED: uncharacterized protein LOC108151677 [Drosophila
>gi|1036767612|ref|XP_017134639.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036767577|ref|XP_017134637.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036808488|ref|XP_017136959.1| PREDICTED: pupal cuticle protein Edg-78E-like [Drosophila mi
>gi|1036808450|ref|XP_017136957.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036808469|ref|XP_017136958.1| PREDICTED: pupal cuticle protein Edg-78E-like [Drosophila mi
>gi|1036808414|ref|XP_017136955.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036808432|ref|XP_017136956.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036808396|ref|XP_017136954.1| PREDICTED: polycomb group protein Pc [Drosophila miranda]
>gi|1036847834|ref|XP_017139136.1| PREDICTED: ras-related protein Rab-26 isoform X2 [Drosophila
>gi|1036847853|ref|XP_017139137.1| PREDICTED: ras-related protein Rab-26 isoform X3 [Drosophila
>gi|1036847817|ref|XP_017139135.1| PREDICTED: ras-related protein Rab-26 isoform X1 [Drosophila
>gi|1036804826|ref|XP_017136757.1| PREDICTED: pre-rRNA-processing protein TSR1 homolog [Drosoph
>gi|1036804807|ref|XP_017136756.1| PREDICTED: TBC1 domain family member 9 isoform X2 [Drosophi
>gi|1036804789|ref|XP_017136755.1| PREDICTED: TBC1 domain family member 9 isoform X1 [Drosophi
>gi|1036838677|ref|XP_017138647.1| PREDICTED: uncharacterized protein LOC108153286 [Drosophila
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>gi|1036838696|ref|XP_017138648.1| PREDICTED: uncharacterized protein LOC108153287 [Drosophila
>gi|1036838712|ref|XP_017138649.1| PREDICTED: uncharacterized protein LOC108153287 [Drosophila
>gi|1036846881|ref|XP_017139083.1| PREDICTED: nuclear transcription factor Y subunit beta [Dros
>gi|1036846900|ref|XP_017139084.1| PREDICTED: uncharacterized protein DDB_G0283697 isoform X1
>gi|1036846919|ref|XP_017139085.1| PREDICTED: uncharacterized protein LOC108153533 isoform X2
>gi|1036760031|ref|XP_017156871.1| PREDICTED: uncharacterized protein LOC108165349, partial [Di
>gi|1036815665|ref|XP_017137348.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1036815684|ref|XP_017137349.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1036815703|ref|XP_017137350.1| PREDICTED: trichohyalin isoform X2 [Drosophila miranda]
>gi|1036815724|ref|XP_017137351.1| PREDICTED: HIV Tat-specific factor 1 [Drosophila miranda]
>gi|1036815743|ref|XP_017137352.1| PREDICTED: uncharacterized protein LOC108152488 [Drosophila
>gi|1036780912|ref|XP_017135400.1| PREDICTED: adult enhancer factor 1 isoform X1 [Drosophila m:
>gi|1036780930|ref|XP_017135402.1| PREDICTED: adult enhancer factor 1 isoform X2 [Drosophila m:
>gi|1036780879|ref|XP_017135398.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036781019|ref|XP_017135408.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1036760050|ref|XP_017156872.1| PREDICTED: cuticle protein 8 [Drosophila miranda]
>gi|1036780983|ref|XP_017135406.1| PREDICTED: uncharacterized protein LOC108151358 [Drosophila
>gi|1036781037|ref|XP_017135409.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036781001|ref|XP_017135407.1| PREDICTED: peroxisome biogenesis factor 2 [Drosophila mirano
>gi|1036780964|ref|XP_017135404.1| PREDICTED: UPF0183 protein CG7083 [Drosophila miranda]
>gi|1036780948|ref|XP_017135403.1| PREDICTED: DNA primase small subunit [Drosophila miranda]
>gi|1036780896|ref|XP_017135399.1| PREDICTED: heat shock 70 kDa protein 14 [Drosophila miranda]
>gi|1036799591|ref|XP_017136462.1| PREDICTED: rab GTPase-activating protein 1-like isoform X1
>gi|1036799612|ref|XP_017136463.1| PREDICTED: rab GTPase-activating protein 1-like isoform X2
>gi|1036799630|ref|XP_017136464.1| PREDICTED: uncharacterized protein LOC108152004 isoform X1
>gi|1036799649|ref|XP_017136465.1| PREDICTED: uncharacterized protein LOC108152004 isoform X2
>gi|1036799726|ref|XP_017136469.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799708|ref|XP_017136468.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799687|ref|XP_017136467.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799668|ref|XP_017136466.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799744|ref|XP_017136470.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799762|ref|XP_017136471.1| PREDICTED: fibronectin-binding protein A-like [Drosophila m:
>gi|1036843726|ref|XP_017138921.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036843743|ref|XP_017138922.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036843762|ref|XP_017138924.1| PREDICTED: uncharacterized protein LOC108153430 isoform X2
>gi|1036763458|ref|XP_017157067.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036821930|ref|XP_017137692.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036821949|ref|XP_017137693.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036821968|ref|XP_017137694.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036821987|ref|XP_017137695.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036822006|ref|XP_017137696.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036822023|ref|XP_017137697.1| PREDICTED: uncharacterized protein LOC108152696 isoform X2
>gi|1036822042|ref|XP_017137698.1| PREDICTED: uncharacterized protein LOC108152696 isoform X3
>gi|1036822061|ref|XP_017137699.1| PREDICTED: uncharacterized protein LOC108152698 [Drosophila
>gi|1036795692|ref|XP_017136243.1| PREDICTED: NADPH-dependent diflavin oxidoreductase 1 [Drosop
>gi|1036795619|ref|XP_017136238.1| PREDICTED: intraflagellar transport protein 122 homolog [Dro
>gi|1036795637|ref|XP_017136239.1| PREDICTED: exocyst complex component 7 [Drosophila miranda]
>gi|1036795656|ref|XP_017136240.1| PREDICTED: exocyst complex component 7 [Drosophila miranda]
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>gi|1036795711|ref|XP_017136244.1| PREDICTED: protein matrimony [Drosophila miranda]
>gi|1036795673|ref|XP_017136242.1| PREDICTED: agrin [Drosophila miranda]
>gi|1036821777|ref|XP_017137683.1| PREDICTED: E3 ubiquitin-protein ligase CBL-B-B [Drosophila m
>gi|1036818077|ref|XP_017137486.1| PREDICTED: cold shock domain-containing protein E1 isoform X
>gi|1036818060|ref|XP_017137485.1| PREDICTED: cold shock domain-containing protein E1 isoform X
>gi|1036804608|ref|XP_017136745.1| PREDICTED: LOW QUALITY PROTEIN: AFG3-like protein 2 [Drosoph
>gi|1036804645|ref|XP_017136747.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804662|ref|XP_017136748.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804699|ref|XP_017136751.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804626|ref|XP_017136746.1| PREDICTED: hormone receptor 4 isoform X1 [Drosophila miranda
>gi|1036804681|ref|XP_017136750.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804733|ref|XP_017136752.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036764499|ref|XP_017157120.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-
>gi|1036764517|ref|XP_017157121.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-
>gi|1036764543|ref|XP_017157122.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-
>gi|1036764563|ref|XP_017157123.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-
>gi|1036830757|ref|XP_017138195.1| PREDICTED: protein rogdi [Drosophila miranda]
>gi|1036830776|ref|XP_017138196.1| PREDICTED: troponin C [Drosophila miranda]
>gi|1036805538|ref|XP_017136798.1| PREDICTED: uncharacterized protein LOC108152178 [Drosophila
>gi|1036805500|ref|XP_017136796.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805479|ref|XP_017136795.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805460|ref|XP_017136793.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805439|ref|XP_017136792.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805421|ref|XP_017136791.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805386|ref|XP_017136789.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805403|ref|XP_017136790.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805367|ref|XP_017136788.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805519|ref|XP_017136797.1| PREDICTED: uncharacterized protein LOC108152177 [Drosophila
>gi|1036760069|ref|XP_017156873.1| PREDICTED: uncharacterized protein LOC108165352 [Drosophila
>gi|1036781772|ref|XP_017135452.1| PREDICTED: putative RNA-binding protein Luc7-like 1 [Drosoph
>gi|1036781754|ref|XP_017135450.1| PREDICTED: enhancer of mRNA-decapping protein 3 [Drosophila
>gi|1036781519|ref|XP_017135437.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X1 [Dr
>gi|1036781590|ref|XP_017135441.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X5 [Dr
>gi|1036781573|ref|XP_017135440.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X4 [Dr
>gi|1036781645|ref|XP_017135444.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X8 [Dr
>gi|1036781681|ref|XP_017135446.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X10 [D
>gi|1036781538|ref|XP_017135438.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X2 [Dr
>gi|1036781737|ref|XP_017135449.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X13 [D
>gi|1036781664|ref|XP_017135445.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X9 [Dr
>gi|1036781554|ref|XP_017135439.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X3 [Dr
>gi|1036781607|ref|XP_017135442.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X6 [Dr
>gi|1036781626|ref|XP_017135443.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X7 [Dr
>gi|1036781699|ref|XP_017135447.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X11 [D
>gi|1036781717|ref|XP_017135448.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X12 [D
>gi|1036823803|ref|XP_017137798.1| PREDICTED: uncharacterized protein LOC108152751 [Drosophila
>gi|1036823761|ref|XP_017137796.1| PREDICTED: insulin-degrading enzyme [Drosophila miranda]
>gi|1036823780|ref|XP_017137797.1| PREDICTED: insulin-degrading enzyme [Drosophila miranda]
>gi|1036820849|ref|XP_017137633.1| PREDICTED: rho-related BTB domain-containing protein 1 isofo
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>gi|1036820887|ref|XP_017137635.1| PREDICTED: rho-related BTB domain-containing protein 1 isofo
>gi|1036820868|ref|XP_017137634.1| PREDICTED: rho-related BTB domain-containing protein 1 isofo
>gi|1036820905|ref|XP_017137636.1| PREDICTED: charged multivesicular body protein 7 [Drosophila
>gi|1036814073|ref|XP_017137259.1| PREDICTED: pantothenate kinase 3 isoform X1 [Drosophila mira
>gi|1036814091|ref|XP_017137260.1| PREDICTED: pantothenate kinase 3 isoform X2 [Drosophila mira
>gi|1036814138|ref|XP_017137263.1| PREDICTED: pantothenate kinase 3 isoform X4 [Drosophila mira
>gi|1036814176|ref|XP_017137265.1| PREDICTED: pantothenate kinase 3 isoform X6 [Drosophila mira
>gi|1036814117|ref|XP_017137262.1| PREDICTED: pantothenate kinase 3 isoform X3 [Drosophila mira
>gi|1036814157|ref|XP_017137264.1| PREDICTED: pantothenate kinase 3 isoform X5 [Drosophila mira
>gi|1036814194|ref|XP_017137266.1| PREDICTED: F-box/WD repeat-containing protein 4 [Drosophila
>gi|1036827193|ref|XP_017137988.1| PREDICTED: CBL-interacting serine/threonine-protein kinase
>gi|1036847567|ref|XP_017139121.1| PREDICTED: cationic amino acid transporter 4 [Drosophila mi
>gi|1036843599|ref|XP_017138914.1| PREDICTED: uncharacterized protein LOC108153422 [Drosophila
>gi|1036843618|ref|XP_017138915.1| PREDICTED: dipeptidase 3 [Drosophila miranda]
>gi|1036838554|ref|XP_017138636.1| PREDICTED: uncharacterized protein C7orf26 homolog [Drosophi
>gi|1036838161|ref|XP_017138613.1| PREDICTED: transmembrane protein 104 homolog [Drosophila mi
>gi|1036781297|ref|XP_017135424.1| PREDICTED: lipoyl synthase, mitochondrial [Drosophila miran
>gi|1036781351|ref|XP_017135427.1| PREDICTED: neuronal membrane glycoprotein M6-a isoform X1 [
>gi|1036781369|ref|XP_017135428.1| PREDICTED: neuronal membrane glycoprotein M6-a isoform X2 [
>gi|1036781335|ref|XP_017135426.1| PREDICTED: serine/threonine-protein kinase PLK4 [Drosophila
>gi|1036781316|ref|XP_017135425.1| PREDICTED: cyclin-dependent kinase 12 [Drosophila miranda]
>gi|1036808324|ref|XP_017136950.1| PREDICTED: WD repeat-containing protein 26 homolog [Drosophi
>gi|1036808342|ref|XP_017136951.1| PREDICTED: WD repeat-containing protein 26 homolog [Drosophi
>gi|1036808360|ref|XP_017136952.1| PREDICTED: WD repeat-containing protein 26 homolog [Drosophi
>gi|1036808378|ref|XP_017136953.1| PREDICTED: protein ARV1 [Drosophila miranda]
>gi|1036827172|ref|XP_017137987.1| PREDICTED: probable E3 ubiquitin-protein ligase makorin-1 [
>gi|1036767050|ref|XP_017134610.1| PREDICTED: pickpocket protein 28 isoform X1 [Drosophila mira
>gi|1036767071|ref|XP_017134611.1| PREDICTED: pickpocket protein 28 isoform X2 [Drosophila mira
>gi|1036833247|ref|XP_017138334.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 4 [
>gi|1036833265|ref|XP_017138335.1| PREDICTED: uncharacterized protein LOC108153066 [Drosophila
>gi|1036847420|ref|XP_017139115.1| PREDICTED: V-type proton ATPase subunit e 2 [Drosophila mira
>gi|1036802904|ref|XP_017136648.1| PREDICTED: serine/threonine-protein kinase WNK1 [Drosophila
>gi|1036802885|ref|XP_017136647.1| PREDICTED: N-acylneuraminate cytidylyltransferase [Drosophil
>gi|1036830998|ref|XP_017138208.1| PREDICTED: neuronal calcium sensor 2 isoform X2 [Drosophila
>gi|1036830981|ref|XP_017138207.1| PREDICTED: neuronal calcium sensor 2 isoform X1 [Drosophila
>gi|1036830962|ref|XP_017138206.1| PREDICTED: neurocalcin homolog [Drosophila miranda]
>gi|1036840070|ref|XP_017138721.1| PREDICTED: 4-aminobutyrate aminotransferase, mitochondrial
>gi|1036840388|ref|XP_017138738.1| PREDICTED: mitochondrial import receptor subunit TOM20 homol
>gi|1036775759|ref|XP_017135093.1| PREDICTED: fat-like cadherin-related tumor suppressor homolo
>gi|1036775778|ref|XP_017135094.1| PREDICTED: uncharacterized protein LOC108151174 [Drosophila
>gi|1036816945|ref|XP_017137423.1| PREDICTED: GDP-Man:Man(3)GlcNAc(2)-PP-Dol alpha-1,2-mannosyl
>gi|1036816964|ref|XP_017137424.1| PREDICTED: uncharacterized protein LOC108152530 [Drosophila
>gi|1036816844|ref|XP_017137417.1| PREDICTED: IQ and AAA domain-containing protein 1-like isofo
>gi|1036816863|ref|XP_017137418.1| PREDICTED: IQ and AAA domain-containing protein 1-like isofo
>gi|1036816903|ref|XP_017137421.1| PREDICTED: IQ and AAA domain-containing protein 1-like isofo
>gi|1036816882|ref|XP_017137420.1| PREDICTED: IQ and AAA domain-containing protein 1-like isofo
>gi|1036816926|ref|XP_017137422.1| PREDICTED: opioid-binding protein/cell adhesion molecule-lik
>gi|1036835442|ref|XP_017138450.1| PREDICTED: uncharacterized protein LOC108153158 [Drosophila
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>gi|1036831455|ref|XP_017138235.1| PREDICTED: membrane-bound transcription factor site-1 prote
>gi|1036845372|ref|XP_017139005.1| PREDICTED: transcription factor AP-2-epsilon isoform X6 [Dro
>gi|1036845336|ref|XP_017139003.1| PREDICTED: transcription factor AP-2-epsilon isoform X4 [Dro
>gi|1036845317|ref|XP_017139001.1| PREDICTED: transcription factor AP-2-epsilon isoform X3 [Dro
>gi|1036845280|ref|XP_017138999.1| PREDICTED: transcription factor AP-2-epsilon isoform X1 [Dro
>gi|1036845355|ref|XP_017139004.1| PREDICTED: transcription factor AP-2-epsilon isoform X5 [Dro
>gi|1036845298|ref|XP_017139000.1| PREDICTED: transcription factor AP-2-epsilon isoform X2 [Dro
>gi|1036841786|ref|XP_017138821.1| PREDICTED: ER membrane protein complex subunit 10 [Drosophil
>gi|1036814985|ref|XP_017137310.1| PREDICTED: uncharacterized protein LOC108152457 [Drosophila
>gi|1036815006|ref|XP_017137311.1| PREDICTED: alanine--glyoxylate aminotransferase 2, mitochond
>gi|1036815027|ref|XP_017137312.1| PREDICTED: alanine--glyoxylate aminotransferase 2, mitochond
>gi|1036815048|ref|XP_017137313.1| PREDICTED: transmembrane protein 64 [Drosophila miranda]
>gi|1036844266|ref|XP_017138950.1| PREDICTED: uncharacterized protein LOC108153445 isoform X1
>gi|1036844285|ref|XP_017138951.1| PREDICTED: uncharacterized protein LOC108153445 isoform X1
>gi|1036844304|ref|XP_017138952.1| PREDICTED: uncharacterized protein LOC108153445 isoform X2
>gi|1036834508|ref|XP_017138394.1| PREDICTED: transcription elongation factor, mitochondrial [
>gi|1036834527|ref|XP_017138395.1| PREDICTED: protein maelstrom 1 [Drosophila miranda]
>gi|1036817710|ref|XP_017137465.1| PREDICTED: serine/arginine repetitive matrix protein 1 [Dros
>gi|1036817674|ref|XP_017137463.1| PREDICTED: ADP-ribosylation factor GTPase-activating protei
>gi|1036817691|ref|XP_017137464.1| PREDICTED: ADP-ribosylation factor GTPase-activating protei
>gi|1036817782|ref|XP_017137469.1| PREDICTED: translation initiation factor IF-2 isoform X4 [D
>gi|1036817763|ref|XP_017137468.1| PREDICTED: salt stress root protein RS1 isoform X3 [Drosoph
>gi|1036817746|ref|XP_017137467.1| PREDICTED: translation initiation factor IF-2 isoform X2 [D
>gi|1036817729|ref|XP_017137466.1| PREDICTED: salt stress root protein RS1 isoform X1 [Drosoph
>gi|1036760087|ref|XP_017156874.1| PREDICTED: cell wall protein TIR3-like [Drosophila miranda]
>gi|1036769608|ref|XP_017134749.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036760104|ref|XP_017156877.1| PREDICTED: protein new-glue 2-like [Drosophila miranda]
>gi|1036768041|ref|XP_017134660.1| PREDICTED: uncharacterized protein LOC108150868 [Drosophila
>gi|1036760124|ref|XP_017156878.1| PREDICTED: uncharacterized protein LOC108165355 [Drosophila
>gi|1036760145|ref|XP_017156879.1| PREDICTED: circadian clock-controlled protein [Drosophila m
>gi|1036839884|ref|XP_017138710.1| PREDICTED: ribonuclease P protein subunit p29-like [Drosoph
>gi|1036839903|ref|XP_017138711.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosop
>gi|1036839922|ref|XP_017138712.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosop
>gi|1036839943|ref|XP_017138713.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosop
>gi|1036763368|ref|XP_017157062.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosop
>gi|1036770903|ref|XP_017134820.1| PREDICTED: uncharacterized protein LOC108150985 [Drosophila
>gi|1036760162|ref|XP_017156880.1| PREDICTED: uncharacterized protein LOC108165357 [Drosophila
>gi|1036760180|ref|XP_017156881.1| PREDICTED: uncharacterized protein LOC108165358 [Drosophila
>gi|1036760200|ref|XP_017156882.1| PREDICTED: uncharacterized protein LOC108165359 [Drosophila
>gi|1036770922|ref|XP_017134821.1| PREDICTED: uncharacterized protein LOC108150986 [Drosophila
>gi|1036760218|ref|XP_017156884.1| PREDICTED: uncharacterized protein LOC108165360 [Drosophila
>gi|1036760237|ref|XP_017156885.1| PREDICTED: uncharacterized protein LOC108165361 [Drosophila
>gi|1036760256|ref|XP_017156886.1| PREDICTED: uncharacterized protein LOC108165362 [Drosophila
>gi|1036760275|ref|XP_017156887.1| PREDICTED: uncharacterized protein LOC108165363 [Drosophila
>gi|1036760296|ref|XP_017156888.1| PREDICTED: uncharacterized protein LOC108165364 [Drosophila
>gi|1036760315|ref|XP_017156889.1| PREDICTED: putative uncharacterized protein DDB_G0282129 [D
>gi|1036852232|ref|XP_017139398.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036850112|ref|XP_017139271.1| PREDICTED: thioredoxin reductase 2, mitochondrial [Drosophil
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>gi|1036766462|ref|XP_017134576.1| PREDICTED: uncharacterized protein LOC108150786 [Drosophila
>gi|1036760334|ref|XP_017156890.1| PREDICTED: uncharacterized protein LOC108165366 [Drosophila
>gi|1036760376|ref|XP_017156892.1| PREDICTED: uncharacterized protein LOC108165367 [Drosophila
>gi|1036852195|ref|XP_017139396.1| PREDICTED: uncharacterized protein LOC108153768 [Drosophila
>gi|1036852214|ref|XP_017139397.1| PREDICTED: uncharacterized protein LOC108153770 [Drosophila
>gi|1036766504|ref|XP_017134578.1| PREDICTED: uncharacterized protein LOC108150788 [Drosophila
>gi|1036760395|ref|XP_017156893.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036851018|ref|XP_017139322.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840279|ref|XP_017138733.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840335|ref|XP_017138736.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840314|ref|XP_017138735.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840296|ref|XP_017138734.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840261|ref|XP_017138732.1| PREDICTED: cilia- and flagella-associated protein 45 [Droso
>gi|1036773303|ref|XP_017134957.1| PREDICTED: teneurin-m isoform X1 [Drosophila miranda]
>gi|1036773322|ref|XP_017134958.1| PREDICTED: teneurin-m isoform X2 [Drosophila miranda]
>gi|1036773341|ref|XP_017134959.1| PREDICTED: teneurin-m isoform X3 [Drosophila miranda]
>gi|1036773360|ref|XP_017134960.1| PREDICTED: teneurin-m isoform X4 [Drosophila miranda]
>gi|1036819799|ref|XP_017137581.1| PREDICTED: isoleucine--tRNA ligase, cytoplasmic [Drosophila
>gi|1036821237|ref|XP_017137654.1| PREDICTED: DDB1- and CUL4-associated factor 8 [Drosophila m
>gi|1036821218|ref|XP_017137652.1| PREDICTED: dynein heavy chain 12, axonemal [Drosophila mira
>gi|1036772576|ref|XP_017134916.1| PREDICTED: ankyrin-1 [Drosophila miranda]
>gi|1036772595|ref|XP_017134917.1| PREDICTED: DNA polymerase delta small subunit [Drosophila m
>gi|1036772637|ref|XP_017134920.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm8 [Drosoph
>gi|1036772616|ref|XP_017134918.1| PREDICTED: 39S ribosomal protein L46, mitochondrial [Drosoph
>gi|1036772538|ref|XP_017134914.1| PREDICTED: spectrin alpha chain isoform X4 [Drosophila mira
>gi|1036772521|ref|XP_017134913.1| PREDICTED: spectrin alpha chain isoform X3 [Drosophila mira
>gi|1036772503|ref|XP_017134912.1| PREDICTED: spectrin alpha chain isoform X2 [Drosophila mira
>gi|1036772486|ref|XP_017134911.1| PREDICTED: spectrin alpha chain isoform X1 [Drosophila mira
>gi|1036772557|ref|XP_017134915.1| PREDICTED: protein disks lost [Drosophila miranda]
>gi|1036836935|ref|XP_017138538.1| PREDICTED: LOW QUALITY PROTEIN: ragulator complex protein LA
>gi|1036836916|ref|XP_017138537.1| PREDICTED: vacuolar fusion protein CCZ1 homolog [Drosophila
>gi|1036844517|ref|XP_017138963.1| PREDICTED: polyubiquitin [Drosophila miranda]
>gi|1036821406|ref|XP_017137662.1| PREDICTED: very-long-chain enoyl-CoA reductase [Drosophila
>gi|1036821385|ref|XP_017137661.1| PREDICTED: anaphase-promoting complex subunit 5 [Drosophila
>gi|1036821425|ref|XP_017137664.1| PREDICTED: mitochondrial import receptor subunit TOM22 homol
>gi|1036839718|ref|XP_017138701.1| PREDICTED: calcium-binding protein E63-1 isoform X2 [Drosoph
>gi|1036839699|ref|XP_017138700.1| PREDICTED: calcium-binding protein E63-1 isoform X1 [Drosoph
>gi|1036839735|ref|XP_017138702.1| PREDICTED: calcium-binding protein E63-1 isoform X3 [Drosoph
>gi|1036839792|ref|XP_017138705.1| PREDICTED: calcium-binding protein E63-1 isoform X6 [Drosoph
>gi|1036839773|ref|XP_017138704.1| PREDICTED: calcium-binding protein E63-1 isoform X5 [Drosoph
>gi|1036839754|ref|XP_017138703.1| PREDICTED: calcium-binding protein E63-1 isoform X4 [Drosoph
>gi|1036845762|ref|XP_017139019.1| PREDICTED: transmembrane protein C5orf28 homolog [Drosophila
>gi|1036843132|ref|XP_017138885.1| PREDICTED: aldose reductase [Drosophila miranda]
>gi|1036843077|ref|XP_017138882.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036843095|ref|XP_017138883.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036843114|ref|XP_017138884.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036770382|ref|XP_017134792.1| PREDICTED: F-box/LRR-repeat protein 17 [Drosophila miranda]
>gi|1036766809|ref|XP_017134595.1| PREDICTED: uncharacterized protein LOC108150802 [Drosophila
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>gi|1036841357|ref|XP_017138795.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841374|ref|XP_017138796.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841393|ref|XP_017138797.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841412|ref|XP_017138798.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841432|ref|XP_017138799.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841451|ref|XP_017138800.1| PREDICTED: zinc finger protein 2 isoform X2 [Drosophila mira
>gi|1036838087|ref|XP_017138609.1| PREDICTED: zinc finger protein rotund [Drosophila miranda]
>gi|1036844972|ref|XP_017138980.1| PREDICTED: MIT domain-containing protein 1 [Drosophila mirar
>gi|1036824637|ref|XP_017137846.1| PREDICTED: fork head domain-containing protein FD2 isoform 2
>gi|1036824658|ref|XP_017137848.1| PREDICTED: fork head domain-containing protein FD2 isoform 2
>gi|1036824677|ref|XP_017137849.1| PREDICTED: mitochondrial ubiquitin ligase activator of NFKB
>gi|1036824617|ref|XP_017137845.1| PREDICTED: FGGY carbohydrate kinase domain-containing prote:
>gi|1036800785|ref|XP_017136530.1| PREDICTED: uncharacterized protein LOC108152042 [Drosophila
>gi|1036800745|ref|XP_017136528.1| PREDICTED: adenosine 3'-phospho 5'-phosphosulfate transporte
>gi|1036800692|ref|XP_017136525.1| PREDICTED: calcium homeostasis endoplasmic reticulum protei
>gi|1036800728|ref|XP_017136527.1| PREDICTED: calcium homeostasis endoplasmic reticulum protei
>gi|1036800710|ref|XP_017136526.1| PREDICTED: calcium homeostasis endoplasmic reticulum protei
>gi|1036800766|ref|XP_017136529.1| PREDICTED: probable malonyl-CoA-acyl carrier protein transac
>gi|1036842385|ref|XP_017138842.1| PREDICTED: uncharacterized protein LOC108153380 [Drosophila
>gi|1036842367|ref|XP_017138841.1| PREDICTED: uncharacterized protein LOC108153379 [Drosophila
>gi|1036795458|ref|XP_017136229.1| PREDICTED: unc-112-related protein [Drosophila miranda]
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>gi|1036795494|ref|XP_017136231.1| PREDICTED: uncharacterized protein LOC108151858 [Drosophila
>gi|1036795439|ref|XP_017136228.1| PREDICTED: ribosome biogenesis protein BMS1 homolog [Drosopl
>gi|1036833429|ref|XP_017138344.1| PREDICTED: coiled-coil domain-containing protein 6 [Drosophi
>gi|1036824982|ref|XP_017137863.1| PREDICTED: acetyl-coenzyme A transporter 1 [Drosophila mirar
>gi|1036825001|ref|XP_017137864.1| PREDICTED: eukaryotic translation initiation factor 3 subun:
>gi|1036811610|ref|XP_017137128.1| PREDICTED: glutamate synthase [NADH], amyloplastic isoform 2
>gi|1036811592|ref|XP_017137126.1| PREDICTED: glutamate synthase [NADH], amyloplastic isoform 2
>gi|1036811629|ref|XP_017137129.1| PREDICTED: glutamate synthase 1 [NADH], chloroplastic isofor
>gi|1036811648|ref|XP_017137130.1| PREDICTED: glutamate synthase 1 [NADH], chloroplastic isofor
>gi|1036811667|ref|XP_017137131.1| PREDICTED: glutamate synthase 1 [NADH], chloroplastic isofor
>gi|1036831019|ref|XP_017138210.1| PREDICTED: nuclear migration protein nudC [Drosophila mirand
>gi|1036831038|ref|XP_017138211.1| PREDICTED: vitellogenin-2 isoform X1 [Drosophila miranda]
>gi|1036831057|ref|XP_017138212.1| PREDICTED: vitellogenin-2 isoform X2 [Drosophila miranda]
>gi|1036831075|ref|XP_017138213.1| PREDICTED: vitellogenin-2 isoform X2 [Drosophila miranda]
>gi|1036796359|ref|XP_017136279.1| PREDICTED: zinc finger protein CG2199 [Drosophila miranda]
>gi|1036796378|ref|XP_017136280.1| PREDICTED: uncharacterized protein LOC108151890 [Drosophila
>gi|1036796397|ref|XP_017136281.1| PREDICTED: uncharacterized protein LOC108151891 [Drosophila
>gi|1036796418|ref|XP_017136282.1| PREDICTED: uncharacterized protein LOC108151892 [Drosophila
>gi|1036796488|ref|XP_017136286.1| PREDICTED: uncharacterized protein LOC108151895 [Drosophila
>gi|1036796456|ref|XP_017136284.1| PREDICTED: acyl carrier protein, mitochondrial isoform X2 [I
>gi|1036796471|ref|XP_017136285.1| PREDICTED: acyl carrier protein, mitochondrial isoform X3 [I
>gi|1036796437|ref|XP_017136283.1| PREDICTED: acyl carrier protein, mitochondrial isoform X1 [I
>gi|1036776981|ref|XP_017135164.1| PREDICTED: myosin-IB isoform X1 [Drosophila miranda]
>gi|1036776998|ref|XP_017135165.1| PREDICTED: myosin-IB isoform X2 [Drosophila miranda]
>gi|1036777069|ref|XP_017135170.1| PREDICTED: annexin A11 isoform X1 [Drosophila miranda]
>gi|1036777087|ref|XP_017135171.1| PREDICTED: annexin A7 isoform X2 [Drosophila miranda]
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>gi|1036776963|ref|XP_017135163.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofo
>gi|1036776945|ref|XP_017135162.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofo
>gi|1036776926|ref|XP_017135161.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofo
>gi|1036776908|ref|XP_017135160.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofo
>gi|1036777050|ref|XP_017135168.1| PREDICTED: lipid droplet-associated hydrolase [Drosophila m
>gi|1036776889|ref|XP_017135159.1| PREDICTED: RNA polymerase-associated protein CTR9 homolog [
>gi|1036777012|ref|XP_017135166.1| PREDICTED: 5'-nucleotidase domain-containing protein 1 [Dros
>gi|1036777031|ref|XP_017135167.1| PREDICTED: acetyl-CoA acetyltransferase, cytosolic [Drosophi
>gi|1036835217|ref|XP_017138436.1| PREDICTED: ras-related protein Rac1 [Drosophila miranda]
>gi|1036841651|ref|XP_017138812.1| PREDICTED: calumenin [Drosophila miranda]
>gi|1036836224|ref|XP_017138497.1| PREDICTED: rab5 GDP/GTP exchange factor [Drosophila miranda]
>gi|1036852560|ref|XP_017139418.1| PREDICTED: E3 ubiquitin-protein ligase msl-2-like [Drosophil
>gi|1036828205|ref|XP_017138047.1| PREDICTED: uncharacterized protein LOC108152907 isoform X2
>gi|1036828148|ref|XP_017138044.1| PREDICTED: uncharacterized protein LOC108152907 isoform X1
>gi|1036828167|ref|XP_017138045.1| PREDICTED: uncharacterized protein LOC108152907 isoform X1
>gi|1036828186|ref|XP_017138046.1| PREDICTED: uncharacterized protein LOC108152907 isoform X1
>gi|1036828224|ref|XP_017138048.1| PREDICTED: uncharacterized protein LOC108152907 isoform X3
>gi|1036850398|ref|XP_017139286.1| PREDICTED: uncharacterized protein LOC108153681 isoform X1
>gi|1036850417|ref|XP_017139288.1| PREDICTED: perlucin isoform X2 [Drosophila miranda]
>gi|1036827133|ref|XP_017137985.1| PREDICTED: uncharacterized protein LOC108152866 [Drosophila
>gi|1036827114|ref|XP_017137983.1| PREDICTED: uncharacterized protein LOC108152865 [Drosophila
>gi|1036827057|ref|XP_017137980.1| PREDICTED: phosphatidylinositide phosphatase SAC1 isoform X
>gi|1036827076|ref|XP_017137981.1| PREDICTED: phosphatidylinositide phosphatase SAC1 isoform X
>gi|1036827095|ref|XP_017137982.1| PREDICTED: phosphatidylinositide phosphatase SAC1 isoform X
>gi|1036807280|ref|XP_017136893.1| PREDICTED: kinesin-like protein Klp61F isoform X2 [Drosophil
>gi|1036807263|ref|XP_017136892.1| PREDICTED: kinesin-like protein Klp61F isoform X1 [Drosophil
>gi|1036807299|ref|XP_017136894.1| PREDICTED: uncharacterized protein LOC108152223 [Drosophila
>gi|1036807317|ref|XP_017136895.1| PREDICTED: uncharacterized protein LOC108152223 [Drosophila
>gi|1036826333|ref|XP_017137938.1| PREDICTED: UBX domain-containing protein 4 [Drosophila mira
>gi|1036826352|ref|XP_017137939.1| PREDICTED: ribonuclease Oy [Drosophila miranda]
>gi|1036847978|ref|XP_017139145.1| PREDICTED: homeotic protein empty spiracles [Drosophila mira
>gi|1036810752|ref|XP_017137081.1| PREDICTED: signal recognition particle 9 kDa protein [Drosop
>gi|1036810564|ref|XP_017137071.1| PREDICTED: JNK-interacting protein 3 isoform X1 [Drosophila
>gi|1036810583|ref|XP_017137072.1| PREDICTED: JNK-interacting protein 3 isoform X2 [Drosophila
>gi|1036810602|ref|XP_017137073.1| PREDICTED: JNK-interacting protein 3 isoform X3 [Drosophila
>gi|1036810677|ref|XP_017137077.1| PREDICTED: JNK-interacting protein 3 isoform X7 [Drosophila
>gi|1036810640|ref|XP_017137075.1| PREDICTED: JNK-interacting protein 3 isoform X5 [Drosophila
>gi|1036810621|ref|XP_017137074.1| PREDICTED: JNK-interacting protein 3 isoform X4 [Drosophila
>gi|1036810696|ref|XP_017137078.1| PREDICTED: JNK-interacting protein 3 isoform X8 [Drosophila
>gi|1036810714|ref|XP_017137079.1| PREDICTED: JNK-interacting protein 3 isoform X9 [Drosophila
>gi|1036810659|ref|XP_017137076.1| PREDICTED: JNK-interacting protein 3 isoform X6 [Drosophila
>gi|1036810733|ref|XP_017137080.1| PREDICTED: JNK-interacting protein 3 isoform X10 [Drosophila
>gi|1036836561|ref|XP_017138518.1| PREDICTED: transmembrane protein 43 homolog [Drosophila mira
>gi|1036836579|ref|XP_017138519.1| PREDICTED: uncharacterized protein LOC108153208 [Drosophila
>gi|1036798749|ref|XP_017136415.1| PREDICTED: protein ECT2 [Drosophila miranda]
>gi|1036798768|ref|XP_017136416.1| PREDICTED: eukaryotic translation initiation factor 4E [Dros
>gi|1036760408|ref|XP_017156894.1| PREDICTED: uncharacterized protein LOC108165369 [Drosophila
>gi|1036810295|ref|XP_017137055.1| PREDICTED: hepatic leukemia factor isoform X11 [Drosophila
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>gi|1036810312|ref|XP_017137056.1| PREDICTED: hepatic leukemia factor isoform X12 [Drosophila r
>gi|1036810329|ref|XP_017137058.1| PREDICTED: hepatic leukemia factor isoform X13 [Drosophila r
>gi|1036810113|ref|XP_017137045.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X1
>gi|1036810150|ref|XP_017137047.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X3
>gi|1036810222|ref|XP_017137051.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X7
>gi|1036810132|ref|XP_017137046.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X2
>gi|1036810169|ref|XP_017137048.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X4
>gi|1036810187|ref|XP_017137049.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X5
>gi|1036810205|ref|XP_017137050.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X6
>gi|1036810278|ref|XP_017137054.1| PREDICTED: DNA N6-methyl adenine demethylase isoform X10 [Di
>gi|1036810242|ref|XP_017137052.1| PREDICTED: transcription factor mef2A isoform X8 [Drosophila
>gi|1036810261|ref|XP_017137053.1| PREDICTED: transcription factor mef2A isoform X9 [Drosophila
>gi|1036832589|ref|XP_017138296.1| PREDICTED: uncharacterized protein LOC108153045 [Drosophila
>gi|1036832606|ref|XP_017138297.1| PREDICTED: uncharacterized protein LOC108153045 [Drosophila
>gi|1036760426|ref|XP_017156895.1| PREDICTED: circadian locomoter output cycles protein kaput
>gi|1036837419|ref|XP_017138568.1| PREDICTED: protein henna [Drosophila miranda]
>gi|1036837401|ref|XP_017138567.1| PREDICTED: uncharacterized protein LOC108153239 isoform X2
>gi|1036837382|ref|XP_017138566.1| PREDICTED: fibrillin-1 isoform X1 [Drosophila miranda]
>gi|1036837435|ref|XP_017138569.1| PREDICTED: heat shock protein 27 [Drosophila miranda]
>gi|1036760445|ref|XP_017156896.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036848732|ref|XP_017139187.1| PREDICTED: sensory neuron membrane protein 2 isoform X2 [Dro
>gi|1036848695|ref|XP_017139185.1| PREDICTED: sensory neuron membrane protein 2 isoform X1 [Dro
>gi|1036848713|ref|XP_017139186.1| PREDICTED: sensory neuron membrane protein 2 isoform X1 [Dro
>gi|1036848750|ref|XP_017139188.1| PREDICTED: sensory neuron membrane protein 2 isoform X3 [Dro
>gi|1036848771|ref|XP_017139189.1| PREDICTED: sensory neuron membrane protein 2 isoform X4 [Dro
>gi|1036847236|ref|XP_017139103.1| PREDICTED: uncharacterized protein LOC108153548 isoform X1
>gi|1036847253|ref|XP_017139104.1| PREDICTED: uncharacterized protein LOC108153548 isoform X2
>gi|1036760462|ref|XP_017156897.1| PREDICTED: probable serine/threonine-protein kinase DDB_G026
>gi|1036806315|ref|XP_017136838.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806333|ref|XP_017136839.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806375|ref|XP_017136841.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806394|ref|XP_017136843.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806352|ref|XP_017136840.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036822288|ref|XP_017137712.1| PREDICTED: WASH complex subunit strumpellin homolog [Drosoph
>gi|1036822307|ref|XP_017137713.1| PREDICTED: phosphoglucomutase [Drosophila miranda]
>gi|1036799926|ref|XP_017136480.1| PREDICTED: LOW QUALITY PROTEIN: nuclear pore complex protein
>gi|1036799944|ref|XP_017136481.1| PREDICTED: translocon-associated protein subunit beta [Drosc
>gi|1036799960|ref|XP_017136482.1| PREDICTED: probable trafficking protein particle complex sul
>gi|1036799907|ref|XP_017136479.1| PREDICTED: ATPase family AAA domain-containing protein 5 [Di
>gi|1036799977|ref|XP_017136484.1| PREDICTED: cytochrome b5 [Drosophila miranda]
>gi|1036779161|ref|XP_017135297.1| PREDICTED: P17/29C-like protein DDB_G0287399 [Drosophila mi]
>gi|1036779181|ref|XP_017135298.1| PREDICTED: P17/29C-like protein DDB_G0287399 [Drosophila mi]
>gi|1036779144|ref|XP_017135296.1| PREDICTED: LOW QUALITY PROTEIN: protein arginine N-methyltra
>gi|1036779443|ref|XP_017135313.1| PREDICTED: uncharacterized protein LOC108151290 isoform X4
>gi|1036779425|ref|XP_017135312.1| PREDICTED: uncharacterized protein LOC108151290 isoform X3
>gi|1036779459|ref|XP_017135314.1| PREDICTED: uncharacterized protein LOC108151290 isoform X5
>gi|1036779406|ref|XP_017135311.1| PREDICTED: uncharacterized protein LOC108151290 isoform X2
>gi|1036779387|ref|XP_017135310.1| PREDICTED: uncharacterized protein LOC108151290 isoform X1
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>gi|1036760478|ref|XP_017156898.1| PREDICTED: uncharacterized protein LOC108165373 [Drosophila
>gi|1036760495|ref|XP_017156900.1| PREDICTED: homeotic protein female sterile [Drosophila miran
>gi|1036779107|ref|XP_017135294.1| PREDICTED: U3 small nucleolar RNA-associated protein 4 homol
>gi|1036779199|ref|XP_017135299.1| PREDICTED: protein disulfide-isomerase TMX3 isoform X1 [Dros
>gi|1036779218|ref|XP_017135300.1| PREDICTED: protein disulfide-isomerase TMX3 isoform X2 [Dros
>gi|1036779604|ref|XP_017135323.1| PREDICTED: programmed cell death protein 5 [Drosophila miran
>gi|1036779641|ref|XP_017135325.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036779348|ref|XP_017135308.1| PREDICTED: iron-sulfur cluster co-chaperone protein HscB, m
>gi|1036779237|ref|XP_017135301.1| PREDICTED: gamma-glutamyl hydrolase isoform X1 [Drosophila r
>gi|1036779256|ref|XP_017135303.1| PREDICTED: gamma-glutamyl hydrolase A isoform X2 [Drosophila
>gi|1036779312|ref|XP_017135306.1| PREDICTED: gamma-glutamyl hydrolase isoform X1 [Drosophila r
>gi|1036779329|ref|XP_017135307.1| PREDICTED: gamma-glutamyl hydrolase A isoform X2 [Drosophila
>gi|1036779293|ref|XP_017135305.1| PREDICTED: homeobox protein 5 [Drosophila miranda]
>gi|1036779274|ref|XP_017135304.1| PREDICTED: dentin sialophosphoprotein [Drosophila miranda]
>gi|1036779819|ref|XP_017135336.1| PREDICTED: uncharacterized protein LOC108151309 isoform X2
>gi|1036779804|ref|XP_017135335.1| PREDICTED: uncharacterized protein LOC108151309 isoform X1
>gi|1036779909|ref|XP_017135342.1| PREDICTED: uncharacterized protein LOC108151314 [Drosophila
>gi|1036779928|ref|XP_017135343.1| PREDICTED: uncharacterized protein LOC108151315 [Drosophila
>gi|1036760511|ref|XP_017156901.1| PREDICTED: uncharacterized protein LOC108165375 [Drosophila
>gi|1036779854|ref|XP_017135339.1| PREDICTED: cuticle protein 38 [Drosophila miranda]
>gi|1036779124|ref|XP_017135295.1| PREDICTED: leucine-rich repeat-containing G-protein coupled
>gi|1036779660|ref|XP_017135327.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mi
>gi|1036779837|ref|XP_017135337.1| PREDICTED: EMI domain-containing protein 1 [Drosophila miran
>gi|1036779945|ref|XP_017135344.1| PREDICTED: uncharacterized protein LOC108151316 [Drosophila
>gi|1036779513|ref|XP_017135318.1| PREDICTED: cuticle protein 10.6 [Drosophila miranda]
>gi|1036779549|ref|XP_017135320.1| PREDICTED: glycine-rich protein [Drosophila miranda]
>gi|1036779873|ref|XP_017135340.1| PREDICTED: uncharacterized protein LOC108151312 [Drosophila
>gi|1036779476|ref|XP_017135316.1| PREDICTED: cuticle protein 38 [Drosophila miranda]
>gi|1036760529|ref|XP_017156902.1| PREDICTED: pollen-specific leucine-rich repeat extensin-like
>gi|1036760545|ref|XP_017156903.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mi
>gi|1036779368|ref|XP_017135309.1| PREDICTED: transcription factor btd [Drosophila miranda]
>gi|1036760563|ref|XP_017156904.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036779568|ref|XP_017135321.1| PREDICTED: cuticle protein 10.6 [Drosophila miranda]
>gi|1036779714|ref|XP_017135330.1| PREDICTED: uncharacterized protein LOC108151305 [Drosophila
>gi|1036779532|ref|XP_017135319.1| PREDICTED: pupal cuticle protein C1B [Drosophila miranda]
>gi|1036779587|ref|XP_017135322.1| PREDICTED: cuticle protein LPCP-23 [Drosophila miranda]
>gi|1036779733|ref|XP_017135331.1| PREDICTED: pupal cuticle protein C1B [Drosophila miranda]
>gi|1036779752|ref|XP_017135332.1| PREDICTED: uncharacterized protein LOC108151307 [Drosophila
>gi|1036779771|ref|XP_017135333.1| PREDICTED: uncharacterized protein LOC108151307 [Drosophila
>gi|1036779789|ref|XP_017135334.1| PREDICTED: uncharacterized protein LOC108151307 [Drosophila
>gi|1036779697|ref|XP_017135329.1| PREDICTED: uncharacterized histidine-rich protein DDB_G02745
>gi|1036779891|ref|XP_017135341.1| PREDICTED: neuropeptide-like 3 [Drosophila miranda]
>gi|1036779679|ref|XP_017135328.1| PREDICTED: uncharacterized protein LOC108151303 [Drosophila
>gi|1036779622|ref|XP_017135324.1| PREDICTED: pupal cuticle protein Edg-84A-like [Drosophila m
>gi|1036779964|ref|XP_017135346.1| PREDICTED: uncharacterized protein LOC108151317 [Drosophila
>gi|1036779494|ref|XP_017135317.1| PREDICTED: cuticle protein LPCP-23 [Drosophila miranda]
>gi|1036768478|ref|XP_017134684.1| PREDICTED: cuticle protein [Drosophila miranda]
>gi|1036766977|ref|XP_017134605.1| PREDICTED: uncharacterized protein LOC108150811 [Drosophila
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>gi|1036767983|ref|XP_017134657.1| PREDICTED: uncharacterized protein LOC108150865 [Drosophila
>gi|1036851525|ref|XP_017139354.1| PREDICTED: E3 ubiquitin-protein ligase TRIM33 [Drosophila m
>gi|1036851506|ref|XP_017139352.1| PREDICTED: trimeric intracellular cation channel type B [Dro
>gi|1036760581|ref|XP_017156905.1| PREDICTED: uncharacterized protein LOC108165379 [Drosophila
>gi|1036851180|ref|XP_017139332.1| PREDICTED: uncharacterized protein LOC108153716 [Drosophila
>gi|1036851199|ref|XP_017139333.1| PREDICTED: uncharacterized protein LOC108153717 [Drosophila
>gi|1036827712|ref|XP_017138019.1| PREDICTED: uncharacterized protein LOC108152891 isoform X2
>gi|1036827694|ref|XP_017138018.1| PREDICTED: uncharacterized protein LOC108152891 isoform X1
>gi|1036760600|ref|XP_017156907.1| PREDICTED: protein anoxia up-regulated [Drosophila miranda]
>gi|1036827750|ref|XP_017138022.1| PREDICTED: uncharacterized protein LOC108152893 [Drosophila
>gi|1036827731|ref|XP_017138020.1| PREDICTED: coiled-coil domain-containing protein 186 [Drosop
>gi|1036837536|ref|XP_017138575.1| PREDICTED: probable tRNA N6-adenosine threonylcarbamoyltrans
>gi|1036837573|ref|XP_017138578.1| PREDICTED: alcohol dehydrogenase isoform X2 [Drosophila mira
>gi|1036837554|ref|XP_017138576.1| PREDICTED: alcohol dehydrogenase isoform X1 [Drosophila mira
>gi|1036837594|ref|XP_017138579.1| PREDICTED: TPPP family protein CG45057 isoform X1 [Drosophil
>gi|1036837611|ref|XP_017138580.1| PREDICTED: TPPP family protein CG45057 isoform X2 [Drosophil
>gi|1036767887|ref|XP_017134651.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036770141|ref|XP_017134777.1| PREDICTED: adult-specific rigid cuticular protein 15.7 [Dros
>gi|1036760618|ref|XP_017156908.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036851237|ref|XP_017139336.1| PREDICTED: alcohol dehydrogenase-like [Drosophila miranda]
>gi|1036779038|ref|XP_017135289.1| PREDICTED: roquin-1 [Drosophila miranda]
>gi|1036779055|ref|XP_017135291.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036779071|ref|XP_017135292.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036778929|ref|XP_017135283.1| PREDICTED: uncharacterized protein LOC108151271 isoform X1
>gi|1036778981|ref|XP_017135286.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036778999|ref|XP_017135287.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036779017|ref|XP_017135288.1| PREDICTED: bromodomain-containing protein 4 [Drosophila mira
>gi|1036778945|ref|XP_017135284.1| PREDICTED: uncharacterized protein LOC108151271 isoform X2
>gi|1036778963|ref|XP_017135285.1| PREDICTED: uncharacterized protein LOC108151271 isoform X3
>gi|1036779088|ref|XP_017135293.1| PREDICTED: uncharacterized protein LOC108151276 [Drosophila
>gi|1036811725|ref|XP_017137133.1| PREDICTED: failed axon connections isoform X2 [Drosophila mi
>gi|1036811706|ref|XP_017137132.1| PREDICTED: failed axon connections isoform X1 [Drosophila mi
>gi|1036775285|ref|XP_017135068.1| PREDICTED: serine incorporator 1 isoform X1 [Drosophila mira
>gi|1036775304|ref|XP_017135069.1| PREDICTED: serine incorporator 1 isoform X2 [Drosophila mira
>gi|1036775322|ref|XP_017135070.1| PREDICTED: serine incorporator 1 isoform X3 [Drosophila mira
>gi|1036775251|ref|XP_017135065.1| PREDICTED: probable glutamate--tRNA ligase, mitochondrial [D
>gi|1036775233|ref|XP_017135064.1| PREDICTED: tyrosine--tRNA ligase, cytoplasmic [Drosophila mi
>gi|1036775161|ref|XP_017135060.1| PREDICTED: elongation factor-like GTPase 1 [Drosophila miran
>gi|1036775178|ref|XP_017135061.1| PREDICTED: uncharacterized protein LOC108151146 [Drosophila
>gi|1036775269|ref|XP_017135066.1| PREDICTED: protein giant-lens [Drosophila miranda]
>gi|1036775215|ref|XP_017135063.1| PREDICTED: proton-coupled folate transporter [Drosophila mi
>gi|1036775379|ref|XP_017135073.1| PREDICTED: pupal cuticle protein C1B-like [Drosophila mirand
>gi|1036775398|ref|XP_017135074.1| PREDICTED: pupal cuticle protein C1B-like [Drosophila mirand
>gi|1036775196|ref|XP_017135062.1| PREDICTED: protein scarlet [Drosophila miranda]
>gi|1036775086|ref|XP_017135056.1| PREDICTED: uncharacterized protein LOC108151144 isoform X1
>gi|1036775105|ref|XP_017135057.1| PREDICTED: adenylate cyclase type 5 isoform X2 [Drosophila m
>gi|1036775124|ref|XP_017135058.1| PREDICTED: adenylate cyclase type 5 isoform X2 [Drosophila m
>gi|1036775143|ref|XP_017135059.1| PREDICTED: adenylate cyclase type 5 isoform X2 [Drosophila m
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>gi|1036775339|ref|XP_017135071.1| PREDICTED: uncharacterized protein LOC108151154 [Drosophila
>gi|1036775358|ref|XP_017135072.1| PREDICTED: uncharacterized protein LOC108151154 [Drosophila
>gi|1036760636|ref|XP_017156909.1| PREDICTED: uncharacterized protein LOC108165382 [Drosophila
>gi|1036789499|ref|XP_017135893.1| PREDICTED: multiple inositol polyphosphate phosphatase 1 [D
>gi|1036789588|ref|XP_017135898.1| PREDICTED: LOW QUALITY PROTEIN: endothelial differentiation-
>gi|1036789606|ref|XP_017135900.1| PREDICTED: nuclear RNA export factor 2 [Drosophila miranda]
>gi|1036789553|ref|XP_017135896.1| PREDICTED: survival motor neuron protein [Drosophila miranda
>gi|1036789537|ref|XP_017135895.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 8 [
>gi|1036789518|ref|XP_017135894.1| PREDICTED: cytochrome b-c1 complex subunit 2, mitochondrial
>gi|1036789571|ref|XP_017135897.1| PREDICTED: syntaxin-8 [Drosophila miranda]
>gi|1036835974|ref|XP_017138482.1| PREDICTED: polycomb group RING finger protein 3 [Drosophila
>gi|1036835989|ref|XP_017138483.1| PREDICTED: female-specific protein transformer [Drosophila m
>gi|1036780737|ref|XP_017135389.1| PREDICTED: probable GPI-anchored adhesin-like protein PGA55
>gi|1036780755|ref|XP_017135390.1| PREDICTED: importin-4 [Drosophila miranda]
>gi|1036780826|ref|XP_017135395.1| PREDICTED: ADP-ribose pyrophosphatase, mitochondrial isofor
>gi|1036780808|ref|XP_017135394.1| PREDICTED: ADP-ribose pyrophosphatase, mitochondrial isofor
>gi|1036780843|ref|XP_017135396.1| PREDICTED: proteasome subunit beta type-1 [Drosophila miran
>gi|1036780861|ref|XP_017135397.1| PREDICTED: protein twisted gastrulation [Drosophila miranda]
>gi|1036780772|ref|XP_017135391.1| PREDICTED: RNA-binding protein 45 isoform X1 [Drosophila mi
>gi|1036780790|ref|XP_017135392.1| PREDICTED: RNA-binding protein 45 isoform X2 [Drosophila mi
>gi|1036811554|ref|XP_017137124.1| PREDICTED: general transcription factor IIE subunit 2 [Dros
>gi|1036811496|ref|XP_017137121.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036811515|ref|XP_017137122.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036811573|ref|XP_017137125.1| PREDICTED: transmembrane protein 50A [Drosophila miranda]
>gi|1036811535|ref|XP_017137123.1| PREDICTED: nucleolin 2 [Drosophila miranda]
>gi|1036795399|ref|XP_017136226.1| PREDICTED: LOW QUALITY PROTEIN: salivary glue protein Sgs-3
>gi|1036795380|ref|XP_017136225.1| PREDICTED: sorting nexin lst-4 [Drosophila miranda]
>gi|1036795418|ref|XP_017136227.1| PREDICTED: COPII coat assembly protein SEC16 [Drosophila mi
>gi|1036822794|ref|XP_017137740.1| PREDICTED: uncharacterized protein LOC108152717 isoform X1
>gi|1036822813|ref|XP_017137742.1| PREDICTED: uncharacterized protein LOC108152717 isoform X2
>gi|1036822870|ref|XP_017137745.1| PREDICTED: trafficking protein particle complex subunit 3 [
>gi|1036822832|ref|XP_017137743.1| PREDICTED: nuclear transcription factor Y subunit alpha [Dro
>gi|1036822851|ref|XP_017137744.1| PREDICTED: nuclear transcription factor Y subunit alpha [Dro
>gi|1036829711|ref|XP_017138131.1| PREDICTED: TOM1-like protein 2 isoform X1 [Drosophila miran
>gi|1036829789|ref|XP_017138136.1| PREDICTED: TOM1-like protein 2 isoform X5 [Drosophila miran
>gi|1036829730|ref|XP_017138132.1| PREDICTED: TOM1-like protein 2 isoform X2 [Drosophila miran
>gi|1036829749|ref|XP_017138133.1| PREDICTED: TOM1-like protein 2 isoform X3 [Drosophila miran
>gi|1036829770|ref|XP_017138134.1| PREDICTED: TOM1-like protein 2 isoform X4 [Drosophila miran
>gi|1036829808|ref|XP_017138137.1| PREDICTED: uncharacterized protein LOC108152949 [Drosophila
>gi|1036778100|ref|XP_017135233.1| PREDICTED: uncharacterized protein LOC108151243 [Drosophila
>gi|1036778020|ref|XP_017135228.1| PREDICTED: uncharacterized protein LOC108151239 [Drosophila
>gi|1036778084|ref|XP_017135232.1| PREDICTED: SAC3 domain-containing protein 1 [Drosophila mira
>gi|1036778068|ref|XP_017135231.1| PREDICTED: GDP-D-glucose phosphorylase 1 [Drosophila miranda
>gi|1036778051|ref|XP_017135230.1| PREDICTED: queuine tRNA-ribosyltransferase subunit QTRTD1 ho
>gi|1036778132|ref|XP_017135235.1| PREDICTED: uncharacterized protein LOC108151244 isoform X2
>gi|1036778116|ref|XP_017135234.1| PREDICTED: probable serine/threonine-protein kinase nek3 iso
>gi|1036787226|ref|XP_017135762.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase isozyme L5
>gi|1036787207|ref|XP_017135761.1| PREDICTED: mucin-19 [Drosophila miranda]
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>gi|1036787243|ref|XP_017135763.1| PREDICTED: F-box only protein 28 [Drosophila miranda]
>gi|1036760655|ref|XP_017156910.1| PREDICTED: uncharacterized protein LOC108165383 [Drosophila
>gi|1036795286|ref|XP_017136220.1| PREDICTED: proton-coupled amino acid transporter 2 isoform )
>gi|1036795305|ref|XP_017136221.1| PREDICTED: proton-coupled amino acid transporter 2 isoform )
>gi|1036795322|ref|XP_017136223.1| PREDICTED: leucine-rich repeat-containing protein 59 [Droso]
>gi|1036795343|ref|XP_017136224.1| PREDICTED: 40S ribosomal protein S9 [Drosophila miranda]
>gi|1036768060|ref|XP_017134661.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036836076|ref|XP_017138488.1| PREDICTED: sodium-independent sulfate anion transporter iso:
>gi|1036836095|ref|XP_017138489.1| PREDICTED: sodium-independent sulfate anion transporter iso:
>gi|1036836059|ref|XP_017138487.1| PREDICTED: sodium-independent sulfate anion transporter iso:
>gi|1036833285|ref|XP_017138336.1| PREDICTED: CCR4-NOT transcription complex subunit 7 isoform
>gi|1036833303|ref|XP_017138337.1| PREDICTED: CCR4-NOT transcription complex subunit 7 isoform
>gi|1036805840|ref|XP_017136815.1| PREDICTED: PAX-interacting protein 1 [Drosophila miranda]
>gi|1036788020|ref|XP_017135809.1| PREDICTED: uncharacterized protein LOC108151621 [Drosophila
>gi|1036788039|ref|XP_017135810.1| PREDICTED: uncharacterized protein LOC108151621 [Drosophila
>gi|1036836954|ref|XP_017138539.1| PREDICTED: nitrogen permease regulator 3-like protein [Droso
>gi|1036835497|ref|XP_017138453.1| PREDICTED: zinc finger protein 391 isoform X1 [Drosophila m:
>gi|1036835516|ref|XP_017138454.1| PREDICTED: zinc finger protein 425 isoform X2 [Drosophila m:
>gi|1036760672|ref|XP_017156911.1| PREDICTED: uncharacterized zinc finger protein CG2678 [Dros
>gi|1036836540|ref|XP_017138515.1| PREDICTED: probable 28S ribosomal protein S6, mitochondrial
>gi|1036836502|ref|XP_017138513.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036836521|ref|XP_017138514.1| PREDICTED: probable RNA methyltransferase CG11342 [Drosophil
>gi|1036843837|ref|XP_017138927.1| PREDICTED: beta-2 adrenergic receptor [Drosophila miranda]
>gi|1036766845|ref|XP_017134597.1| PREDICTED: uncharacterized protein LOC108150805 [Drosophila
>gi|1036844247|ref|XP_017138949.1| PREDICTED: uncharacterized protein LOC108153444 isoform X2
>gi|1036844228|ref|XP_017138948.1| PREDICTED: uncharacterized protein LOC108153444 isoform X1
>gi|1036835592|ref|XP_017138459.1| PREDICTED: syntaxin-17 [Drosophila miranda]
>gi|1036835611|ref|XP_017138460.1| PREDICTED: protein LLP homolog [Drosophila miranda]
>gi|1036824507|ref|XP_017137839.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824526|ref|XP_017137840.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824545|ref|XP_017137841.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824562|ref|XP_017137842.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824488|ref|XP_017137838.1| PREDICTED: uncharacterized protein LOC108152771 [Drosophila
>gi|1036842053|ref|XP_017138826.1| PREDICTED: acetylcholine receptor subunit beta-like 1 isofo:
>gi|1036842034|ref|XP_017138825.1| PREDICTED: acetylcholine receptor subunit beta-like 1 isofo:
>gi|1036851781|ref|XP_017139369.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036767905|ref|XP_017134652.1| PREDICTED: cyclin-dependent kinase inhibitor 1C [Drosophila
>gi|1036767944|ref|XP_017134654.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036760691|ref|XP_017156912.1| PREDICTED: early nodulin-75 [Drosophila miranda]
>gi|1036760709|ref|XP_017156913.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036760728|ref|XP_017156914.1| PREDICTED: uncharacterized protein LOC108165388 [Drosophila
>gi|1036851033|ref|XP_017139324.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036851069|ref|XP_017139326.1| PREDICTED: uncharacterized protein LOC108153710 isoform X1
>gi|1036851088|ref|XP_017139327.1| PREDICTED: uncharacterized protein LOC108153710 isoform X2
>gi|1036851050|ref|XP_017139325.1| PREDICTED: skin secretory protein xP2 [Drosophila miranda]
>gi|1036852490|ref|XP_017139413.1| PREDICTED: uncharacterized protein LOC108153782 isoform X1
>gi|1036852508|ref|XP_017139414.1| PREDICTED: uncharacterized protein LOC108153782 isoform X2
>gi|1036760744|ref|XP_017156915.1| PREDICTED: uncharacterized protein LOC108165389 [Drosophila
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>gi|1036851164|ref|XP_017139331.1| PREDICTED: LOW QUALITY PROTEIN: extensin-2 [Drosophila miran
>gi|1036846273|ref|XP_017139047.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036769665|ref|XP_017134753.1| PREDICTED: glycine and tyrosine-rich protein [Drosophila mir
>gi|1036760763|ref|XP_017156916.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036850470|ref|XP_017139291.1| PREDICTED: uncharacterized protein LOC108153684 isoform X1
>gi|1036850487|ref|XP_017139292.1| PREDICTED: uncharacterized protein LOC108153684 isoform X1
>gi|1036850506|ref|XP_017139293.1| PREDICTED: uncharacterized protein LOC108153684 isoform X1
>gi|1036850542|ref|XP_017139295.1| PREDICTED: uncharacterized protein LOC108153684 isoform X3
>gi|1036850525|ref|XP_017139294.1| PREDICTED: uncharacterized protein LOC108153684 isoform X2
>gi|1036850559|ref|XP_017139296.1| PREDICTED: uncharacterized protein LOC108153684 isoform X4
>gi|1036844062|ref|XP_017138938.1| PREDICTED: LOW QUALITY PROTEIN: mucin-5AC, partial [Drosoph
>gi|1036760782|ref|XP_017156918.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036819139|ref|XP_017137546.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036819160|ref|XP_017137547.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036819181|ref|XP_017137548.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036819219|ref|XP_017137551.1| PREDICTED: uncharacterized protein LOC108152603 isoform X1
>gi|1036819240|ref|XP_017137552.1| PREDICTED: uncharacterized protein LOC108152603 isoform X2
>gi|1036819200|ref|XP_017137549.1| PREDICTED: beta-1,3-galactosyltransferase 5 [Drosophila mir
>gi|1036844097|ref|XP_017138940.1| PREDICTED: band 7 protein AGAP004871 isoform X2 [Drosophila
>gi|1036844081|ref|XP_017138939.1| PREDICTED: band 7 protein AGAP004871 isoform X1 [Drosophila
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>gi|1036844190|ref|XP_017138946.1| PREDICTED: band 7 protein AGAP004871 isoform X7 [Drosophila
>gi|1036844115|ref|XP_017138941.1| PREDICTED: band 7 protein AGAP004871 isoform X3 [Drosophila
>gi|1036844134|ref|XP_017138942.1| PREDICTED: band 7 protein AGAP004871 isoform X4 [Drosophila
>gi|1036805165|ref|XP_017136776.1| PREDICTED: tyrosine-protein kinase Src64B isoform X2 [Droso
>gi|1036805146|ref|XP_017136775.1| PREDICTED: tyrosine-protein kinase Src64B isoform X1 [Droso
>gi|1036826750|ref|XP_017137963.1| PREDICTED: histone deacetylase Rpd3 isoform X1 [Drosophila
>gi|1036826769|ref|XP_017137964.1| PREDICTED: histone deacetylase Rpd3 isoform X2 [Drosophila
>gi|1036819572|ref|XP_017137568.1| PREDICTED: uncharacterized protein LOC108152614 isoform X1
>gi|1036819591|ref|XP_017137569.1| PREDICTED: uncharacterized protein LOC108152614 isoform X1
>gi|1036819666|ref|XP_017137574.1| PREDICTED: contactin-associated protein-like 2 isoform X5 [
>gi|1036819742|ref|XP_017137578.1| PREDICTED: CASP-like protein 4U1 [Drosophila miranda]
>gi|1036819628|ref|XP_017137572.1| PREDICTED: uncharacterized protein LOC108152614 isoform X3
>gi|1036819610|ref|XP_017137570.1| PREDICTED: uncharacterized protein LOC108152614 isoform X2
>gi|1036819647|ref|XP_017137573.1| PREDICTED: contactin-associated protein-like 2 isoform X4 [
>gi|1036819685|ref|XP_017137575.1| PREDICTED: probable tubulin polyglutamylase TTLL1 [Drosophi
>gi|1036819704|ref|XP_017137576.1| PREDICTED: uncharacterized protein LOC108152616 [Drosophila
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>gi|1036851998|ref|XP_017139384.1| PREDICTED: platelet glycoprotein V isoform X2 [Drosophila mi
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>gi|1036851981|ref|XP_017139383.1| PREDICTED: platelet glycoprotein V isoform X1 [Drosophila mi
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>gi|1036794285|ref|XP_017136163.1| PREDICTED: uncharacterized oxidoreductase YjmC [Drosophila m
>gi|1036794247|ref|XP_017136161.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X3 [Dros
>gi|1036794266|ref|XP_017136162.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X4 [Dros
>gi|1036794211|ref|XP_017136159.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X1 [Dros
>gi|1036794229|ref|XP_017136160.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X2 [Dros
>gi|1036828986|ref|XP_017138091.1| PREDICTED: probable leucine--tRNA ligase, mitochondrial [Dr
>gi|1036828966|ref|XP_017138090.1| PREDICTED: uncharacterized protein LOC108152921 [Drosophila
>gi|1036848806|ref|XP_017139191.1| PREDICTED: uncharacterized protein LOC108153606 isoform X2
>gi|1036848787|ref|XP_017139190.1| PREDICTED: uncharacterized protein LOC108153606 isoform X1
>gi|1036839865|ref|XP_017138709.1| PREDICTED: dynein heavy chain 3, axonemal [Drosophila miran
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>gi|1036848457|ref|XP_017139171.1| PREDICTED: ammonium transporter Rh type B [Drosophila miran
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>gi|1036842499|ref|XP_017138849.1| PREDICTED: ubiquitin-conjugating enzyme E2 L3 [Drosophila m
>gi|1036842517|ref|XP_017138850.1| PREDICTED: uncharacterized protein LOC108153387 [Drosophila
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>gi|1036760820|ref|XP_017156920.1| PREDICTED: uncharacterized protein LOC108165393 [Drosophila
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>gi|1036815278|ref|XP_017137327.1| PREDICTED: lysozyme S [Drosophila miranda]
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>gi|1036815240|ref|XP_017137324.1| PREDICTED: lysozyme B-like [Drosophila miranda]
>gi|1036815296|ref|XP_017137328.1| PREDICTED: lysozyme D-like [Drosophila miranda]
>gi|1036815395|ref|XP_017137333.1| PREDICTED: lysozyme B-like [Drosophila miranda]
>gi|1036815374|ref|XP_017137332.1| PREDICTED: lysozyme D-like isoform X2 [Drosophila miranda]
>gi|1036815355|ref|XP_017137331.1| PREDICTED: lysozyme D-like isoform X1 [Drosophila miranda]
>gi|1036815202|ref|XP_017137322.1| PREDICTED: lysozyme A/C-like [Drosophila miranda]
>gi|1036815220|ref|XP_017137323.1| PREDICTED: lysozyme A/C-like [Drosophila miranda]
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>gi|1036815183|ref|XP_017137321.1| PREDICTED: ester hydrolase C11orf54 homolog [Drosophila mira
>gi|1036797501|ref|XP_017136344.1| PREDICTED: mitochondrial carrier homolog 2 [Drosophila mira
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>gi|1036797393|ref|XP_017136338.1| PREDICTED: uncharacterized protein LOC108151926 isoform X2
>gi|1036797335|ref|XP_017136335.1| PREDICTED: uncharacterized protein LOC108151926 isoform X1
>gi|1036797356|ref|XP_017136336.1| PREDICTED: uncharacterized protein LOC108151926 isoform X1
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>gi|1036797464|ref|XP_017136342.1| PREDICTED: arrestin domain-containing protein 4 [Drosophila
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>gi|1036818759|ref|XP_017137524.1| PREDICTED: breast cancer anti-estrogen resistance protein 1
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>gi|1036818797|ref|XP_017137526.1| PREDICTED: breast cancer anti-estrogen resistance protein 1
>gi|1036818816|ref|XP_017137527.1| PREDICTED: WD repeat-containing protein 78 [Drosophila mirar
>gi|1036807754|ref|XP_017136918.1| PREDICTED: 3-phosphoinositide-dependent protein kinase 1 iso
>gi|1036807770|ref|XP_017136919.1| PREDICTED: 3-phosphoinositide-dependent protein kinase 1 iso
>gi|1036852322|ref|XP_017139404.1| PREDICTED: glycine-rich protein 5 [Drosophila miranda]
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>gi|1036781892|ref|XP_017135458.1| PREDICTED: G-protein coupled receptor Mth2 isoform X5 [Drosc
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>gi|1036775857|ref|XP_017135098.1| PREDICTED: nicotinamide riboside kinase 2 [Drosophila mirand
>gi|1036775948|ref|XP_017135103.1| PREDICTED: dolichol phosphate-mannose biosynthesis regulato)
>gi|1036775874|ref|XP_017135099.1| PREDICTED: phosphatidylinositol N-acetylglucosaminyltransfer
>gi|1036775837|ref|XP_017135097.1| PREDICTED: LOW QUALITY PROTEIN: zinc/cadmium resistance prot
>gi|1036760877|ref|XP_017156923.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
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>gi|1036774133|ref|XP_017135007.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso)
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>gi|1036777361|ref|XP_017135187.1| PREDICTED: DNA replication complex GINS protein PSF1 [Drosop
>gi|1036777342|ref|XP_017135186.1| PREDICTED: staphylococcal nuclease domain-containing protei
>gi|1036777397|ref|XP_017135189.1| PREDICTED: uncharacterized protein LOC108151219 [Drosophila
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>gi|1036765224|ref|XP_017157158.1| PREDICTED: uncharacterized protein LOC108165595 [Drosophila
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>gi|1036785536|ref|XP_017135670.1| PREDICTED: glycerol kinase [Drosophila miranda]
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>gi|1036796958|ref|XP_017136313.1| PREDICTED: PHD finger protein rhinoceros [Drosophila miranda
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>gi|1036810451|ref|XP_017137065.1| PREDICTED: serine/threonine-protein kinase NLK isoform X1 [I
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>gi|1036833893|ref|XP_017138361.1| PREDICTED: UBX domain-containing protein 1 [Drosophila miran
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>gi|1036801433|ref|XP_017136568.1| PREDICTED: major facilitator superfamily domain-containing p
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>gi|1036801469|ref|XP_017136570.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801488|ref|XP_017136571.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801507|ref|XP_017136572.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801526|ref|XP_017136573.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801544|ref|XP_017136574.1| PREDICTED: uncharacterized protein LOC108152065 [Drosophila
>gi|1036801615|ref|XP_017136579.1| PREDICTED: signal recognition particle 19 kDa protein [Droso
>gi|1036801580|ref|XP_017136576.1| PREDICTED: geranylgeranyl pyrophosphate synthase isoform X1
>gi|1036801596|ref|XP_017136577.1| PREDICTED: geranylgeranyl pyrophosphate synthase isoform X2
>gi|1036829556|ref|XP_017138122.1| PREDICTED: transcriptional repressor CTCF [Drosophila mirand
>gi|1036810405|ref|XP_017137062.1| PREDICTED: regulator of chromosome condensation [Drosophila
>gi|1036810392|ref|XP_017137061.1| PREDICTED: LOW QUALITY PROTEIN: motile sperm domain-containi
>gi|1036810419|ref|XP_017137063.1| PREDICTED: motile sperm domain-containing protein 2 [Drosoph
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>gi|1036810434|ref|XP_017137064.1| PREDICTED: motile sperm domain-containing protein 2 [Drosop
>gi|1036849251|ref|XP_017139217.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036852436|ref|XP_017139411.1| PREDICTED: uncharacterized protein LOC108153780 [Drosophila
>gi|1036760931|ref|XP_017156927.1| PREDICTED: organic cation transporter protein [Drosophila mi
>gi|1036833841|ref|XP_017138357.1| PREDICTED: glutaminyl-peptide cyclotransferase-like protein
>gi|1036833805|ref|XP_017138355.1| PREDICTED: glutaminyl-peptide cyclotransferase-like [Drosop
>gi|1036833823|ref|XP_017138356.1| PREDICTED: ribosome biogenesis regulatory protein homolog [
>gi|1036766082|ref|XP_017134557.1| PREDICTED: solute carrier family 22 member 2 [Drosophila mi
>gi|1036838372|ref|XP_017138625.1| PREDICTED: DNA polymerase epsilon subunit 2 [Drosophila mira
>gi|1036827576|ref|XP_017138012.1| PREDICTED: transcription factor btd [Drosophila miranda]
>gi|1036827538|ref|XP_017138010.1| PREDICTED: protein vein isoform X4 [Drosophila miranda]
>gi|1036827520|ref|XP_017138008.1| PREDICTED: protein vein isoform X3 [Drosophila miranda]
>gi|1036827501|ref|XP_017138007.1| PREDICTED: protein vein isoform X2 [Drosophila miranda]
>gi|1036827557|ref|XP_017138011.1| PREDICTED: protein vein isoform X5 [Drosophila miranda]
>gi|1036827489|ref|XP_017138006.1| PREDICTED: protein vein isoform X1 [Drosophila miranda]
>gi|1036792862|ref|XP_017136084.1| PREDICTED: mitotic spindle assembly checkpoint protein MAD2A
>gi|1036792809|ref|XP_017136082.1| PREDICTED: uracil phosphoribosyltransferase homolog [Drosopl
>gi|1036792792|ref|XP_017136081.1| PREDICTED: regulator of chromosome condensation-like [Drosop
>gi|1036792773|ref|XP_017136080.1| PREDICTED: ribosomal protein S6 kinase beta-2 [Drosophila mi
>gi|1036792754|ref|XP_017136079.1| PREDICTED: uncharacterized protein LOC108151752 [Drosophila
>gi|1036851800|ref|XP_017139370.1| PREDICTED: tektin-1 [Drosophila miranda]
>gi|1036821294|ref|XP_017137657.1| PREDICTED: E3 ubiquitin-protein ligase Bre1 [Drosophila mira
>gi|1036813489|ref|XP_017137225.1| PREDICTED: odorant receptor 67b [Drosophila miranda]
>gi|1036813449|ref|XP_017137223.1| PREDICTED: laminin subunit gamma-1 [Drosophila miranda]
>gi|1036813466|ref|XP_017137224.1| PREDICTED: laminin subunit gamma-1 [Drosophila miranda]
>gi|1036760947|ref|XP_017156928.1| PREDICTED: uncharacterized protein LOC108165400 [Drosophila
>gi|1036844022|ref|XP_017138936.1| PREDICTED: MORN repeat-containing protein 5 [Drosophila mira
>gi|1036844041|ref|XP_017138937.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036760965|ref|XP_017156929.1| PREDICTED: uncharacterized protein LOC108165401 [Drosophila
>gi|1036835955|ref|XP_017138481.1| PREDICTED: capon-like protein isoform X4 [Drosophila miranda
>gi|1036835938|ref|XP_017138480.1| PREDICTED: uncharacterized protein LOC108153181 isoform X3
>gi|1036835904|ref|XP_017138477.1| PREDICTED: capon-like protein isoform X2 [Drosophila miranda
>gi|1036835921|ref|XP_017138478.1| PREDICTED: capon-like protein isoform X2 [Drosophila miranda
>gi|1036835887|ref|XP_017138476.1| PREDICTED: capon-like protein isoform X1 [Drosophila miranda
>gi|1036760983|ref|XP_017156930.1| PREDICTED: uncharacterized protein LOC108165402 [Drosophila
>gi|1036761002|ref|XP_017156931.1| PREDICTED: uncharacterized protein LOC108165403 [Drosophila
>gi|1036825921|ref|XP_017137914.1| PREDICTED: DNA replication licensing factor Mcm7-like [Drosc
>gi|1036825807|ref|XP_017137907.1| PREDICTED: protein TANC1 isoform X1 [Drosophila miranda]
>gi|1036825825|ref|XP_017137908.1| PREDICTED: protein TANC1 isoform X1 [Drosophila miranda]
>gi|1036825863|ref|XP_017137910.1| PREDICTED: protein TANC2 isoform X3 [Drosophila miranda]
>gi|1036825844|ref|XP_017137909.1| PREDICTED: protein TANC1 isoform X2 [Drosophila miranda]
>gi|1036825882|ref|XP_017137912.1| PREDICTED: protein TANC2 isoform X4 [Drosophila miranda]
>gi|1036825903|ref|XP_017137913.1| PREDICTED: protein TANC2 isoform X5 [Drosophila miranda]
>gi|1036758102|ref|XP_017156757.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036842109|ref|XP_017138829.1| PREDICTED: isovaleryl-CoA dehydrogenase, mitochondrial [Dros
>gi|1036815963|ref|XP_017137364.1| PREDICTED: uncharacterized protein LOC108152501 isoform X1
>gi|1036815981|ref|XP_017137365.1| PREDICTED: uncharacterized protein LOC108152501 isoform X1
>gi|1036816038|ref|XP_017137369.1| PREDICTED: uncharacterized protein LOC108152501 isoform X4
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>gi|1036816002|ref|XP_017137367.1| PREDICTED: uncharacterized protein LOC108152501 isoform X2
>gi|1036816114|ref|XP_017137373.1| PREDICTED: uncharacterized protein LOC108152501 isoform X7
>gi|1036816077|ref|XP_017137371.1| PREDICTED: uncharacterized protein LOC108152501 isoform X6
>gi|1036816095|ref|XP_017137372.1| PREDICTED: uncharacterized protein LOC108152501 isoform X6
>gi|1036816019|ref|XP_017137368.1| PREDICTED: uncharacterized protein LOC108152501 isoform X3
>gi|1036816056|ref|XP_017137370.1| PREDICTED: uncharacterized protein LOC108152501 isoform X5
>gi|1036816133|ref|XP_017137374.1| PREDICTED: negative elongation factor E [Drosophila miranda]
>gi|1036829693|ref|XP_017138130.1| PREDICTED: 60S ribosomal protein L14 [Drosophila miranda]
>gi|1036829674|ref|XP_017138129.1| PREDICTED: LETM1 domain-containing protein 1 [Drosophila mi
>gi|1036818740|ref|XP_017137523.1| PREDICTED: uncharacterized protein LOC108152587 [Drosophila
>gi|1036818639|ref|XP_017137517.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1036818656|ref|XP_017137518.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1036818677|ref|XP_017137519.1| PREDICTED: TBC1 domain family member 13 isoform X1 [Drosophi
>gi|1036818719|ref|XP_017137522.1| PREDICTED: TBC1 domain family member 13 isoform X3 [Drosophi
>gi|1036818698|ref|XP_017137520.1| PREDICTED: TBC1 domain family member 13 isoform X2 [Drosophi
>gi|1036802318|ref|XP_017136619.1| PREDICTED: uncharacterized protein LOC108152090 isoform X1
>gi|1036802337|ref|XP_017136620.1| PREDICTED: uncharacterized protein LOC108152090 isoform X2
>gi|1036802391|ref|XP_017136623.1| PREDICTED: uncharacterized protein LOC108152090 isoform X5
>gi|1036802409|ref|XP_017136624.1| PREDICTED: uncharacterized protein LOC108152090 isoform X6
>gi|1036802355|ref|XP_017136621.1| PREDICTED: uncharacterized protein LOC108152090 isoform X3
>gi|1036802373|ref|XP_017136622.1| PREDICTED: PDZ and LIM domain protein 2 isoform X4 [Drosophi
>gi|1036802428|ref|XP_017136625.1| PREDICTED: PDZ and LIM domain protein 4 isoform X7 [Drosophi
>gi|1036802447|ref|XP_017136626.1| PREDICTED: uncharacterized protein LOC108152090 isoform X8
>gi|1036802466|ref|XP_017136627.1| PREDICTED: uncharacterized protein LOC108152090 isoform X9
>gi|1036802485|ref|XP_017136628.1| PREDICTED: uncharacterized protein LOC108152090 isoform X10
>gi|1036802299|ref|XP_017136618.1| PREDICTED: cell division control protein 6 homolog [Drosophi
>gi|1036802628|ref|XP_017136633.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036802680|ref|XP_017136635.1| PREDICTED: putative uncharacterized protein DDB_G0294196 iso
>gi|1036802661|ref|XP_017136634.1| PREDICTED: putative uncharacterized protein DDB_G0294196 iso
>gi|1036802572|ref|XP_017136630.1| PREDICTED: phosrestin-1 [Drosophila miranda]
>gi|1036802609|ref|XP_017136632.1| PREDICTED: peroxisomal targeting signal 2 receptor [Drosophi
>gi|1036802590|ref|XP_017136631.1| PREDICTED: protein hairy [Drosophila miranda]
>gi|1036761020|ref|XP_017156932.1| PREDICTED: selenide, water dikinase-like [Drosophila miranda
>gi|1036802110|ref|XP_017136607.1| PREDICTED: signal recognition particle receptor subunit beta
>gi|1036802129|ref|XP_017136608.1| PREDICTED: tail-anchored protein insertion receptor WRB [Dro
>gi|1036802057|ref|XP_017136606.1| PREDICTED: erythroid differentiation-related factor 1 [Droso
>gi|1036802147|ref|XP_017136609.1| PREDICTED: chorion protein S18 [Drosophila miranda]
>gi|1036766994|ref|XP_017134606.1| PREDICTED: chorion protein S15 [Drosophila miranda]
>gi|1036766884|ref|XP_017134600.1| PREDICTED: chorion protein S19 [Drosophila miranda]
>gi|1036766406|ref|XP_017134571.1| PREDICTED: chorion protein S16 [Drosophila miranda]
>gi|1036836335|ref|XP_017138503.1| PREDICTED: paramyosin, long form isoform X1 [Drosophila mira
>gi|1036836354|ref|XP_017138505.1| PREDICTED: paramyosin, short form isoform X2 [Drosophila mi
>gi|1036836373|ref|XP_017138506.1| PREDICTED: uncharacterized protein LOC108153198 [Drosophila
>gi|1036836392|ref|XP_017138507.1| PREDICTED: uncharacterized protein LOC108153198 [Drosophila
>gi|1036836411|ref|XP_017138508.1| PREDICTED: uncharacterized protein LOC108153198 [Drosophila
>gi|1036809164|ref|XP_017136995.1| PREDICTED: uncharacterized protein LOC108152288 isoform X1
>gi|1036809221|ref|XP_017136997.1| PREDICTED: uncharacterized protein LOC108152288 isoform X3
>gi|1036809200|ref|XP_017136996.1| PREDICTED: uncharacterized protein LOC108152288 isoform X2
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>gi|1036809297|ref|XP_017137000.1| PREDICTED: uncharacterized protein LOC108152289 [Drosophila
>gi|1036809242|ref|XP_017136998.1| PREDICTED: FH1/FH2 domain-containing protein 1 isoform X4 [I
>gi|1036809278|ref|XP_017136999.1| PREDICTED: FH1/FH2 domain-containing protein 3 isoform X5 [I
>gi|1036809316|ref|XP_017137001.1| PREDICTED: acyl-CoA-binding protein [Drosophila miranda]
>gi|1036761038|ref|XP_017156933.1| PREDICTED: uncharacterized protein LOC108165406 [Drosophila
>gi|1036761052|ref|XP_017156934.1| PREDICTED: thioredoxin-related transmembrane protein 1-like
>gi|1036764149|ref|XP_017157103.1| PREDICTED: uncharacterized protein LOC108165540 [Drosophila
>gi|1036761068|ref|XP_017156935.1| PREDICTED: mucin-5AC-like [Drosophila miranda]
>gi|1036847066|ref|XP_017139094.1| PREDICTED: uncharacterized protein LOC108153539 [Drosophila
>gi|1036847047|ref|XP_017139092.1| PREDICTED: spore coat protein SP96 [Drosophila miranda]
>gi|1036850895|ref|XP_017139316.1| PREDICTED: uncharacterized protein LOC108153700 [Drosophila
>gi|1036850913|ref|XP_017139317.1| PREDICTED: uncharacterized protein LOC108153701 [Drosophila
>gi|1036816630|ref|XP_017137405.1| PREDICTED: division abnormally delayed protein [Drosophila r
>gi|1036816611|ref|XP_017137404.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036826844|ref|XP_017137968.1| PREDICTED: DNA replication licensing factor Mcm7 [Drosophila
>gi|1036826825|ref|XP_017137967.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036826806|ref|XP_017137966.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036826788|ref|XP_017137965.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036849397|ref|XP_017139226.1| PREDICTED: fer-1-like protein 6 [Drosophila miranda]
>gi|1036836244|ref|XP_017138498.1| PREDICTED: tetraspanin-9 [Drosophila miranda]
>gi|1036836262|ref|XP_017138499.1| PREDICTED: tetraspanin-9 [Drosophila miranda]
>gi|1036783249|ref|XP_017135537.1| PREDICTED: peptidyl-prolyl cis-trans isomerase D [Drosophila
>gi|1036783174|ref|XP_017135533.1| PREDICTED: protein furry isoform X7 [Drosophila miranda]
>gi|1036783155|ref|XP_017135532.1| PREDICTED: protein furry isoform X6 [Drosophila miranda]
>gi|1036783117|ref|XP_017135530.1| PREDICTED: protein furry isoform X4 [Drosophila miranda]
>gi|1036783099|ref|XP_017135529.1| PREDICTED: protein furry isoform X3 [Drosophila miranda]
>gi|1036783083|ref|XP_017135527.1| PREDICTED: protein furry isoform X2 [Drosophila miranda]
>gi|1036783068|ref|XP_017135526.1| PREDICTED: protein furry isoform X1 [Drosophila miranda]
>gi|1036783212|ref|XP_017135535.1| PREDICTED: protein furry isoform X9 [Drosophila miranda]
>gi|1036783136|ref|XP_017135531.1| PREDICTED: protein furry isoform X5 [Drosophila miranda]
>gi|1036783193|ref|XP_017135534.1| PREDICTED: protein furry isoform X8 [Drosophila miranda]
>gi|1036783266|ref|XP_017135538.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036783286|ref|XP_017135539.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036783230|ref|XP_017135536.1| PREDICTED: uncharacterized protein LOC108151437 [Drosophila
>gi|1036830828|ref|XP_017138198.1| PREDICTED: UPF0046 protein C25E10.12 [Drosophila miranda]
>gi|1036830809|ref|XP_017138197.1| PREDICTED: tubulin alpha-4 chain [Drosophila miranda]
>gi|1036830847|ref|XP_017138199.1| PREDICTED: uncharacterized protein LOC108152984 [Drosophila
>gi|1036817059|ref|XP_017137429.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X4
>gi|1036817040|ref|XP_017137428.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X3
>gi|1036817098|ref|XP_017137431.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X6
>gi|1036817079|ref|XP_017137430.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X5
>gi|1036817021|ref|XP_017137427.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X2
>gi|1036817003|ref|XP_017137426.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X1
>gi|1036817135|ref|XP_017137434.1| PREDICTED: sperm flagellar protein 1-like [Drosophila mirand
>gi|1036817117|ref|XP_017137432.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa [Drosophil
>gi|1036816983|ref|XP_017137425.1| PREDICTED: uncharacterized protein KIAA1841 homolog [Drosoph
>gi|1036834860|ref|XP_017138416.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
>gi|1036834823|ref|XP_017138414.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
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>gi|1036834768|ref|XP_017138410.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
>gi|1036834841|ref|XP_017138415.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
>gi|1036834787|ref|XP_017138411.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
>gi|1036834879|ref|XP_017138417.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
>gi|1036834806|ref|XP_017138413.1| PREDICTED: eukaryotic translation initiation factor 4E isofo
>gi|1036834898|ref|XP_017138418.1| PREDICTED: sentrin-specific protease 1-like [Drosophila mira
>gi|1036828817|ref|XP_017138081.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1036803828|ref|XP_017136700.1| PREDICTED: tyrosine-protein kinase Abl isoform X1 [Drosophil
>gi|1036803901|ref|XP_017136704.1| PREDICTED: tyrosine-protein kinase Abl isoform X5 [Drosophil
>gi|1036803862|ref|XP_017136702.1| PREDICTED: tyrosine-protein kinase Abl isoform X3 [Drosophil
>gi|1036803844|ref|XP_017136701.1| PREDICTED: tyrosine-protein kinase Abl isoform X2 [Drosophil
>gi|1036803881|ref|XP_017136703.1| PREDICTED: tyrosine-protein kinase Abl isoform X4 [Drosophil
>gi|1036819929|ref|XP_017137587.1| PREDICTED: protein Mo25 [Drosophila miranda]
>gi|1036819966|ref|XP_017137589.1| PREDICTED: protein anon-73B1 [Drosophila miranda]
>gi|1036819947|ref|XP_017137588.1| PREDICTED: prostaglandin E synthase 2 [Drosophila miranda]
>gi|1036820833|ref|XP_017137632.1| PREDICTED: ribosome biogenesis protein BRX1 homolog [Drosoph
>gi|1036820815|ref|XP_017137631.1| PREDICTED: CTL-like protein 1 [Drosophila miranda]
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>gi|1036835079|ref|XP_017138429.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036835098|ref|XP_017138430.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036835116|ref|XP_017138431.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036835134|ref|XP_017138432.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036828909|ref|XP_017138087.1| PREDICTED: V-type proton ATPase subunit e 2 [Drosophila mira
>gi|1036828894|ref|XP_017138086.1| PREDICTED: zinc finger matrin-type protein 2 [Drosophila mi
>gi|1036828876|ref|XP_017138084.1| PREDICTED: protein ABHD16A [Drosophila miranda]
>gi|1036763440|ref|XP_017157066.1| PREDICTED: uncharacterized protein LOC108165519 isoform X2
>gi|1036763425|ref|XP_017157065.1| PREDICTED: uncharacterized protein LOC108165519 isoform X1
>gi|1036761085|ref|XP_017156937.1| PREDICTED: uncharacterized protein LOC108165409 [Drosophila
>gi|1036780681|ref|XP_017135386.1| PREDICTED: NF-X1-type zinc finger protein NFXL1 [Drosophila
>gi|1036780700|ref|XP_017135387.1| PREDICTED: PQ-loop repeat-containing protein 3 [Drosophila m
>gi|1036780662|ref|XP_017135385.1| PREDICTED: F-box/WD repeat-containing protein 7 [Drosophila
>gi|1036780718|ref|XP_017135388.1| PREDICTED: kinesin-like protein KIF23 [Drosophila miranda]
>gi|1036843149|ref|XP_017138886.1| PREDICTED: sulfide:quinone oxidoreductase, mitochondrial [Dr
>gi|1036785666|ref|XP_017135677.1| PREDICTED: neural/ectodermal development factor IMP-L2 isofo
>gi|1036785685|ref|XP_017135678.1| PREDICTED: neural/ectodermal development factor IMP-L2 isofo
>gi|1036785647|ref|XP_017135676.1| PREDICTED: neurobeachin-like protein 1 isoform X2 [Drosophil
>gi|1036785628|ref|XP_017135675.1| PREDICTED: neurobeachin-like protein 1 isoform X1 [Drosophil
>gi|1036785704|ref|XP_017135679.1| PREDICTED: larval cuticle protein A3A [Drosophila miranda]
>gi|1036785742|ref|XP_017135682.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1036785759|ref|XP_017135683.1| PREDICTED: larval cuticle protein A3A [Drosophila miranda]
>gi|1036785723|ref|XP_017135681.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1036758120|ref|XP_017156758.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036807493|ref|XP_017136904.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807512|ref|XP_017136905.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807531|ref|XP_017136906.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807552|ref|XP_017136907.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807644|ref|XP_017136912.1| PREDICTED: V-type proton ATPase subunit e [Drosophila miran
>gi|1036807571|ref|XP_017136908.1| PREDICTED: uncharacterized protein LOC108152231 [Drosophila
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>gi|1036807626|ref|XP_017136911.1| PREDICTED: protein tipE isoform X2 [Drosophila miranda]
>gi|1036807607|ref|XP_017136910.1| PREDICTED: protein tipE isoform X1 [Drosophila miranda]
>gi|1036807407|ref|XP_017136899.1| PREDICTED: uncharacterized protein LOC108152228 isoform X1
>gi|1036807426|ref|XP_017136900.1| PREDICTED: uncharacterized protein LOC108152228 isoform X2
>gi|1036761102|ref|XP_017156938.1| PREDICTED: UPF0769 protein CG18675 [Drosophila miranda]
>gi|1036807444|ref|XP_017136901.1| PREDICTED: ero1-like protein [Drosophila miranda]
>gi|1036807460|ref|XP_017136903.1| PREDICTED: ero1-like protein [Drosophila miranda]
>gi|1036838500|ref|XP_017138633.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036848898|ref|XP_017139197.1| PREDICTED: NGFI-A-binding protein homolog isoform X1 [Drosop
>gi|1036848914|ref|XP_017139198.1| PREDICTED: NGFI-A-binding protein homolog isoform X2 [Drosop
>gi|1036821619|ref|XP_017137674.1| PREDICTED: venom serine protease Bi-VSP isoform X2 [Drosophi
>gi|1036821656|ref|XP_017137677.1| PREDICTED: venom protease isoform X4 [Drosophila miranda]
>gi|1036821637|ref|XP_017137676.1| PREDICTED: venom protease isoform X3 [Drosophila miranda]
>gi|1036821600|ref|XP_017137673.1| PREDICTED: venom serine protease Bi-VSP isoform X1 [Drosophi
>gi|1036821757|ref|XP_017137682.1| PREDICTED: venom protease isoform X2 [Drosophila miranda]
>gi|1036821734|ref|XP_017137681.1| PREDICTED: proclotting enzyme isoform X1 [Drosophila miranda
>gi|1036821715|ref|XP_017137680.1| PREDICTED: adipokinetic hormone [Drosophila miranda]
>gi|1036821694|ref|XP_017137679.1| PREDICTED: ras-like protein 2 [Drosophila miranda]
>gi|1036821675|ref|XP_017137678.1| PREDICTED: replication factor C subunit 2 [Drosophila miran
>gi|1036804292|ref|XP_017136727.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036804368|ref|XP_017136731.1| PREDICTED: PHD finger protein rhinoceros isoform X5 [Drosoph
>gi|1036804424|ref|XP_017136734.1| PREDICTED: ensconsin isoform X8 [Drosophila miranda]
>gi|1036804460|ref|XP_017136737.1| PREDICTED: ensconsin isoform X10 [Drosophila miranda]
>gi|1036804349|ref|XP_017136730.1| PREDICTED: flocculation protein FLO11 isoform X4 [Drosophila
>gi|1036804405|ref|XP_017136733.1| PREDICTED: ensconsin isoform X7 [Drosophila miranda]
>gi|1036804386|ref|XP_017136732.1| PREDICTED: protein split ends isoform X6 [Drosophila miranda
>gi|1036804443|ref|XP_017136736.1| PREDICTED: ensconsin isoform X9 [Drosophila miranda]
>gi|1036804498|ref|XP_017136739.1| PREDICTED: ensconsin isoform X12 [Drosophila miranda]
>gi|1036804311|ref|XP_017136728.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036804479|ref|XP_017136738.1| PREDICTED: ensconsin isoform X11 [Drosophila miranda]
>gi|1036804515|ref|XP_017136740.1| PREDICTED: MAP7 domain-containing protein 1 isoform X13 [Dro
>gi|1036804534|ref|XP_017136741.1| PREDICTED: ensconsin isoform X14 [Drosophila miranda]
>gi|1036804330|ref|XP_017136729.1| PREDICTED: flocculation protein FLO11 isoform X3 [Drosophila
>gi|1036804553|ref|XP_017136742.1| PREDICTED: ensconsin isoform X15 [Drosophila miranda]
>gi|1036835461|ref|XP_017138451.1| PREDICTED: myophilin isoform X1 [Drosophila miranda]
>gi|1036835480|ref|XP_017138452.1| PREDICTED: myophilin isoform X2 [Drosophila miranda]
>gi|1036796737|ref|XP_017136299.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036796718|ref|XP_017136298.1| PREDICTED: uncharacterized protein F23B12.7 [Drosophila mir
>gi|1036796062|ref|XP_017136263.1| PREDICTED: SPRY domain-containing SOCS box protein 3 [Drosop
>gi|1036796043|ref|XP_017136262.1| PREDICTED: ATP-dependent helicase brm isoform X2 [Drosophila
>gi|1036796025|ref|XP_017136261.1| PREDICTED: ATP-dependent helicase brm isoform X1 [Drosophila
>gi|1036796102|ref|XP_017136265.1| PREDICTED: inositol monophosphatase 1 [Drosophila miranda]
>gi|1036796142|ref|XP_017136268.1| PREDICTED: inositol monophosphatase 2 [Drosophila miranda]
>gi|1036796123|ref|XP_017136267.1| PREDICTED: inositol monophosphatase ttx-7 [Drosophila miran
>gi|1036796081|ref|XP_017136264.1| PREDICTED: inositol monophosphatase 2 [Drosophila miranda]
>gi|1036819118|ref|XP_017137545.1| PREDICTED: ADP-ribosylation factor-like protein 1 [Drosophi
>gi|1036819099|ref|XP_017137544.1| PREDICTED: DNA polymerase delta catalytic subunit [Drosophil
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>gi|1036821795|ref|XP_017137684.1| PREDICTED: palmitoyltransferase Hip14 [Drosophila miranda]
>gi|1036831960|ref|XP_017138261.1| PREDICTED: carboxy-terminal domain RNA polymerase II polypep
>gi|1036831979|ref|XP_017138262.1| PREDICTED: carboxy-terminal domain RNA polymerase II polypep
>gi|1036822269|ref|XP_017137711.1| PREDICTED: 28S ribosomal protein S31, mitochondrial [Drosoph
>gi|1036822250|ref|XP_017137710.1| PREDICTED: E3 ubiquitin-protein ligase mind-bomb [Drosophila
>gi|1036826203|ref|XP_017137930.1| PREDICTED: uncharacterized protein LOC108152836 isoform X2
>gi|1036826180|ref|XP_017137929.1| PREDICTED: uncharacterized protein LOC108152836 isoform X1
>gi|1036761120|ref|XP_017156939.1| PREDICTED: cytosol aminopeptidase-like, partial [Drosophila
>gi|1036761138|ref|XP_017156940.1| PREDICTED: isocitrate dehydrogenase [NAD] catalytic subunit
>gi|1036826259|ref|XP_017137934.1| PREDICTED: hydroxyacylglutathione hydrolase, mitochondrial
>gi|1036826239|ref|XP_017137933.1| PREDICTED: hydroxyacylglutathione hydrolase, mitochondrial
>gi|1036826278|ref|XP_017137935.1| PREDICTED: hydroxyacylglutathione hydrolase, mitochondrial
>gi|1036826221|ref|XP_017137931.1| PREDICTED: uncharacterized protein LOC108152837 [Drosophila
>gi|1036796788|ref|XP_017136303.1| PREDICTED: homeobox protein SIX6 [Drosophila miranda]
>gi|1036796838|ref|XP_017136306.1| PREDICTED: uncharacterized protein LOC108151906 isoform X2
>gi|1036796821|ref|XP_017136305.1| PREDICTED: uncharacterized protein LOC108151906 isoform X1
>gi|1036796771|ref|XP_017136301.1| PREDICTED: trafficking protein particle complex subunit 12
>gi|1036796754|ref|XP_017136300.1| PREDICTED: phosphorylated CTD-interacting factor 1 [Drosoph
>gi|1036796804|ref|XP_017136304.1| PREDICTED: peroxisomal membrane protein PEX14 [Drosophila mi
>gi|1036788984|ref|XP_017135863.1| PREDICTED: peroxisomal membrane protein PEX16 [Drosophila mi
>gi|1036788931|ref|XP_017135860.1| PREDICTED: serine/threonine-protein kinase PITSLRE [Drosoph
>gi|1036789059|ref|XP_017135868.1| PREDICTED: ATP synthase subunit s-like protein [Drosophila
>gi|1036789097|ref|XP_017135870.1| PREDICTED: coiled-coil-helix-coiled-coil-helix domain-conta
>gi|1036789003|ref|XP_017135864.1| PREDICTED: Down syndrome critical region protein 3 homolog
>gi|1036788947|ref|XP_017135861.1| PREDICTED: uncharacterized protein LOC108151653 [Drosophila
>gi|1036788966|ref|XP_017135862.1| PREDICTED: uncharacterized protein LOC108151653 [Drosophila
>gi|1036789116|ref|XP_017135871.1| PREDICTED: putative gustatory receptor 77a [Drosophila mira
>gi|1036789021|ref|XP_017135866.1| PREDICTED: uncharacterized protein LOC108151657 [Drosophila
>gi|1036789040|ref|XP_017135867.1| PREDICTED: uncharacterized protein LOC108151657 [Drosophila
>gi|1036789078|ref|XP_017135869.1| PREDICTED: uncharacterized protein LOC108151659 [Drosophila
>gi|1036843582|ref|XP_017138913.1| PREDICTED: uncharacterized protein LOC108153421 [Drosophila
>gi|1036761158|ref|XP_017156941.1| PREDICTED: extracellular sulfatase SULF-1 homolog [Drosophil
>gi|1036843564|ref|XP_017138911.1| PREDICTED: zygotic gap protein knirps [Drosophila miranda]
>gi|1036761178|ref|XP_017156942.1| PREDICTED: uncharacterized protein LOC108165414 [Drosophila
>gi|1036832902|ref|XP_017138313.1| PREDICTED: knirps-related protein [Drosophila miranda]
>gi|1036829827|ref|XP_017138138.1| PREDICTED: cell wall protein IFF6 [Drosophila miranda]
>gi|1036829863|ref|XP_017138140.1| PREDICTED: uncharacterized protein LOC108152952 [Drosophila
>gi|1036829844|ref|XP_017138139.1| PREDICTED: cytosolic Fe-S cluster assembly factor NUBP2 hom
>gi|1036785395|ref|XP_017135661.1| PREDICTED: phosphatidylserine synthase 1 [Drosophila miranda
>gi|1036785483|ref|XP_017135666.1| PREDICTED: vesicle transport protein SFT2A [Drosophila mira
>gi|1036785433|ref|XP_017135663.1| PREDICTED: transcription termination factor 3, mitochondrial
>gi|1036785447|ref|XP_017135664.1| PREDICTED: uncharacterized protein LOC108151526 isoform X1
>gi|1036785464|ref|XP_017135665.1| PREDICTED: uncharacterized protein LOC108151526 isoform X2
>gi|1036785377|ref|XP_017135660.1| PREDICTED: uncharacterized protein LOC108151520 isoform X3
>gi|1036785340|ref|XP_017135658.1| PREDICTED: uncharacterized protein LOC108151520 isoform X1
>gi|1036785358|ref|XP_017135659.1| PREDICTED: C2 domain-containing protein 3 isoform X2 [Drosop
>gi|1036785414|ref|XP_017135662.1| PREDICTED: hippocampus abundant transcript-like protein 1 [
>gi|1036839101|ref|XP_017138671.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
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>gi|1036800954|ref|XP_017136539.1| PREDICTED: zinc transporter 1 [Drosophila miranda]
>gi|1036800992|ref|XP_017136541.1| PREDICTED: glutaminyl-peptide cyclotransferase isoform X1 [
>gi|1036801010|ref|XP_017136543.1| PREDICTED: glutaminyl-peptide cyclotransferase isoform X2 [
>gi|1036801045|ref|XP_017136545.1| PREDICTED: vacuolar ATPase assembly integral membrane prote
>gi|1036801026|ref|XP_017136544.1| PREDICTED: uncharacterized protein LOC108152051 [Drosophila
>gi|1036800973|ref|XP_017136540.1| PREDICTED: L-threonine 3-dehydrogenase, mitochondrial [Dros
>gi|1036839049|ref|XP_017138669.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1036839009|ref|XP_017138667.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1036839028|ref|XP_017138668.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1036829250|ref|XP_017138107.1| PREDICTED: uncharacterized protein LOC108152931 [Drosophila
>gi|1036829273|ref|XP_017138108.1| PREDICTED: uncharacterized protein LOC108152931 [Drosophila
>gi|1036829292|ref|XP_017138109.1| PREDICTED: uncharacterized protein LOC108152932 [Drosophila
>gi|1036829311|ref|XP_017138110.1| PREDICTED: uncharacterized protein LOC108152932 [Drosophila
>gi|1036829328|ref|XP_017138112.1| PREDICTED: uncharacterized protein LOC108152932 [Drosophila
>gi|1036793663|ref|XP_017136126.1| PREDICTED: 4-hydroxyphenylpyruvate dioxygenase [Drosophila r
>gi|1036793624|ref|XP_017136124.1| PREDICTED: uncharacterized protein LOC108151787 isoform X1
>gi|1036793643|ref|XP_017136125.1| PREDICTED: uncharacterized protein LOC108151787 isoform X2
>gi|1036793682|ref|XP_017136127.1| PREDICTED: 39S ribosomal protein L15, mitochondrial [Drosopl
>gi|1036793605|ref|XP_017136123.1| PREDICTED: presenilin homolog isoform X3 [Drosophila miranda
>gi|1036793587|ref|XP_017136122.1| PREDICTED: presenilin homolog isoform X2 [Drosophila miranda
>gi|1036793572|ref|XP_017136121.1| PREDICTED: presenilin homolog isoform X1 [Drosophila miranda
>gi|1036793501|ref|XP_017136117.1| PREDICTED: Golgi apparatus protein 1 isoform X1 [Drosophila
>gi|1036793519|ref|XP_017136118.1| PREDICTED: Golgi apparatus protein 1 isoform X2 [Drosophila
>gi|1036793536|ref|XP_017136119.1| PREDICTED: glutactin isoform X1 [Drosophila miranda]
>gi|1036793555|ref|XP_017136120.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1036831997|ref|XP_017138264.1| PREDICTED: nuclear hormone receptor HR78 isoform X1 [Drosopl
>gi|1036832015|ref|XP_017138265.1| PREDICTED: nuclear hormone receptor HR78 isoform X2 [Drosopl
>gi|1036832053|ref|XP_017138267.1| PREDICTED: protein PBDC1 [Drosophila miranda]
>gi|1036832034|ref|XP_017138266.1| PREDICTED: uncharacterized protein LOC108153023 [Drosophila
>gi|1036800160|ref|XP_017136494.1| PREDICTED: uncharacterized protein LOC108152024 [Drosophila
>gi|1036800068|ref|XP_017136489.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800085|ref|XP_017136490.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800104|ref|XP_017136491.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800123|ref|XP_017136492.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800142|ref|XP_017136493.1| PREDICTED: speract receptor-like isoform X2 [Drosophila mira
>gi|1036813202|ref|XP_017137208.1| PREDICTED: mucolipin-3 [Drosophila miranda]
>gi|1036813276|ref|XP_017137213.1| PREDICTED: uncharacterized protein LOC108152411 [Drosophila
>gi|1036813220|ref|XP_017137209.1| PREDICTED: uncharacterized protein LOC108152410 isoform X1
>gi|1036813238|ref|XP_017137211.1| PREDICTED: uncharacterized protein LOC108152410 isoform X2
>gi|1036813257|ref|XP_017137212.1| PREDICTED: uncharacterized protein LOC108152410 isoform X3
>gi|1036813295|ref|XP_017137214.1| PREDICTED: guanine nucleotide-binding protein subunit beta-2
>gi|1036835535|ref|XP_017138455.1| PREDICTED: uncharacterized protein LOC108153162 [Drosophila
>gi|1036848474|ref|XP_017139172.1| PREDICTED: protein Teyrha-meyrha isoform X1 [Drosophila mira
>gi|1036848551|ref|XP_017139176.1| PREDICTED: protein Teyrha-meyrha isoform X5 [Drosophila mira
>gi|1036848493|ref|XP_017139173.1| PREDICTED: protein Teyrha-meyrha isoform X2 [Drosophila mira
>gi|1036848567|ref|XP_017139177.1| PREDICTED: protein Teyrha-meyrha isoform X6 [Drosophila mira
>gi|1036848513|ref|XP_017139174.1| PREDICTED: protein Teyrha-meyrha isoform X3 [Drosophila mira
>gi|1036848532|ref|XP_017139175.1| PREDICTED: protein Teyrha-meyrha isoform X4 [Drosophila mira
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>gi|1036848585|ref|XP_017139178.1| PREDICTED: protein Teyrha-meyrha isoform X7 [Drosophila mira
>gi|1036848603|ref|XP_017139179.1| PREDICTED: protein Teyrha-meyrha isoform X8 [Drosophila mira
>gi|1036848621|ref|XP_017139180.1| PREDICTED: protein Teyrha-meyrha isoform X9 [Drosophila mira
>gi|1036848145|ref|XP_017139153.1| PREDICTED: ornithine aminotransferase, mitochondrial [Drosop
>gi|1036766120|ref|XP_017134559.1| PREDICTED: uncharacterized protein LOC108150774 [Drosophila
>gi|1036817801|ref|XP_017137470.1| PREDICTED: E3 ubiquitin-protein ligase rnf146 isoform X1 [Dr
>gi|1036817821|ref|XP_017137471.1| PREDICTED: E3 ubiquitin-protein ligase RNF146 isoform X2 [Dr
>gi|1036761196|ref|XP_017156943.1| PREDICTED: E3 ubiquitin-protein ligase rnf146-like [Drosophi
>gi|1036783322|ref|XP_017135542.1| PREDICTED: mitogen-activated protein kinase kinase kinase 13
>gi|1036783341|ref|XP_017135543.1| PREDICTED: mitogen-activated protein kinase kinase kinase 13
>gi|1036783360|ref|XP_017135544.1| PREDICTED: mitogen-activated protein kinase kinase kinase 13
>gi|1036783379|ref|XP_017135545.1| PREDICTED: uncharacterized protein LOC108151444 [Drosophila
>gi|1036783417|ref|XP_017135547.1| PREDICTED: histone chaperone asf1 [Drosophila miranda]
>gi|1036783304|ref|XP_017135540.1| PREDICTED: trafficking protein particle complex subunit 8 [
>gi|1036783398|ref|XP_017135546.1| PREDICTED: glycosylphosphatidylinositol anchor biosynthesis
>gi|1036777143|ref|XP_017135174.1| PREDICTED: LOW QUALITY PROTEIN: lon protease homolog, mitoch
>gi|1036777215|ref|XP_017135178.1| PREDICTED: venom protease [Drosophila miranda]
>gi|1036777268|ref|XP_017135181.1| PREDICTED: uncharacterized protein LOC108151213 isoform X2
>gi|1036777249|ref|XP_017135180.1| PREDICTED: uncharacterized protein LOC108151213 isoform X1
>gi|1036777286|ref|XP_017135183.1| PREDICTED: general odorant-binding protein lush [Drosophila
>gi|1036777180|ref|XP_017135176.1| PREDICTED: transcription initiation factor TFIID subunit 6
>gi|1036777198|ref|XP_017135177.1| PREDICTED: transcription initiation factor TFIID subunit 6
>gi|1036777162|ref|XP_017135175.1| PREDICTED: transcription initiation factor TFIID subunit 6
>gi|1036777104|ref|XP_017135172.1| PREDICTED: histone-lysine N-methyltransferase ash1 isoform
>gi|1036777123|ref|XP_017135173.1| PREDICTED: histone-lysine N-methyltransferase ash1 isoform
>gi|1036777230|ref|XP_017135179.1| PREDICTED: proteasome subunit beta type-1-like [Drosophila
>gi|1036810076|ref|XP_017137043.1| PREDICTED: rRNA methyltransferase 3, mitochondrial [Drosoph
>gi|1036809980|ref|XP_017137038.1| PREDICTED: ATP-binding cassette sub-family F member 3 [Dros
>gi|1036810094|ref|XP_017137044.1| PREDICTED: UPF0389 protein GA21628 [Drosophila miranda]
>gi|1036809999|ref|XP_017137039.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036810017|ref|XP_017137040.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036810038|ref|XP_017137041.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036810057|ref|XP_017137042.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036775510|ref|XP_017135080.1| PREDICTED: uncharacterized protein LOC108151161 [Drosophila
>gi|1036775416|ref|XP_017135075.1| PREDICTED: E3 ubiquitin-protein ligase listerin [Drosophila
>gi|1036775624|ref|XP_017135086.1| PREDICTED: uncharacterized protein LOC108151168 [Drosophila
>gi|1036775434|ref|XP_017135076.1| PREDICTED: dynactin subunit 1 [Drosophila miranda]
>gi|1036775453|ref|XP_017135077.1| PREDICTED: fibroin heavy chain [Drosophila miranda]
>gi|1036775472|ref|XP_017135078.1| PREDICTED: fibroin heavy chain [Drosophila miranda]
>gi|1036761214|ref|XP_017156944.1| PREDICTED: LOW QUALITY PROTEIN: MICOS complex subunit Mic10-
>gi|1036775548|ref|XP_017135082.1| PREDICTED: uncharacterized protein LOC108151163 [Drosophila
>gi|1036758138|ref|XP_017156759.1| PREDICTED: adult-specific cuticular protein ACP-20 [Drosoph
>gi|1036775645|ref|XP_017135087.1| PREDICTED: cuticle protein 19.8 [Drosophila miranda]
>gi|1036775567|ref|XP_017135083.1| PREDICTED: venom acid phosphatase Acph-1 [Drosophila miranda
>gi|1036775586|ref|XP_017135084.1| PREDICTED: venom acid phosphatase Acph-1 [Drosophila miranda
>gi|1036775605|ref|XP_017135085.1| PREDICTED: venom acid phosphatase Acph-1 [Drosophila miranda
>gi|1036775529|ref|XP_017135081.1| PREDICTED: sulfhydryl oxidase 1-like [Drosophila miranda]
>gi|1036775491|ref|XP_017135079.1| PREDICTED: polycystin-2 [Drosophila miranda]
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>gi|1036761232|ref|XP_017156945.1| PREDICTED: glutaconyl-CoA decarboxylase subunit gamma-like
>gi|1036758155|ref|XP_017156760.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036761251|ref|XP_017156946.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036761270|ref|XP_017156948.1| PREDICTED: cyclin-dependent kinase inhibitor 1C-like [Droso
>gi|1036761287|ref|XP_017156949.1| PREDICTED: cyclin-dependent kinase inhibitor 1C [Drosophila
>gi|1036849064|ref|XP_017139207.1| PREDICTED: uncharacterized protein LOC108153617 [Drosophila
>gi|1036820061|ref|XP_017137595.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036820003|ref|XP_017137592.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036820042|ref|XP_017137594.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036819985|ref|XP_017137590.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036820023|ref|XP_017137593.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036758171|ref|XP_017156761.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [D
>gi|1036820134|ref|XP_017137599.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [D
>gi|1036820171|ref|XP_017137601.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [D
>gi|1036820153|ref|XP_017137600.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like, p
>gi|1036820116|ref|XP_017137598.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [D
>gi|1036820080|ref|XP_017137596.1| PREDICTED: deoxynucleoside triphosphate triphosphohydrolase
>gi|1036820098|ref|XP_017137597.1| PREDICTED: deoxynucleoside triphosphate triphosphohydrolase
>gi|1036833503|ref|XP_017138348.1| PREDICTED: glycogen-binding subunit 76A [Drosophila miranda]
>gi|1036833520|ref|XP_017138349.1| PREDICTED: glycogen-binding subunit 76A [Drosophila miranda]
>gi|1036833538|ref|XP_017138350.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036761305|ref|XP_017156950.1| PREDICTED: collagen alpha-1(IX) chain [Drosophila miranda]
>gi|1036770105|ref|XP_017134775.1| PREDICTED: uncharacterized protein LOC108150952 [Drosophila
>gi|1036761324|ref|XP_017156951.1| PREDICTED: uncharacterized protein LOC108165423 [Drosophila
>gi|1036798272|ref|XP_017136387.1| PREDICTED: putative aldehyde dehydrogenase family 7 member A
>gi|1036798383|ref|XP_017136394.1| PREDICTED: methyltransferase-like protein 5 [Drosophila mira
>gi|1036798457|ref|XP_017136399.1| PREDICTED: SOSS complex subunit C homolog [Drosophila mirand
>gi|1036798402|ref|XP_017136395.1| PREDICTED: protein max isoform X1 [Drosophila miranda]
>gi|1036798420|ref|XP_017136396.1| PREDICTED: protein max isoform X2 [Drosophila miranda]
>gi|1036798438|ref|XP_017136398.1| PREDICTED: protein max isoform X3 [Drosophila miranda]
>gi|1036798475|ref|XP_017136400.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798494|ref|XP_017136401.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798512|ref|XP_017136402.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798529|ref|XP_017136403.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798547|ref|XP_017136404.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798311|ref|XP_017136389.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798328|ref|XP_017136390.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798345|ref|XP_017136391.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798364|ref|XP_017136393.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798292|ref|XP_017136388.1| PREDICTED: ruvB-like helicase 2 [Drosophila miranda]
>gi|1036844935|ref|XP_017138979.1| PREDICTED: 39S ribosomal protein L21, mitochondrial [Drosoph
>gi|1036768023|ref|XP_017134659.1| PREDICTED: uncharacterized protein LOC108150867 [Drosophila
>gi|1036851745|ref|XP_017139367.1| PREDICTED: protein IWS1 homolog [Drosophila miranda]
>gi|1036851762|ref|XP_017139368.1| PREDICTED: uncharacterized protein LOC108153749 [Drosophila
>gi|1036761343|ref|XP_017156952.1| PREDICTED: uncharacterized protein LOC108165425 [Drosophila
>gi|1036815846|ref|XP_017137358.1| PREDICTED: frizzled-2 [Drosophila miranda]
>gi|1036815864|ref|XP_017137359.1| PREDICTED: putative glycine-rich cell wall structural protei
>gi|1036834233|ref|XP_017138382.1| PREDICTED: serpin B6-like isoform X1 [Drosophila miranda]
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>gi|1036834252|ref|XP_017138383.1| PREDICTED: serpin B3-like isoform X2 [Drosophila miranda]
>gi|1036761362|ref|XP_017156953.1| PREDICTED: LOW QUALITY PROTEIN: accessory gland protein Acp⁊
>gi|1036834214|ref|XP_017138381.1| PREDICTED: uncharacterized protein LOC108153106 [Drosophila
>gi|1036823262|ref|XP_017137768.1| PREDICTED: UPF0489 protein C5orf22 homolog [Drosophila miraᵣ
>gi|1036823318|ref|XP_017137772.1| PREDICTED: protein canopy 4 [Drosophila miranda]
>gi|1036823282|ref|XP_017137770.1| PREDICTED: uncharacterized protein LOC108152736 [Drosophila
>gi|1036823300|ref|XP_017137771.1| PREDICTED: protein Gemin2 [Drosophila miranda]
>gi|1036769533|ref|XP_017134745.1| PREDICTED: dual specificity protein phosphatase Mpk3 [Drosoᵣ
>gi|1036769551|ref|XP_017134746.1| PREDICTED: sideroflexin-2 [Drosophila miranda]
>gi|1036769474|ref|XP_017134741.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1036769359|ref|XP_017134735.1| PREDICTED: protein naked cuticle isoform X1 [Drosophila miraᵣ
>gi|1036769380|ref|XP_017134736.1| PREDICTED: protein naked cuticle isoform X2 [Drosophila miraᵣ
>gi|1036769397|ref|XP_017134737.1| PREDICTED: protein naked cuticle isoform X3 [Drosophila miraᵣ
>gi|1036769515|ref|XP_017134743.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036769497|ref|XP_017134742.1| PREDICTED: tRNA (uracil-5-)-methyltransferase homolog A [Drᵣ
>gi|1036769456|ref|XP_017134740.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 iᵣ
>gi|1036769435|ref|XP_017134739.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 iᵣ
>gi|1036769571|ref|XP_017134747.1| PREDICTED: uncharacterized protein LOC108150927 [Drosophila
>gi|1036769416|ref|XP_017134738.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [ᵣ
>gi|1036769590|ref|XP_017134748.1| PREDICTED: uncharacterized protein LOC108150928 [Drosophila
>gi|1036845930|ref|XP_017139029.1| PREDICTED: organic solute transporter alpha-like protein [Dᵣ
>gi|1036845949|ref|XP_017139030.1| PREDICTED: organic solute transporter alpha-like protein [Dᵣ
>gi|1036845965|ref|XP_017139031.1| PREDICTED: organic solute transporter alpha-like protein [Dᵣ
>gi|1036845984|ref|XP_017139032.1| PREDICTED: organic solute transporter alpha-like protein [Dᵣ
>gi|1036846003|ref|XP_017139033.1| PREDICTED: uncharacterized protein LOC108153494 [Drosophila
>gi|1036846020|ref|XP_017139034.1| PREDICTED: uncharacterized protein LOC108153494 [Drosophila
>gi|1036846037|ref|XP_017139035.1| PREDICTED: uncharacterized protein LOC108153494 [Drosophila
>gi|1036801139|ref|XP_017136551.1| PREDICTED: uncharacterized protein LOC108152057 [Drosophila
>gi|1036761383|ref|XP_017156954.1| PREDICTED: mite allergen Eur m 3 [Drosophila miranda]
>gi|1036801120|ref|XP_017136549.1| PREDICTED: uncharacterized protein LOC108152056 [Drosophila
>gi|1036761402|ref|XP_017156956.1| PREDICTED: uncharacterized protein LOC108165428 [Drosophila
>gi|1036801083|ref|XP_017136547.1| PREDICTED: pre-mRNA-processing factor 6 [Drosophila miranda]
>gi|1036801101|ref|XP_017136548.1| PREDICTED: COP9 signalosome complex subunit 1b [Drosophila ᵣ
>gi|1036801158|ref|XP_017136552.1| PREDICTED: corepressor interacting with RBPJ 1 [Drosophila ᵣ
>gi|1036833963|ref|XP_017138365.1| PREDICTED: gametocyte-specific factor 1 homolog [Drosophila
>gi|1036833981|ref|XP_017138367.1| PREDICTED: 60S ribosomal protein L26 [Drosophila miranda]
>gi|1036833998|ref|XP_017138368.1| PREDICTED: 60S ribosomal protein L26 [Drosophila miranda]
>gi|1036829091|ref|XP_017138098.1| PREDICTED: uncharacterized protein LOC108152926 [Drosophila
>gi|1036829111|ref|XP_017138099.1| PREDICTED: cell death-inducing p53-target protein 1 [Drosopʰ
>gi|1036829132|ref|XP_017138100.1| PREDICTED: cell death-inducing p53-target protein 1 [Drosopʰ
>gi|1036829153|ref|XP_017138101.1| PREDICTED: cell death-inducing p53-target protein 1 [Drosopʰ
>gi|1036831378|ref|XP_017138230.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1036831397|ref|XP_017138231.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1036831357|ref|XP_017138229.1| PREDICTED: integrator complex subunit 3 homolog isoform X1
>gi|1036803424|ref|XP_017136679.1| PREDICTED: uncharacterized protein LOC108152115 [Drosophila
>gi|1036803406|ref|XP_017136678.1| PREDICTED: vacuolar protein-sorting-associated protein 36 [ᵣ
>gi|1036826090|ref|XP_017137924.1| PREDICTED: liprin-beta-1 [Drosophila miranda]
>gi|1036825997|ref|XP_017137918.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
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>gi|1036826015|ref|XP_017137919.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826034|ref|XP_017137921.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826052|ref|XP_017137922.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826071|ref|XP_017137923.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036839502|ref|XP_017138688.1| PREDICTED: probable WRKY transcription factor protein 1 [Dro
>gi|1036839483|ref|XP_017138687.1| PREDICTED: CUGBP Elav-like family member 4 isoform X9 [Droso
>gi|1036839428|ref|XP_017138684.1| PREDICTED: CUGBP Elav-like family member 4 isoform X6 [Droso
>gi|1036839392|ref|XP_017138681.1| PREDICTED: CUGBP Elav-like family member 4 isoform X4 [Droso
>gi|1036839356|ref|XP_017138679.1| PREDICTED: CUGBP Elav-like family member 4 isoform X2 [Droso
>gi|1036839464|ref|XP_017138686.1| PREDICTED: CUGBP Elav-like family member 4 isoform X8 [Droso
>gi|1036839409|ref|XP_017138683.1| PREDICTED: CUGBP Elav-like family member 4 isoform X5 [Droso
>gi|1036839375|ref|XP_017138680.1| PREDICTED: CUGBP Elav-like family member 4 isoform X3 [Droso
>gi|1036839335|ref|XP_017138678.1| PREDICTED: CUGBP Elav-like family member 4 isoform X1 [Droso
>gi|1036839447|ref|XP_017138685.1| PREDICTED: CUGBP Elav-like family member 4 isoform X7 [Droso
>gi|1036852108|ref|XP_017139391.1| PREDICTED: uncharacterized protein LOC108153763 [Drosophila
>gi|1036767503|ref|XP_017134633.1| PREDICTED: ell-associated factor Eaf-like isoform X1 [Drosop
>gi|1036767522|ref|XP_017134634.1| PREDICTED: ell-associated factor Eaf-like isoform X1 [Drosop
>gi|1036767540|ref|XP_017134635.1| PREDICTED: mastermind-like domain-containing protein 1 isofo
>gi|1036818023|ref|XP_017137483.1| PREDICTED: ERAD-associated E3 ubiquitin-protein ligase HRD1A
>gi|1036818004|ref|XP_017137482.1| PREDICTED: protein panoramix [Drosophila miranda]
>gi|1036761420|ref|XP_017156957.1| PREDICTED: protein FAM210A-like [Drosophila miranda]
>gi|1036768739|ref|XP_017134698.1| PREDICTED: protein new-glue 2-like [Drosophila miranda]
>gi|1036768721|ref|XP_017134697.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036767427|ref|XP_017134629.1| PREDICTED: uncharacterized protein LOC108150833 [Drosophila
>gi|1036834733|ref|XP_017138408.1| PREDICTED: hemocytin [Drosophila miranda]
>gi|1036840869|ref|XP_017138766.1| PREDICTED: GPN-loop GTPase 2 [Drosophila miranda]
>gi|1036830865|ref|XP_017138201.1| PREDICTED: actin-interacting protein 1 [Drosophila miranda]
>gi|1036843908|ref|XP_017138929.1| PREDICTED: uncharacterized protein LOC108153435 [Drosophila
>gi|1036843927|ref|XP_017138930.1| PREDICTED: uncharacterized protein LOC108153435 [Drosophila
>gi|1036844003|ref|XP_017138935.1| PREDICTED: uncharacterized protein LOC108153438 [Drosophila
>gi|1036843948|ref|XP_017138931.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036843966|ref|XP_017138932.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036843984|ref|XP_017138933.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036765677|ref|XP_017134533.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036765487|ref|XP_017157174.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036844444|ref|XP_017138959.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036844463|ref|XP_017138960.1| PREDICTED: uncharacterized protein LOC108153451 [Drosophila
>gi|1036781054|ref|XP_017135410.1| PREDICTED: uncharacterized protein LOC108151362 [Drosophila
>gi|1036781147|ref|XP_017135415.1| PREDICTED: jmjC domain-containing protein 7 [Drosophila mira
>gi|1036781165|ref|XP_017135416.1| PREDICTED: GSK3-beta interaction protein [Drosophila miranda
>gi|1036781129|ref|XP_017135414.1| PREDICTED: ATP-dependent RNA helicase Ddx1 [Drosophila miran
>gi|1036781073|ref|XP_017135411.1| PREDICTED: guanine nucleotide exchange factor subunit Rich
>gi|1036781091|ref|XP_017135412.1| PREDICTED: guanine nucleotide exchange factor subunit Rich
>gi|1036781110|ref|XP_017135413.1| PREDICTED: guanine nucleotide exchange factor subunit Rich
>gi|1036820962|ref|XP_017137639.1| PREDICTED: MOXD1 homolog 2 [Drosophila miranda]
>gi|1036771162|ref|XP_017134835.1| PREDICTED: 60S acidic ribosomal protein P0 [Drosophila miran
>gi|1036771032|ref|XP_017134828.1| PREDICTED: NEDD4-binding protein 2 [Drosophila miranda]
>gi|1036771051|ref|XP_017134829.1| PREDICTED: NEDD4-binding protein 2 [Drosophila miranda]
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>gi|1036771195|ref|XP_017134838.1| PREDICTED: uncharacterized protein LOC108150997 isoform X2
>gi|1036771177|ref|XP_017134837.1| PREDICTED: uncharacterized protein LOC108150997 isoform X1
>gi|1036771214|ref|XP_017134839.1| PREDICTED: tyramine/octopamine receptor [Drosophila miranda]
>gi|1036771125|ref|XP_017134833.1| PREDICTED: glucose-6-phosphate 1-dehydrogenase [Drosophila r
>gi|1036771144|ref|XP_017134834.1| PREDICTED: actin, larval muscle [Drosophila miranda]
>gi|1036771070|ref|XP_017134830.1| PREDICTED: delta-1-pyrroline-5-carboxylate synthase [Drosoph
>gi|1036771013|ref|XP_017134827.1| PREDICTED: uncharacterized protein LOC108150988 isoform X2
>gi|1036770993|ref|XP_017134826.1| PREDICTED: uncharacterized protein LOC108150988 isoform X1
>gi|1036771088|ref|XP_017134831.1| PREDICTED: organic cation transporter protein [Drosophila m
>gi|1036771106|ref|XP_017134832.1| PREDICTED: solute carrier family 22 member 3-like [Drosophil
>gi|1036770941|ref|XP_017134822.1| PREDICTED: uncharacterized protein LOC108150987 [Drosophila
>gi|1036770959|ref|XP_017134823.1| PREDICTED: uncharacterized protein LOC108150987 [Drosophila
>gi|1036770976|ref|XP_017134825.1| PREDICTED: uncharacterized protein LOC108150987 [Drosophila
>gi|1036761438|ref|XP_017156958.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036794440|ref|XP_017136171.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794516|ref|XP_017136175.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794535|ref|XP_017136176.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794552|ref|XP_017136177.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794569|ref|XP_017136178.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
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>gi|1036794459|ref|XP_017136172.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794478|ref|XP_017136173.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794584|ref|XP_017136179.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794600|ref|XP_017136180.1| PREDICTED: dyslexia-associated protein KIAA0319 [Drosophila
>gi|1036794675|ref|XP_017136185.1| PREDICTED: uncharacterized protein LOC108151827 [Drosophila
>gi|1036794618|ref|XP_017136181.1| PREDICTED: uncharacterized protein LOC108151826 isoform X1
>gi|1036794637|ref|XP_017136182.1| PREDICTED: uncharacterized protein LOC108151826 isoform X1
>gi|1036794656|ref|XP_017136183.1| PREDICTED: uncharacterized protein LOC108151826 isoform X2
>gi|1036765908|ref|XP_017134546.1| PREDICTED: borealin-like [Drosophila miranda]
>gi|1036848316|ref|XP_017139163.1| PREDICTED: JNK-interacting protein 1 isoform X2 [Drosophila
>gi|1036848297|ref|XP_017139162.1| PREDICTED: JNK-interacting protein 1 isoform X1 [Drosophila
>gi|1036847365|ref|XP_017139111.1| PREDICTED: tryptophan 5-hydroxylase 1 [Drosophila miranda]
>gi|1036847383|ref|XP_017139112.1| PREDICTED: uncharacterized protein LOC108153553 [Drosophila
>gi|1036826865|ref|XP_017137969.1| PREDICTED: protein bric-a-brac 2 isoform X1 [Drosophila mira
>gi|1036826884|ref|XP_017137970.1| PREDICTED: protein bric-a-brac 2 isoform X1 [Drosophila mira
>gi|1036826904|ref|XP_017137972.1| PREDICTED: protein bric-a-brac 2 isoform X2 [Drosophila mira
>gi|1036761456|ref|XP_017156959.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036847142|ref|XP_017139098.1| PREDICTED: protein bric-a-brac 1 isoform X1 [Drosophila mira
>gi|1036847161|ref|XP_017139099.1| PREDICTED: protein bric-a-brac 1 isoform X2 [Drosophila mira
>gi|1036845594|ref|XP_017139009.1| PREDICTED: oxysterol-binding protein-related protein 11 [Dro
>gi|1036788185|ref|XP_017135817.1| PREDICTED: triple functional domain protein isoform X1 [Dros
>gi|1036788204|ref|XP_017135818.1| PREDICTED: triple functional domain protein isoform X2 [Dros
>gi|1036788223|ref|XP_017135819.1| PREDICTED: triple functional domain protein isoform X3 [Dros
>gi|1036788242|ref|XP_017135821.1| PREDICTED: rho guanine nucleotide exchange factor 25 isoform
>gi|1036788300|ref|XP_017135824.1| PREDICTED: lipase member I [Drosophila miranda]
>gi|1036788280|ref|XP_017135823.1| PREDICTED: microfibril-associated glycoprotein 4 isoform X2
>gi|1036788261|ref|XP_017135822.1| PREDICTED: fibrinogen C domain-containing protein 1 isoform
>gi|1036796658|ref|XP_017136295.1| PREDICTED: cyclin-dependent kinase 14 isoform X3 [Drosophila
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>gi|1036796640|ref|XP_017136294.1| PREDICTED: cyclin-dependent kinase 14 isoform X2 [Drosophila
>gi|1036796581|ref|XP_017136291.1| PREDICTED: cyclin-dependent kinase 14 isoform X1 [Drosophila
>gi|1036796600|ref|XP_017136292.1| PREDICTED: cyclin-dependent kinase 14 isoform X1 [Drosophila
>gi|1036796619|ref|XP_017136293.1| PREDICTED: cyclin-dependent kinase 14 isoform X1 [Drosophila
>gi|1036796693|ref|XP_017136297.1| PREDICTED: cyclin-dependent kinase 14 isoform X5 [Drosophila
>gi|1036796676|ref|XP_017136296.1| PREDICTED: cyclin-dependent kinase 14 isoform X4 [Drosophila
>gi|1036843168|ref|XP_017138888.1| PREDICTED: 20-hydroxyecdysone protein [Drosophila miranda]
>gi|1036837087|ref|XP_017138547.1| PREDICTED: uncharacterized protein LOC108153227 isoform X2
>gi|1036837068|ref|XP_017138546.1| PREDICTED: uncharacterized protein LOC108153227 isoform X1
>gi|1036837104|ref|XP_017138548.1| PREDICTED: uncharacterized protein LOC108153227 isoform X3
>gi|1036837136|ref|XP_017138550.1| PREDICTED: uncharacterized protein LOC108153228 [Drosophila
>gi|1036796159|ref|XP_017136269.1| PREDICTED: solute carrier family 35 member C2 [Drosophila m
>gi|1036796177|ref|XP_017136271.1| PREDICTED: solute carrier family 35 member C2 [Drosophila m
>gi|1036796215|ref|XP_017136273.1| PREDICTED: cyclin-J-like isoform X2 [Drosophila miranda]
>gi|1036796196|ref|XP_017136272.1| PREDICTED: cyclin-J-like isoform X1 [Drosophila miranda]
>gi|1036796272|ref|XP_017136276.1| PREDICTED: uncharacterized protein LOC108151887 [Drosophila
>gi|1036796253|ref|XP_017136275.1| PREDICTED: cyclin-J-like isoform X2 [Drosophila miranda]
>gi|1036796234|ref|XP_017136274.1| PREDICTED: cyclin-J-like isoform X1 [Drosophila miranda]
>gi|1036796290|ref|XP_017136277.1| PREDICTED: probable RNA helicase armi [Drosophila miranda]
>gi|1036820796|ref|XP_017137629.1| PREDICTED: eukaryotic translation initiation factor 5B [Dros
>gi|1036827452|ref|XP_017138004.1| PREDICTED: protein sprouty [Drosophila miranda]
>gi|1036827470|ref|XP_017138005.1| PREDICTED: protein sprouty [Drosophila miranda]
>gi|1036826626|ref|XP_017137955.1| PREDICTED: iduronate 2-sulfatase [Drosophila miranda]
>gi|1036826643|ref|XP_017137956.1| PREDICTED: brain protein I3 [Drosophila miranda]
>gi|1036826607|ref|XP_017137954.1| PREDICTED: cell division cycle protein 48 homolog [Drosophil
>gi|1036824580|ref|XP_017137843.1| PREDICTED: YTH domain-containing protein 1 [Drosophila mira
>gi|1036824597|ref|XP_017137844.1| PREDICTED: osteopetrosis-associated transmembrane protein 1
>gi|1036782975|ref|XP_017135520.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036782994|ref|XP_017135521.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036783031|ref|XP_017135523.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036783050|ref|XP_017135525.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036783012|ref|XP_017135522.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036768461|ref|XP_017134683.1| PREDICTED: uncharacterized protein LOC108150885 [Drosophila
>gi|1036790307|ref|XP_017135939.1| PREDICTED: calcium uniporter protein, mitochondrial isoform
>gi|1036790326|ref|XP_017135940.1| PREDICTED: calcium uniporter protein, mitochondrial isoform
>gi|1036790345|ref|XP_017135941.1| PREDICTED: calcium uniporter protein, mitochondrial isoform
>gi|1036790288|ref|XP_017135938.1| PREDICTED: uncharacterized protein LOC108151696 [Drosophila
>gi|1036790364|ref|XP_017135942.1| PREDICTED: serine protease nudel [Drosophila miranda]
>gi|1036790382|ref|XP_017135943.1| PREDICTED: small RNA degrading nuclease 5 [Drosophila miran
>gi|1036790400|ref|XP_017135944.1| PREDICTED: innexin inx4 [Drosophila miranda]
>gi|1036845115|ref|XP_017138989.1| PREDICTED: uncharacterized protein LOC108153468 [Drosophila
>gi|1036851110|ref|XP_017139328.1| PREDICTED: glutamate receptor 1 [Drosophila miranda]
>gi|1036766014|ref|XP_017134552.1| PREDICTED: eukaryotic translation initiation factor 4E [Dros
>gi|1036761472|ref|XP_017156960.1| PREDICTED: uncharacterized protein CG32395 [Drosophila mira
>gi|1036841317|ref|XP_017138793.1| PREDICTED: uncharacterized protein LOC108153347 [Drosophila
>gi|1036841336|ref|XP_017138794.1| PREDICTED: uncharacterized protein LOC108153347 [Drosophila
>gi|1036850875|ref|XP_017139315.1| PREDICTED: ras-related and estrogen-regulated growth inhibit
>gi|1036847779|ref|XP_017139134.1| PREDICTED: tyrosine 3-monooxygenase [Drosophila miranda]
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>gi|1036788370|ref|XP_017135826.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036788318|ref|XP_017135825.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036797211|ref|XP_017136328.1| PREDICTED: uncharacterized protein LOC108151921 isoform X1
>gi|1036797226|ref|XP_017136329.1| PREDICTED: uncharacterized protein LOC108151921 isoform X2
>gi|1036797245|ref|XP_017136330.1| PREDICTED: uncharacterized protein LOC108151921 isoform X3
>gi|1036797282|ref|XP_017136332.1| PREDICTED: ribosome maturation protein SBDS [Drosophila mira
>gi|1036797263|ref|XP_017136331.1| PREDICTED: LOW QUALITY PROTEIN: IST1 homolog [Drosophila mi
>gi|1036827359|ref|XP_017137998.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036827321|ref|XP_017137996.1| PREDICTED: LOW QUALITY PROTEIN: POU domain protein CF1A [Dro
>gi|1036827340|ref|XP_017137997.1| PREDICTED: amidophosphoribosyltransferase [Drosophila miran
>gi|1036807133|ref|XP_017136884.1| PREDICTED: transcription factor SPT20 homolog isoform X3 [D
>gi|1036807152|ref|XP_017136885.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1036807099|ref|XP_017136882.1| PREDICTED: transcription factor SPT20 homolog isoform X1 [D
>gi|1036807118|ref|XP_017136883.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036807083|ref|XP_017136881.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila mira
>gi|1036850635|ref|XP_017139301.1| PREDICTED: uncharacterized protein LOC108153688 [Drosophila
>gi|1036846639|ref|XP_017139069.1| PREDICTED: guanine nucleotide-binding protein G(f) subunit a
>gi|1036842128|ref|XP_017138830.1| PREDICTED: coatomer subunit zeta-1 [Drosophila miranda]
>gi|1036850128|ref|XP_017139272.1| PREDICTED: coiled-coil domain-containing protein 13 isoform
>gi|1036850146|ref|XP_017139273.1| PREDICTED: coiled-coil domain-containing protein 13 isoform
>gi|1036814831|ref|XP_017137302.1| PREDICTED: POU domain, class 3, transcription factor 2 [Dros
>gi|1036814810|ref|XP_017137301.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X4 [Droso
>gi|1036814792|ref|XP_017137300.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X3 [Droso
>gi|1036814773|ref|XP_017137299.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X2 [Droso
>gi|1036814754|ref|XP_017137298.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X1 [Droso
>gi|1036810847|ref|XP_017137087.1| PREDICTED: protein disabled [Drosophila miranda]
>gi|1036810866|ref|XP_017137088.1| PREDICTED: uncharacterized protein LOC108152337 [Drosophila
>gi|1036768756|ref|XP_017134699.1| PREDICTED: cuticlin-2 [Drosophila miranda]
>gi|1036827652|ref|XP_017138016.1| PREDICTED: uncharacterized protein LOC108152888 [Drosophila
>gi|1036827633|ref|XP_017138015.1| PREDICTED: peptidoglycan-recognition protein SB2 [Drosophila
>gi|1036827614|ref|XP_017138014.1| PREDICTED: peptidoglycan-recognition protein SB1 [Drosophila
>gi|1036827595|ref|XP_017138013.1| PREDICTED: probable ATP-dependent RNA helicase Dbp73D [Droso
>gi|1036844534|ref|XP_017138964.1| PREDICTED: myrosinase 1 [Drosophila miranda]
>gi|1036789625|ref|XP_017135901.1| PREDICTED: neurotactin isoform X1 [Drosophila miranda]
>gi|1036789644|ref|XP_017135902.1| PREDICTED: neurotactin isoform X1 [Drosophila miranda]
>gi|1036789662|ref|XP_017135903.1| PREDICTED: neurotactin isoform X2 [Drosophila miranda]
>gi|1036789680|ref|XP_017135904.1| PREDICTED: E3 ubiquitin-protein ligase RFWD3 [Drosophila mi
>gi|1036789698|ref|XP_017135905.1| PREDICTED: cold shock domain-containing protein CG9705 [Dros
>gi|1036789716|ref|XP_017135906.1| PREDICTED: cold shock domain-containing protein CG9705 [Dros
>gi|1036789735|ref|XP_017135907.1| PREDICTED: cold shock domain-containing protein CG9705 [Dros
>gi|1036817427|ref|XP_017137448.1| PREDICTED: uncharacterized protein LOC108152547 [Drosophila
>gi|1036826463|ref|XP_017137946.1| PREDICTED: uncharacterized protein LOC108152846 isoform X1
>gi|1036826481|ref|XP_017137947.1| PREDICTED: uncharacterized protein LOC108152846 isoform X1
>gi|1036826552|ref|XP_017137951.1| PREDICTED: ephexin-1 isoform X3 [Drosophila miranda]
>gi|1036826570|ref|XP_017137952.1| PREDICTED: ephexin-1 isoform X4 [Drosophila miranda]
>gi|1036826497|ref|XP_017137948.1| PREDICTED: uncharacterized protein LOC108152846 isoform X2
>gi|1036826515|ref|XP_017137949.1| PREDICTED: uncharacterized protein LOC108152846 isoform X2
>gi|1036826534|ref|XP_017137950.1| PREDICTED: uncharacterized protein LOC108152846 isoform X2
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>gi|1036807866|ref|XP_017136924.1| PREDICTED: 2-oxoglutarate dehydrogenase, mitochondrial isof
>gi|1036807788|ref|XP_017136920.1| PREDICTED: 2-oxoglutarate dehydrogenase-like, mitochondrial
>gi|1036807847|ref|XP_017136923.1| PREDICTED: 2-oxoglutarate dehydrogenase, mitochondrial isof
>gi|1036807807|ref|XP_017136921.1| PREDICTED: 2-oxoglutarate dehydrogenase-like, mitochondrial
>gi|1036807828|ref|XP_017136922.1| PREDICTED: 2-oxoglutarate dehydrogenase-like, mitochondrial
>gi|1036761493|ref|XP_017156962.1| PREDICTED: general odorant-binding protein 70 [Drosophila mi
>gi|1036807885|ref|XP_017136925.1| PREDICTED: fibroin heavy chain isoform X1 [Drosophila mirand
>gi|1036807907|ref|XP_017136926.1| PREDICTED: keratin, type I cytoskeletal 9 isoform X2 [Drosop
>gi|1036761512|ref|XP_017156963.1| PREDICTED: uncharacterized protein LOC108165435 [Drosophila
>gi|1036848014|ref|XP_017139148.1| PREDICTED: uncharacterized protein LOC108153578 isoform X2
>gi|1036848033|ref|XP_017139149.1| PREDICTED: uncharacterized protein LOC108153578 isoform X3
>gi|1036847997|ref|XP_017139147.1| PREDICTED: uncharacterized protein LOC108153578 isoform X1
>gi|1036764820|ref|XP_017157138.1| PREDICTED: protein app1 [Drosophila miranda]
>gi|1036767699|ref|XP_017134641.1| PREDICTED: uncharacterized protein LOC108150848 [Drosophila
>gi|1036804065|ref|XP_017136714.1| PREDICTED: prickle-like protein 3 isoform X5 [Drosophila mi
>gi|1036804046|ref|XP_017136713.1| PREDICTED: four and a half LIM domains protein 2 isoform X4
>gi|1036804083|ref|XP_017136715.1| PREDICTED: prickle-like protein 3 isoform X6 [Drosophila mi
>gi|1036804011|ref|XP_017136710.1| PREDICTED: trichohyalin isoform X2 [Drosophila miranda]
>gi|1036804105|ref|XP_017136716.1| PREDICTED: four and a half LIM domains protein 2 isoform X7
>gi|1036804122|ref|XP_017136717.1| PREDICTED: four and a half LIM domains protein 2 isoform X8
>gi|1036804029|ref|XP_017136711.1| PREDICTED: trichohyalin isoform X3 [Drosophila miranda]
>gi|1036803992|ref|XP_017136709.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1036804141|ref|XP_017136718.1| PREDICTED: peptidyl-prolyl cis-trans isomerase CYP95 [Droso
>gi|1036840242|ref|XP_017138731.1| PREDICTED: coiled-coil domain-containing protein 22 homolog
>gi|1036787262|ref|XP_017135764.1| PREDICTED: opsin Rh4 [Drosophila miranda]
>gi|1036787297|ref|XP_017135765.1| PREDICTED: probable E3 ubiquitin-protein ligase sinah [Dros
>gi|1036787318|ref|XP_017135767.1| PREDICTED: E3 ubiquitin-protein ligase sina [Drosophila mira
>gi|1036787337|ref|XP_017135768.1| PREDICTED: E3 ubiquitin-protein ligase sina [Drosophila mira
>gi|1036761531|ref|XP_017156964.1| PREDICTED: uncharacterized protein LOC108165436 [Drosophila
>gi|1036787353|ref|XP_017135769.1| PREDICTED: uncharacterized protein LOC108151600 [Drosophila
>gi|1036802830|ref|XP_017136643.1| PREDICTED: transmembrane protein 258 homolog [Drosophila mi
>gi|1036802792|ref|XP_017136641.1| PREDICTED: tumor susceptibility gene 101 protein isoform X1
>gi|1036802811|ref|XP_017136642.1| PREDICTED: tumor susceptibility gene 101 protein isoform X2
>gi|1036802866|ref|XP_017136646.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 6
>gi|1036802848|ref|XP_017136645.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 6
>gi|1036838660|ref|XP_017138646.1| PREDICTED: uncharacterized protein LOC108153285 [Drosophila
>gi|1036838641|ref|XP_017138645.1| PREDICTED: rhythmically expressed gene 2 protein [Drosophila
>gi|1036800197|ref|XP_017136496.1| PREDICTED: UDP-glucose 4-epimerase [Drosophila miranda]
>gi|1036800216|ref|XP_017136497.1| PREDICTED: UDP-glucose 4-epimerase [Drosophila miranda]
>gi|1036800253|ref|XP_017136499.1| PREDICTED: uncharacterized protein C45G9.7 [Drosophila mira
>gi|1036800235|ref|XP_017136498.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036800178|ref|XP_017136495.1| PREDICTED: DNA repair protein REV1 [Drosophila miranda]
>gi|1036834067|ref|XP_017138372.1| PREDICTED: uncharacterized protein LOC108153099 [Drosophila
>gi|1036842235|ref|XP_017138837.1| PREDICTED: uncharacterized protein LOC108153377 [Drosophila
>gi|1036842254|ref|XP_017138838.1| PREDICTED: uncharacterized protein LOC108153377 [Drosophila
>gi|1036836483|ref|XP_017138512.1| PREDICTED: uncharacterized protein LOC108153202 [Drosophila
>gi|1036809010|ref|XP_017136986.1| PREDICTED: retinoid-inducible serine carboxypeptidase [Dros
>gi|1036809028|ref|XP_017136987.1| PREDICTED: retinoid-inducible serine carboxypeptidase [Dros
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>gi|1036809064|ref|XP_017136989.1| PREDICTED: ras-related protein Rab-14 [Drosophila miranda]
>gi|1036809046|ref|XP_017136988.1| PREDICTED: vesicle transport through interaction with t-SNAR
>gi|1036808993|ref|XP_017136985.1| PREDICTED: LOW QUALITY PROTEIN: mediator of RNA polymerase
>gi|1036832798|ref|XP_017138308.1| PREDICTED: uncharacterized protein LOC108153047 isoform X2
>gi|1036832720|ref|XP_017138304.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036832741|ref|XP_017138305.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036832760|ref|XP_017138306.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036832779|ref|XP_017138307.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036851636|ref|XP_017139360.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036851656|ref|XP_017139361.1| PREDICTED: cuticle protein 38-like [Drosophila miranda]
>gi|1036851420|ref|XP_017139347.1| PREDICTED: cuticle protein 38-like [Drosophila miranda]
>gi|1036851402|ref|XP_017139345.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036816806|ref|XP_017137415.1| PREDICTED: histone-lysine N-methyltransferase 2D [Drosophila
>gi|1036816825|ref|XP_017137416.1| PREDICTED: protein extra-macrochaetae [Drosophila miranda]
>gi|1036798180|ref|XP_017136382.1| PREDICTED: splicing factor 3B subunit 3 isoform X1 [Drosoph
>gi|1036798199|ref|XP_017136383.1| PREDICTED: splicing factor 3B subunit 3 isoform X2 [Drosoph
>gi|1036798218|ref|XP_017136384.1| PREDICTED: splicing factor 3B subunit 3 isoform X3 [Drosoph
>gi|1036798253|ref|XP_017136386.1| PREDICTED: uncharacterized protein LOC108151958 [Drosophila
>gi|1036798235|ref|XP_017136385.1| PREDICTED: kidney mitochondrial carrier protein 1 [Drosophil
>gi|1036765762|ref|XP_017134538.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036826695|ref|XP_017137960.1| PREDICTED: uncharacterized protein LOC108152853 [Drosophila
>gi|1036826711|ref|XP_017137961.1| PREDICTED: uncharacterized protein LOC108152853 [Drosophila
>gi|1036826729|ref|XP_017137962.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036826660|ref|XP_017137957.1| PREDICTED: gustatory receptor 68a [Drosophila miranda]
>gi|1036826677|ref|XP_017137959.1| PREDICTED: gustatory receptor 68a [Drosophila miranda]
>gi|1036761550|ref|XP_017156965.1| PREDICTED: LOW QUALITY PROTEIN: 2-oxoglutarate dehydrogenase
>gi|1036767830|ref|XP_017134648.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036807170|ref|XP_017136886.1| PREDICTED: phosphatidylinositol 4-phosphate 3-kinase C2 doma
>gi|1036807187|ref|XP_017136887.1| PREDICTED: phosphatidylinositol 4-phosphate 3-kinase C2 doma
>gi|1036807206|ref|XP_017136888.1| PREDICTED: phosphatidylinositol 4-phosphate 3-kinase C2 doma
>gi|1036807244|ref|XP_017136891.1| PREDICTED: uncharacterized protein LOC108152221 [Drosophila
>gi|1036807225|ref|XP_017136889.1| PREDICTED: uncharacterized protein LOC108152220 isoform X2
>gi|1036782214|ref|XP_017135475.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036782252|ref|XP_017135478.1| PREDICTED: peptidyl-prolyl cis-trans isomerase CWC27 homolog
>gi|1036782232|ref|XP_017135476.1| PREDICTED: kinesin-like protein Klp68D [Drosophila miranda]
>gi|1036782288|ref|XP_017135480.1| PREDICTED: 2-amino-3-ketobutyrate coenzyme A ligase, mitocho
>gi|1036782175|ref|XP_017135473.1| PREDICTED: cubilin homolog isoform X1 [Drosophila miranda]
>gi|1036782196|ref|XP_017135474.1| PREDICTED: cubilin homolog isoform X2 [Drosophila miranda]
>gi|1036782305|ref|XP_017135481.1| PREDICTED: uncharacterized protein LOC108151404 [Drosophila
>gi|1036782270|ref|XP_017135479.1| PREDICTED: uncharacterized protein LOC108151402 [Drosophila
>gi|1036782324|ref|XP_017135482.1| PREDICTED: 60S ribosomal protein L10a-2 [Drosophila miranda]
>gi|1036793404|ref|XP_017136111.1| PREDICTED: uncharacterized protein LOC108151779 [Drosophila
>gi|1036793465|ref|XP_017136115.1| PREDICTED: NADH-cytochrome b5 reductase 3 isoform X1 [Drosop
>gi|1036793484|ref|XP_017136116.1| PREDICTED: NADH-cytochrome b5 reductase 3 isoform X2 [Drosop
>gi|1036793423|ref|XP_017136112.1| PREDICTED: G2/mitotic-specific cyclin-A [Drosophila miranda]
>gi|1036793446|ref|XP_017136114.1| PREDICTED: RIB43A-like with coiled-coils protein 2 [Drosophi
>gi|1036843187|ref|XP_017138889.1| PREDICTED: alpha-ketoglutarate-dependent dioxygenase alkB ho
>gi|1036843262|ref|XP_017138893.1| PREDICTED: BPTI/Kunitz domain-containing protein-like [Droso
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>gi|1036843223|ref|XP_017138891.1| PREDICTED: uncharacterized protein LOC108153408 isoform X1
>gi|1036843241|ref|XP_017138892.1| PREDICTED: BPTI/Kunitz domain-containing protein-like isofo
>gi|1036843206|ref|XP_017138890.1| PREDICTED: kunitz-type serine protease inhibitor NACI [Dros
>gi|1036761570|ref|XP_017156966.1| PREDICTED: palmitoleoyl-protein carboxylesterase NOTUM [Dros
>gi|1036808506|ref|XP_017136961.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808524|ref|XP_017136962.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808542|ref|XP_017136963.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808560|ref|XP_017136964.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808579|ref|XP_017136965.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808618|ref|XP_017136967.1| PREDICTED: uncharacterized protein LOC108152263 isoform X3
>gi|1036808599|ref|XP_017136966.1| PREDICTED: probable RNA-binding protein orb2 isoform X2 [Dr
>gi|1036808636|ref|XP_017136968.1| PREDICTED: gram-negative bacteria-binding protein 3 [Drosopl
>gi|1036793983|ref|XP_017136142.1| PREDICTED: 50S ribosomal protein L7/L12 [Drosophila miranda]
>gi|1036793945|ref|XP_017136140.1| PREDICTED: uncharacterized protein LOC108151800 [Drosophila
>gi|1036793925|ref|XP_017136139.1| PREDICTED: valine--tRNA ligase, mitochondrial [Drosophila m:
>gi|1036793887|ref|XP_017136137.1| PREDICTED: coatomer subunit alpha [Drosophila miranda]
>gi|1036793906|ref|XP_017136138.1| PREDICTED: coatomer subunit alpha [Drosophila miranda]
>gi|1036793964|ref|XP_017136141.1| PREDICTED: protein big brother [Drosophila miranda]
>gi|1036831812|ref|XP_017138253.1| PREDICTED: methyltransferase-like protein isoform X2 [Droso]
>gi|1036831793|ref|XP_017138252.1| PREDICTED: methyltransferase-like protein isoform X1 [Droso]
>gi|1036831846|ref|XP_017138256.1| PREDICTED: vesicle-associated membrane protein 2 isoform X2
>gi|1036831865|ref|XP_017138257.1| PREDICTED: vesicle-associated membrane protein 2 isoform X3
>gi|1036831829|ref|XP_017138255.1| PREDICTED: vesicle-associated membrane protein 2 isoform X1
>gi|1036831901|ref|XP_017138258.1| PREDICTED: vesicle-associated membrane protein 2 isoform X4
>gi|1036793173|ref|XP_017136101.1| PREDICTED: uncharacterized protein LOC108151768 [Drosophila
>gi|1036793062|ref|XP_017136095.1| PREDICTED: vitamin K-dependent gamma-carboxylase isoform X1
>gi|1036793079|ref|XP_017136096.1| PREDICTED: vitamin K-dependent gamma-carboxylase isoform X2
>gi|1036793043|ref|XP_017136094.1| PREDICTED: ATP-binding cassette sub-family B member 7, mitoc
>gi|1036793025|ref|XP_017136093.1| PREDICTED: ATP-binding cassette sub-family B member 7, mitoc
>gi|1036793154|ref|XP_017136100.1| PREDICTED: SET domain-containing protein 4 [Drosophila mirar
>gi|1036793191|ref|XP_017136102.1| PREDICTED: protein OPI10 homolog [Drosophila miranda]
>gi|1036793006|ref|XP_017136092.1| PREDICTED: 2-oxoglutarate dehydrogenase, mitochondrial [Dros
>gi|1036792988|ref|XP_017136091.1| PREDICTED: uncharacterized protein LOC108151760 [Drosophila
>gi|1036793118|ref|XP_017136098.1| PREDICTED: uncharacterized protein LOC108151765 isoform X2
>gi|1036793099|ref|XP_017136097.1| PREDICTED: uncharacterized protein LOC108151765 isoform X1
>gi|1036793135|ref|XP_017136099.1| PREDICTED: uncharacterized protein LOC108151765 isoform X3
>gi|1036795601|ref|XP_017136237.1| PREDICTED: TOX high mobility group box family member 4-like
>gi|1036761589|ref|XP_017156967.1| PREDICTED: TOX high mobility group box family member 4-like
>gi|1036795513|ref|XP_017136232.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795531|ref|XP_017136233.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795550|ref|XP_017136234.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795566|ref|XP_017136235.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795584|ref|XP_017136236.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036851274|ref|XP_017139338.1| PREDICTED: uncharacterized protein LOC108153721 isoform X1
>gi|1036851292|ref|XP_017139339.1| PREDICTED: ninjurin-2 isoform X2 [Drosophila miranda]
>gi|1036837030|ref|XP_017138544.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036837049|ref|XP_017138545.1| PREDICTED: uncharacterized protein LOC108153226 [Drosophila
>gi|1036837011|ref|XP_017138543.1| PREDICTED: guanylate kinase [Drosophila miranda]
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>gi|1036761608|ref|XP_017156968.1| PREDICTED: LOW QUALITY PROTEIN: DNA polymerase delta small s
>gi|1036842162|ref|XP_017138832.1| PREDICTED: WD repeat-containing protein 35 isoform X2 [Dros
>gi|1036842144|ref|XP_017138831.1| PREDICTED: WD repeat-containing protein 35 isoform X1 [Dros
>gi|1036851364|ref|XP_017139343.1| PREDICTED: uncharacterized protein At4g17910 isoform X1 [Dr
>gi|1036851383|ref|XP_017139344.1| PREDICTED: uncharacterized protein At4g17910 isoform X2 [Dr
>gi|1036835025|ref|XP_017138425.1| PREDICTED: glycerol kinase [Drosophila miranda]
>gi|1036835043|ref|XP_017138427.1| PREDICTED: glycerol kinase [Drosophila miranda]
>gi|1036817538|ref|XP_017137455.1| PREDICTED: uncharacterized protein LOC108152552 isoform X1
>gi|1036817559|ref|XP_017137456.1| PREDICTED: uncharacterized protein LOC108152552 isoform X2
>gi|1036817578|ref|XP_017137457.1| PREDICTED: uncharacterized protein LOC108152552 isoform X3
>gi|1036817597|ref|XP_017137458.1| PREDICTED: uncharacterized protein LOC108152552 isoform X4
>gi|1036817615|ref|XP_017137459.1| PREDICTED: probable chitinase 2 isoform X1 [Drosophila mira
>gi|1036817635|ref|XP_017137460.1| PREDICTED: probable chitinase 2 isoform X2 [Drosophila mira
>gi|1036817655|ref|XP_017137462.1| PREDICTED: probable chitinase 2 isoform X2 [Drosophila mira
>gi|1036813090|ref|XP_017137202.1| PREDICTED: membrane-associated protein Hem [Drosophila mira
>gi|1036813184|ref|XP_017137207.1| PREDICTED: SRSF protein kinase 3 isoform X5 [Drosophila mira
>gi|1036813145|ref|XP_017137205.1| PREDICTED: SRSF protein kinase 1 isoform X3 [Drosophila mira
>gi|1036813166|ref|XP_017137206.1| PREDICTED: SRSF protein kinase 3 isoform X4 [Drosophila mira
>gi|1036813126|ref|XP_017137204.1| PREDICTED: SRSF protein kinase 1 isoform X2 [Drosophila mira
>gi|1036813108|ref|XP_017137203.1| PREDICTED: SRSF protein kinase 1 isoform X1 [Drosophila mira
>gi|1036823186|ref|XP_017137764.1| PREDICTED: dnaJ homolog subfamily C member 5 homolog isoform
>gi|1036823168|ref|XP_017137763.1| PREDICTED: dnaJ homolog subfamily C member 5 homolog isoform
>gi|1036837740|ref|XP_017138588.1| PREDICTED: pancreatic lipase-related protein 3 [Drosophila m
>gi|1036837646|ref|XP_017138582.1| PREDICTED: speract receptor isoform X1 [Drosophila miranda]
>gi|1036837683|ref|XP_017138584.1| PREDICTED: speract receptor isoform X3 [Drosophila miranda]
>gi|1036837664|ref|XP_017138583.1| PREDICTED: speract receptor isoform X2 [Drosophila miranda]
>gi|1036837702|ref|XP_017138585.1| PREDICTED: speract receptor isoform X4 [Drosophila miranda]
>gi|1036837721|ref|XP_017138587.1| PREDICTED: dual specificity protein phosphatase 22-like [Dro
>gi|1036845743|ref|XP_017139018.1| PREDICTED: uncharacterized protein LOC108153488 isoform X2
>gi|1036845724|ref|XP_017139017.1| PREDICTED: uncharacterized protein LOC108153488 isoform X1
>gi|1036818834|ref|XP_017137530.1| PREDICTED: BAG domain-containing protein Samui isoform X1 [
>gi|1036818873|ref|XP_017137532.1| PREDICTED: BAG domain-containing protein Samui isoform X3 [
>gi|1036818854|ref|XP_017137531.1| PREDICTED: BAG domain-containing protein Samui isoform X2 [
>gi|1036818892|ref|XP_017137533.1| PREDICTED: BAG domain-containing protein Samui isoform X4 [
>gi|1036818913|ref|XP_017137534.1| PREDICTED: BAG domain-containing protein Samui isoform X5 [
>gi|1036818931|ref|XP_017137535.1| PREDICTED: BAG domain-containing protein Samui isoform X5 [
>gi|1036818951|ref|XP_017137536.1| PREDICTED: drebrin-like protein [Drosophila miranda]
>gi|1036785030|ref|XP_017135640.1| PREDICTED: protein suppressor of hairy wing [Drosophila mira
>gi|1036785012|ref|XP_017135639.1| PREDICTED: protein-cysteine N-palmitoyltransferase Rasp [Dro
>gi|1036784993|ref|XP_017135638.1| PREDICTED: uncharacterized protein LOC108151504 [Drosophila
>gi|1036785049|ref|XP_017135641.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prote
>gi|1036784938|ref|XP_017135634.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X1
>gi|1036784974|ref|XP_017135636.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036784956|ref|XP_017135635.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036792881|ref|XP_017136085.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 5 [Drosophi
>gi|1036792899|ref|XP_017136086.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 5 [Drosophi
>gi|1036792917|ref|XP_017136087.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 5 [Drosophi
>gi|1036792971|ref|XP_017136090.1| PREDICTED: alpha-1,3/1,6-mannosyltransferase ALG2 [Drosophil
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>gi|1036792953|ref|XP_017136089.1| PREDICTED: hippocampus abundant transcript 1 protein isofor
>gi|1036792934|ref|XP_017136088.1| PREDICTED: hippocampus abundant transcript 1 protein isofor
>gi|1036847642|ref|XP_017139126.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1036805614|ref|XP_017136802.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036805633|ref|XP_017136803.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036805652|ref|XP_017136804.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036805689|ref|XP_017136806.1| PREDICTED: prominin-like protein isoform X3 [Drosophila mira
>gi|1036805670|ref|XP_017136805.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1036805708|ref|XP_017136807.1| PREDICTED: prominin-like protein isoform X4 [Drosophila mira
>gi|1036805727|ref|XP_017136809.1| PREDICTED: prominin-like protein isoform X5 [Drosophila mira
>gi|1036845685|ref|XP_017139015.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036840034|ref|XP_017138719.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036840051|ref|XP_017138720.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1036811818|ref|XP_017137139.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036811799|ref|XP_017137138.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036811780|ref|XP_017137137.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036811837|ref|XP_017137140.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036780403|ref|XP_017135371.1| PREDICTED: cysteine and histidine-rich protein 1 homolog [D
>gi|1036780442|ref|XP_017135373.1| PREDICTED: protein FAM207A [Drosophila miranda]
>gi|1036780365|ref|XP_017135368.1| PREDICTED: cyclin-dependent kinase 8 [Drosophila miranda]
>gi|1036780330|ref|XP_017135367.1| PREDICTED: putative glycerol kinase 5 [Drosophila miranda]
>gi|1036780422|ref|XP_017135372.1| PREDICTED: CRAL-TRIO domain-containing protein C3H8.02 [Dros
>gi|1036780296|ref|XP_017135365.1| PREDICTED: sphingosine kinase 2 [Drosophila miranda]
>gi|1036780311|ref|XP_017135366.1| PREDICTED: dnaJ homolog subfamily C member 2 [Drosophila mi
>gi|1036780384|ref|XP_017135369.1| PREDICTED: islet cell autoantigen 1 [Drosophila miranda]
>gi|1036823112|ref|XP_017137759.1| PREDICTED: uncharacterized protein LOC108152730 [Drosophila
>gi|1036823129|ref|XP_017137760.1| PREDICTED: uncharacterized protein LOC108152730 [Drosophila
>gi|1036823147|ref|XP_017137761.1| PREDICTED: uncharacterized protein LOC108152731 [Drosophila
>gi|1036822457|ref|XP_017137721.1| PREDICTED: dnaJ protein homolog 1-like isoform X1 [Drosophil
>gi|1036822476|ref|XP_017137722.1| PREDICTED: dnaJ protein homolog 1-like isoform X2 [Drosophil
>gi|1036822497|ref|XP_017137724.1| PREDICTED: LOW QUALITY PROTEIN: dnaJ protein homolog 1-like
>gi|1036761627|ref|XP_017156969.1| PREDICTED: enhancer of rudimentary homolog [Drosophila mira
>gi|1036845556|ref|XP_017139007.1| PREDICTED: uncharacterized protein LOC108153479 [Drosophila
>gi|1036845575|ref|XP_017139008.1| PREDICTED: uncharacterized protein LOC108153479 [Drosophila
>gi|1036764858|ref|XP_017157140.1| PREDICTED: phosphoglycolate phosphatase 2 [Drosophila miran
>gi|1036848219|ref|XP_017139158.1| PREDICTED: glycerol-3-phosphate phosphatase [Drosophila mira
>gi|1036848200|ref|XP_017139157.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1036850267|ref|XP_017139279.1| PREDICTED: FMRFamide receptor [Drosophila miranda]
>gi|1036822696|ref|XP_017137735.1| PREDICTED: carnitine O-palmitoyltransferase 2, mitochondrial
>gi|1036822773|ref|XP_017137739.1| PREDICTED: odorant receptor 63a [Drosophila miranda]
>gi|1036822538|ref|XP_017137726.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822519|ref|XP_017137725.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822673|ref|XP_017137734.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822654|ref|XP_017137732.1| PREDICTED: uncharacterized protein LOC108152711 isoform X8
>gi|1036822635|ref|XP_017137731.1| PREDICTED: uncharacterized protein LOC108152711 isoform X7
>gi|1036822595|ref|XP_017137729.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822614|ref|XP_017137730.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822576|ref|XP_017137728.1| PREDICTED: potassium voltage-gated channel protein Shab iso
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>gi|1036822557|ref|XP_017137727.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822715|ref|XP_017137736.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036761646|ref|XP_017156970.1| PREDICTED: uncharacterized protein LOC108165443 [Drosophila
>gi|1036822734|ref|XP_017137737.1| PREDICTED: uncharacterized protein LOC108152714 [Drosophila
>gi|1036822754|ref|XP_017137738.1| PREDICTED: uncharacterized protein LOC108152714 [Drosophila
>gi|1036786583|ref|XP_017135729.1| PREDICTED: LOW QUALITY PROTEIN: DNA N6-methyl adenine demetl
>gi|1036818382|ref|XP_017137503.1| PREDICTED: N-acetylgalactosaminyltransferase 6 [Drosophila r
>gi|1036818400|ref|XP_017137504.1| PREDICTED: 28S ribosomal protein S35, mitochondrial [Drosopl
>gi|1036809104|ref|XP_017136992.1| PREDICTED: pyridoxal kinase [Drosophila miranda]
>gi|1036809143|ref|XP_017136994.1| PREDICTED: uncharacterized protein LOC108152287 [Drosophila
>gi|1036809083|ref|XP_017136990.1| PREDICTED: kinesin-like protein KIF18A [Drosophila miranda]
>gi|1036809123|ref|XP_017136993.1| PREDICTED: ribosome-recycling factor, mitochondrial [Drosopl
>gi|1036831095|ref|XP_017138214.1| PREDICTED: protein FAM76A isoform X1 [Drosophila miranda]
>gi|1036831115|ref|XP_017138215.1| PREDICTED: uncharacterized protein LOC108152991 isoform X2
>gi|1036831151|ref|XP_017138217.1| PREDICTED: adrenodoxin-like protein, mitochondrial [Drosophi
>gi|1036831133|ref|XP_017138216.1| PREDICTED: heat shock protein 67B3 [Drosophila miranda]
>gi|1036845134|ref|XP_017138990.1| PREDICTED: heat shock protein 67B2 [Drosophila miranda]
>gi|1036845153|ref|XP_017138991.1| PREDICTED: heat shock protein 22 [Drosophila miranda]
>gi|1036832955|ref|XP_017138317.1| PREDICTED: heat shock protein 27 [Drosophila miranda]
>gi|1036832938|ref|XP_017138316.1| PREDICTED: heat shock protein 26 [Drosophila miranda]
>gi|1036832919|ref|XP_017138314.1| PREDICTED: uncharacterized protein LOC108153054 [Drosophila
>gi|1036845613|ref|XP_017139010.1| PREDICTED: heat shock protein 23 [Drosophila miranda]
>gi|1036816787|ref|XP_017137414.1| PREDICTED: LOW QUALITY PROTEIN: heat shock protein 27 [Dros
>gi|1036816764|ref|XP_017137413.1| PREDICTED: uncharacterized protein LOC108152523 [Drosophila
>gi|1036831436|ref|XP_017138233.1| PREDICTED: 40S ribosomal protein S17 [Drosophila miranda]
>gi|1036831417|ref|XP_017138232.1| PREDICTED: SHC-transforming protein 2 [Drosophila miranda]
>gi|1036798693|ref|XP_017136412.1| PREDICTED: phosphoserine phosphatase [Drosophila miranda]
>gi|1036798565|ref|XP_017136405.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [I
>gi|1036798600|ref|XP_017136406.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [I
>gi|1036798619|ref|XP_017136407.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [I
>gi|1036798637|ref|XP_017136408.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [I
>gi|1036798655|ref|XP_017136409.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X2 [I
>gi|1036798674|ref|XP_017136410.1| PREDICTED: ataxin-2 homolog [Drosophila miranda]
>gi|1036798711|ref|XP_017136413.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036798730|ref|XP_017136414.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036818252|ref|XP_017137495.1| PREDICTED: uncharacterized protein LOC108152569 [Drosophila
>gi|1036818271|ref|XP_017137497.1| PREDICTED: uncharacterized protein LOC108152570 [Drosophila
>gi|1036764038|ref|XP_017157101.1| PREDICTED: sentrin-specific protease 1-like [Drosophila mir
>gi|1036761663|ref|XP_017156971.1| PREDICTED: NADH-quinone oxidoreductase subunit C-like, part
>gi|1036790531|ref|XP_017135951.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X1
>gi|1036790550|ref|XP_017135952.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X1
>gi|1036790569|ref|XP_017135953.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X2
>gi|1036790640|ref|XP_017135957.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X6
>gi|1036790657|ref|XP_017135958.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X7
>gi|1036790676|ref|XP_017135959.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036790588|ref|XP_017135954.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X3
>gi|1036790769|ref|XP_017135964.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X1
>gi|1036790695|ref|XP_017135960.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
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>gi|1036790751|ref|XP_017135963.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X
>gi|1036790604|ref|XP_017135955.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X4
>gi|1036790622|ref|XP_017135956.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X5
>gi|1036790714|ref|XP_017135961.1| PREDICTED: mitogen-activated protein kinase kinase kinase k
>gi|1036790733|ref|XP_017135962.1| PREDICTED: misshapen-like kinase 1 isoform X11 [Drosophila m
>gi|1036841577|ref|XP_017138808.1| PREDICTED: 60S ribosomal protein L8 [Drosophila miranda]
>gi|1036794021|ref|XP_017136147.1| PREDICTED: protein daughter of sevenless [Drosophila miranda
>gi|1036794054|ref|XP_017136149.1| PREDICTED: enkurin [Drosophila miranda]
>gi|1036794038|ref|XP_017136148.1| PREDICTED: uncharacterized protein LOC108151806 [Drosophila
>gi|1036794088|ref|XP_017136151.1| PREDICTED: acyl-coenzyme A thioesterase 13-like [Drosophila
>gi|1036794071|ref|XP_017136150.1| PREDICTED: acyl-coenzyme A thioesterase 13-like [Drosophila
>gi|1036794105|ref|XP_017136152.1| PREDICTED: acyl-coenzyme A thioesterase 13 [Drosophila miran
>gi|1036794123|ref|XP_017136153.1| PREDICTED: acyl-coenzyme A thioesterase 13 [Drosophila miran
>gi|1036794002|ref|XP_017136143.1| PREDICTED: peroxidasin [Drosophila miranda]
>gi|1036804921|ref|XP_017136763.1| PREDICTED: uncharacterized protein KIAA1143 homolog [Drosopl
>gi|1036804845|ref|XP_017136758.1| PREDICTED: uncharacterized protein LOC108152153 isoform X1
>gi|1036804864|ref|XP_017136759.1| PREDICTED: uncharacterized protein LOC108152153 isoform X1
>gi|1036804883|ref|XP_017136760.1| PREDICTED: uncharacterized protein LOC108152153 isoform X2
>gi|1036804902|ref|XP_017136762.1| PREDICTED: uncharacterized protein LOC108152154 [Drosophila
>gi|1036851963|ref|XP_017139382.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036840015|ref|XP_017138718.1| PREDICTED: uncharacterized protein LOC108153317 [Drosophila
>gi|1036839981|ref|XP_017138715.1| PREDICTED: probable proline--tRNA ligase, mitochondrial isoi
>gi|1036840000|ref|XP_017138716.1| PREDICTED: probable proline--tRNA ligase, mitochondrial isoi
>gi|1036831265|ref|XP_017138224.1| PREDICTED: snurportin-1 [Drosophila miranda]
>gi|1036831244|ref|XP_017138223.1| PREDICTED: uncharacterized protein LOC108152996 [Drosophila
>gi|1036831284|ref|XP_017138225.1| PREDICTED: uncharacterized protein LOC108152998 [Drosophila
>gi|1036833786|ref|XP_017138354.1| PREDICTED: signal recognition particle 54 kDa protein [Droso
>gi|1036772751|ref|XP_017134926.1| PREDICTED: flap endonuclease GEN [Drosophila miranda]
>gi|1036772677|ref|XP_017134922.1| PREDICTED: alanine--tRNA ligase, mitochondrial [Drosophila m
>gi|1036772789|ref|XP_017134928.1| PREDICTED: FIT family protein CG10671 [Drosophila miranda]
>gi|1036772849|ref|XP_017134930.1| PREDICTED: ganglioside-induced differentiation-associated pi
>gi|1036772868|ref|XP_017134932.1| PREDICTED: ganglioside-induced differentiation-associated pi
>gi|1036772828|ref|XP_017134929.1| PREDICTED: ganglioside-induced differentiation-associated pi
>gi|1036772887|ref|XP_017134933.1| PREDICTED: dehydrogenase/reductase SDR family member 4 [Dros
>gi|1036772937|ref|XP_017134936.1| PREDICTED: charged multivesicular body protein 2b-B [Drosopl
>gi|1036772920|ref|XP_017134935.1| PREDICTED: TP53-regulating kinase [Drosophila miranda]
>gi|1036772770|ref|XP_017134927.1| PREDICTED: pentatricopeptide repeat-containing protein 1, m
>gi|1036772954|ref|XP_017134937.1| PREDICTED: protein Asterix [Drosophila miranda]
>gi|1036772903|ref|XP_017134934.1| PREDICTED: ubiquitin thioesterase OTU1 [Drosophila miranda]
>gi|1036773045|ref|XP_017134941.1| PREDICTED: uncharacterized protein LOC108151073 [Drosophila
>gi|1036773020|ref|XP_017134940.1| PREDICTED: uncharacterized protein LOC108151072 [Drosophila
>gi|1036761681|ref|XP_017156972.1| PREDICTED: uncharacterized protein LOC108165445 [Drosophila
>gi|1036761700|ref|XP_017156973.1| PREDICTED: LOW QUALITY PROTEIN: myb-like protein AA [Drosopl
>gi|1036772969|ref|XP_017134938.1| PREDICTED: uncharacterized protein LOC108151070 [Drosophila
>gi|1036772986|ref|XP_017134939.1| PREDICTED: uncharacterized protein LOC108151070 [Drosophila
>gi|1036772696|ref|XP_017134923.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036772715|ref|XP_017134924.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036772733|ref|XP_017134925.1| PREDICTED: proteoglycan 4 isoform X2 [Drosophila miranda]
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>gi|1036772658|ref|XP_017134921.1| PREDICTED: helicase POLQ-like [Drosophila miranda]
>gi|1036792714|ref|XP_017136077.1| PREDICTED: protein IWS1 homolog A-like [Drosophila miranda]
>gi|1036792700|ref|XP_017136075.1| PREDICTED: uncharacterized ATP-dependent helicase C29A10.10
>gi|1036792733|ref|XP_017136078.1| PREDICTED: phosphatidate cytidylyltransferase, photorecepto
>gi|1036827995|ref|XP_017138035.1| PREDICTED: SUMO-activating enzyme subunit 2 [Drosophila mir
>gi|1036799198|ref|XP_017136441.1| PREDICTED: uncharacterized protein LOC108151987 [Drosophila
>gi|1036799179|ref|XP_017136440.1| PREDICTED: 2-aminoethanethiol dioxygenase [Drosophila miran
>gi|1036799161|ref|XP_017136439.1| PREDICTED: lariat debranching enzyme [Drosophila miranda]
>gi|1036799143|ref|XP_017136438.1| PREDICTED: importin-7 [Drosophila miranda]
>gi|1036812528|ref|XP_017137177.1| PREDICTED: actin-related protein 3 [Drosophila miranda]
>gi|1036812510|ref|XP_017137176.1| PREDICTED: transcription factor 25 [Drosophila miranda]
>gi|1036818098|ref|XP_017137487.1| PREDICTED: DENN domain-containing protein 5B isoform X1 [Dro
>gi|1036818115|ref|XP_017137488.1| PREDICTED: DENN domain-containing protein 5B isoform X1 [Dro
>gi|1036818134|ref|XP_017137489.1| PREDICTED: DENN domain-containing protein 5B isoform X1 [Dro
>gi|1036818155|ref|XP_017137490.1| PREDICTED: DENN domain-containing protein 5B isoform X2 [Dro
>gi|1036837150|ref|XP_017138551.1| PREDICTED: protein phosphatase PTC7 homolog [Drosophila mir
>gi|1036779983|ref|XP_017135347.1| PREDICTED: DEP domain-containing protein 5 isoform X1 [Drosc
>gi|1036780004|ref|XP_017135348.1| PREDICTED: DEP domain-containing protein 5 isoform X2 [Drosc
>gi|1036780023|ref|XP_017135349.1| PREDICTED: DEP domain-containing protein 5 isoform X3 [Drosc
>gi|1036780042|ref|XP_017135350.1| PREDICTED: DEP domain-containing protein 5 isoform X4 [Drosc
>gi|1036780061|ref|XP_017135351.1| PREDICTED: DEP domain-containing protein 5 isoform X5 [Drosc
>gi|1036780079|ref|XP_017135352.1| PREDICTED: DEP domain-containing protein 5 isoform X6 [Drosc
>gi|1036780096|ref|XP_017135353.1| PREDICTED: DEP domain-containing protein 5 isoform X7 [Drosc
>gi|1036780114|ref|XP_017135354.1| PREDICTED: DEP domain-containing protein 5 isoform X8 [Drosc
>gi|1036780129|ref|XP_017135355.1| PREDICTED: DEP domain-containing protein 5 isoform X9 [Drosc
>gi|1036780148|ref|XP_017135356.1| PREDICTED: DEP domain-containing protein 5 isoform X10 [Dros
>gi|1036780277|ref|XP_017135364.1| PREDICTED: uncharacterized protein LOC108151323 [Drosophila
>gi|1036780167|ref|XP_017135357.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X1 [Di
>gi|1036780239|ref|XP_017135362.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X3 [Di
>gi|1036780185|ref|XP_017135359.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X2 [Di
>gi|1036780201|ref|XP_017135360.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X2 [Di
>gi|1036780220|ref|XP_017135361.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X2 [Di
>gi|1036780258|ref|XP_017135363.1| PREDICTED: protein cueball [Drosophila miranda]
>gi|1036836205|ref|XP_017138496.1| PREDICTED: uncharacterized protein LOC108153190 [Drosophila
>gi|1036761717|ref|XP_017156974.1| PREDICTED: protein TonB-like [Drosophila miranda]
>gi|1036838351|ref|XP_017138624.1| PREDICTED: protein trachealess-like [Drosophila miranda]
>gi|1036838239|ref|XP_017138618.1| PREDICTED: protein trachealess isoform X2 [Drosophila miran
>gi|1036838258|ref|XP_017138619.1| PREDICTED: protein trachealess isoform X3 [Drosophila miran
>gi|1036838220|ref|XP_017138617.1| PREDICTED: protein trachealess isoform X1 [Drosophila miran
>gi|1036838332|ref|XP_017138623.1| PREDICTED: protein trachealess isoform X7 [Drosophila miran
>gi|1036838296|ref|XP_017138621.1| PREDICTED: protein trachealess isoform X5 [Drosophila miran
>gi|1036838313|ref|XP_017138622.1| PREDICTED: protein trachealess isoform X6 [Drosophila miran
>gi|1036838277|ref|XP_017138620.1| PREDICTED: protein trachealess isoform X4 [Drosophila miran
>gi|1036774020|ref|XP_017135001.1| PREDICTED: uncharacterized protein LOC108151110 isoform X2
>gi|1036774039|ref|XP_017135002.1| PREDICTED: uncharacterized protein LOC108151110 isoform X3
>gi|1036774001|ref|XP_017135000.1| PREDICTED: uncharacterized protein LOC108151110 isoform X1
>gi|1036774058|ref|XP_017135003.1| PREDICTED: uncharacterized protein LOC108151110 isoform X4
>gi|1036774077|ref|XP_017135004.1| PREDICTED: nuclear anchorage protein 1 isoform X5 [Drosophil
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>gi|1036774114|ref|XP_017135006.1| PREDICTED: uncharacterized protein LOC108151112 [Drosophila
>gi|1036758187|ref|XP_017156762.1| PREDICTED: uncharacterized protein LOC108165234, partial [Dr
>gi|1036774095|ref|XP_017135005.1| PREDICTED: uncharacterized protein LOC108151111 [Drosophila
>gi|1036842071|ref|XP_017138827.1| PREDICTED: uncharacterized protein C17orf59 homolog [Drosoph
>gi|1036842090|ref|XP_017138828.1| PREDICTED: peptidyl-prolyl cis-trans isomerase-like 1 [Droso
>gi|1036847703|ref|XP_017139129.1| PREDICTED: BTB/POZ domain-containing adapter for CUL3-mediat
>gi|1036811761|ref|XP_017137136.1| PREDICTED: LOW QUALITY PROTEIN: homeodomain-interacting prot
>gi|1036823205|ref|XP_017137765.1| PREDICTED: probable protein phosphatase 2C T23F11.1 [Drosoph
>gi|1036838537|ref|XP_017138635.1| PREDICTED: probable protein phosphatase 2C T23F11.1 [Drosoph
>gi|1036838518|ref|XP_017138634.1| PREDICTED: zinc finger protein 333 [Drosophila miranda]
>gi|1036807662|ref|XP_017136913.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036807681|ref|XP_017136914.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036807718|ref|XP_017136917.1| PREDICTED: SEC14-like protein 2 isoform X2 [Drosophila miran
>gi|1036807699|ref|XP_017136916.1| PREDICTED: SEC14-like protein 2 isoform X1 [Drosophila miran
>gi|1036852268|ref|XP_017139401.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036852284|ref|XP_017139402.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036852303|ref|XP_017139403.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036761736|ref|XP_017156975.1| PREDICTED: DC-STAMP domain-containing protein 2 [Drosophila
>gi|1036851707|ref|XP_017139365.1| PREDICTED: uncharacterized protein LOC108153746 [Drosophila
>gi|1036851599|ref|XP_017139358.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036850984|ref|XP_017139320.1| PREDICTED: cohesin subunit SA-1 [Drosophila miranda]
>gi|1036851001|ref|XP_017139321.1| PREDICTED: cohesin subunit SA-1 [Drosophila miranda]
>gi|1036841669|ref|XP_017138813.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036767373|ref|XP_017134626.1| PREDICTED: N-alpha-acetyltransferase 30 [Drosophila miranda]
>gi|1036829347|ref|XP_017138113.1| PREDICTED: protein rhomboid isoform X1 [Drosophila miranda]
>gi|1036829423|ref|XP_017138117.1| PREDICTED: protein rhomboid isoform X4 [Drosophila miranda]
>gi|1036829366|ref|XP_017138114.1| PREDICTED: protein rhomboid isoform X2 [Drosophila miranda]
>gi|1036829385|ref|XP_017138115.1| PREDICTED: protein rhomboid isoform X3 [Drosophila miranda]
>gi|1036829404|ref|XP_017138116.1| PREDICTED: protein rhomboid isoform X3 [Drosophila miranda]
>gi|1036831592|ref|XP_017138242.1| PREDICTED: protein zer-1 homolog isoform X2 [Drosophila mira
>gi|1036831533|ref|XP_017138239.1| PREDICTED: protein zer-1 homolog isoform X1 [Drosophila mira
>gi|1036831554|ref|XP_017138240.1| PREDICTED: protein zer-1 homolog isoform X1 [Drosophila mira
>gi|1036831573|ref|XP_017138241.1| PREDICTED: protein zer-1 homolog isoform X1 [Drosophila mira
>gi|1036825280|ref|XP_017137881.1| PREDICTED: poly(U)-binding-splicing factor half pint [Drosop
>gi|1036791210|ref|XP_017135990.1| PREDICTED: cell division cycle 5-like protein [Drosophila mi
>gi|1036791247|ref|XP_017135992.1| PREDICTED: RING-box protein 1B [Drosophila miranda]
>gi|1036791229|ref|XP_017135991.1| PREDICTED: uncharacterized protein LOC108151725 [Drosophila
>gi|1036791191|ref|XP_017135988.1| PREDICTED: bifunctional heparan sulfate N-deacetylase/N-sul
>gi|1036799780|ref|XP_017136473.1| PREDICTED: vacuolar protein sorting-associated protein 8 hom
>gi|1036799888|ref|XP_017136478.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036799817|ref|XP_017136475.1| PREDICTED: uncharacterized protein C18orf8 isoform X2 [Droso
>gi|1036799835|ref|XP_017136476.1| PREDICTED: uncharacterized protein C18orf8 isoform X3 [Droso
>gi|1036799798|ref|XP_017136474.1| PREDICTED: uncharacterized protein C18orf8 isoform X1 [Droso
>gi|1036799853|ref|XP_017136477.1| PREDICTED: zinc finger protein 93 [Drosophila miranda]
>gi|1036805182|ref|XP_017136778.1| PREDICTED: zinc finger protein 253-like isoform X1 [Drosophi
>gi|1036805202|ref|XP_017136779.1| PREDICTED: zinc finger protein 675-like isoform X2 [Drosophi
>gi|1036805220|ref|XP_017136780.1| PREDICTED: uncharacterized protein LOC108152169 [Drosophila
>gi|1036805239|ref|XP_017136782.1| PREDICTED: ubiquitin-conjugating enzyme E2 D4-like [Drosophi
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>gi|1036841716|ref|XP_017138817.1| PREDICTED: putative cyclin-dependent serine/threonine-prote
>gi|1036841731|ref|XP_017138818.1| PREDICTED: uncharacterized protein LOC108153362 [Drosophila
>gi|1036761754|ref|XP_017156976.1| PREDICTED: odorant receptor 65a-like [Drosophila miranda]
>gi|1036761770|ref|XP_017156977.1| PREDICTED: ubiquitin-conjugating enzyme E2 D4-like, partial
>gi|1036766030|ref|XP_017134553.1| PREDICTED: collagenase-like [Drosophila miranda]
>gi|1036761788|ref|XP_017156978.1| PREDICTED: uncharacterized protein LOC108165452 [Drosophila
>gi|1036761809|ref|XP_017156980.1| PREDICTED: protein phosphatase inhibitor 2-like [Drosophila
>gi|1036761827|ref|XP_017156981.1| PREDICTED: protein phosphatase inhibitor 2-like [Drosophila
>gi|1036847347|ref|XP_017139110.1| PREDICTED: protein phosphatase inhibitor 2-like isoform X3
>gi|1036847309|ref|XP_017139108.1| PREDICTED: protein phosphatase inhibitor 2-like isoform X1
>gi|1036847328|ref|XP_017139109.1| PREDICTED: protein phosphatase inhibitor 2-like isoform X2
>gi|1036762929|ref|XP_017157042.1| PREDICTED: L-lactate dehydrogenase [Drosophila miranda]
>gi|1036762908|ref|XP_017157041.1| PREDICTED: phospholipase A1 member A [Drosophila miranda]
>gi|1036761847|ref|XP_017156982.1| PREDICTED: uncharacterized protein LOC108165455 [Drosophila
>gi|1036761865|ref|XP_017156983.1| PREDICTED: uncharacterized protein LOC108165456 [Drosophila
>gi|1036762659|ref|XP_017157026.1| PREDICTED: multidrug resistance protein homolog 65 [Drosoph
>gi|1036762640|ref|XP_017157025.1| PREDICTED: multidrug resistance protein homolog 65 [Drosoph
>gi|1036762995|ref|XP_017157046.1| PREDICTED: uncharacterized protein LOC108165499 [Drosophila
>gi|1036763012|ref|XP_017157047.1| PREDICTED: uncharacterized protein LOC108165500 [Drosophila
>gi|1036762956|ref|XP_017157043.1| PREDICTED: uncharacterized protein LOC108165497 [Drosophila
>gi|1036763031|ref|XP_017157048.1| PREDICTED: uncharacterized protein LOC108165501 [Drosophila
>gi|1036762574|ref|XP_017157021.1| PREDICTED: laminin subunit alpha [Drosophila miranda]
>gi|1036762676|ref|XP_017157027.1| PREDICTED: transportin-1 [Drosophila miranda]
>gi|1036762975|ref|XP_017157045.1| PREDICTED: splicing factor 3B subunit 6-like protein [Droso
>gi|1036762731|ref|XP_017157030.1| PREDICTED: zinc finger protein 879 isoform X2 [Drosophila m
>gi|1036762712|ref|XP_017157029.1| PREDICTED: zinc finger protein 708 isoform X1 [Drosophila m
>gi|1036762694|ref|XP_017157028.1| PREDICTED: zinc finger protein 668 [Drosophila miranda]
>gi|1036762750|ref|XP_017157031.1| PREDICTED: protein ERGIC-53 [Drosophila miranda]
>gi|1036762593|ref|XP_017157022.1| PREDICTED: talin-2 isoform X1 [Drosophila miranda]
>gi|1036762603|ref|XP_017157023.1| PREDICTED: talin-2 isoform X2 [Drosophila miranda]
>gi|1036762620|ref|XP_017157024.1| PREDICTED: ATP-dependent DNA helicase Q4 [Drosophila miranda
>gi|1036762789|ref|XP_017157034.1| PREDICTED: glycylpeptide N-tetradecanoyltransferase [Drosoph
>gi|1036762804|ref|XP_017157035.1| PREDICTED: UDP-glucuronic acid decarboxylase 1 [Drosophila
>gi|1036762773|ref|XP_017157033.1| PREDICTED: steroid hormone receptor ERR2 [Drosophila miranda
>gi|1036762827|ref|XP_017157036.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762841|ref|XP_017157037.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762875|ref|XP_017157039.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762859|ref|XP_017157038.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762890|ref|XP_017157040.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036761883|ref|XP_017156984.1| PREDICTED: chaperone protein DnaJ, partial [Drosophila miran
>gi|1036815069|ref|XP_017137315.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036815088|ref|XP_017137316.1| PREDICTED: nedd8-conjugating enzyme Ubc12 [Drosophila miran
>gi|1036839119|ref|XP_017138673.1| PREDICTED: probable deoxyhypusine synthase isoform X1 [Dros
>gi|1036839138|ref|XP_017138674.1| PREDICTED: probable deoxyhypusine synthase isoform X2 [Dros
>gi|1036849321|ref|XP_017139221.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036763387|ref|XP_017157063.1| PREDICTED: uncharacterized protein LOC108165517 [Drosophila
>gi|1036850837|ref|XP_017139313.1| PREDICTED: uncharacterized protein LOC108153698 isoform X1
>gi|1036850856|ref|XP_017139314.1| PREDICTED: uncharacterized protein LOC108153698 isoform X2
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>gi|1036766424|ref|XP_017134573.1| PREDICTED: uncharacterized protein LOC108150784 [Drosophila
>gi|1036850763|ref|XP_017139308.1| PREDICTED: pupal cuticle protein C1B [Drosophila miranda]
>gi|1036850782|ref|XP_017139309.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mi
>gi|1036850746|ref|XP_017139307.1| PREDICTED: eukaryotic translation initiation factor 4E [Dros
>gi|1036765435|ref|XP_017157170.1| PREDICTED: uncharacterized protein LOC108165606 [Drosophila
>gi|1036765453|ref|XP_017157171.1| PREDICTED: uncharacterized protein LOC108165606 [Drosophila
>gi|1036847439|ref|XP_017139116.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036847458|ref|XP_017139117.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036847476|ref|XP_017139118.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036784789|ref|XP_017135625.1| PREDICTED: collagen type IV alpha-3-binding protein [Drosopl
>gi|1036784805|ref|XP_017135626.1| PREDICTED: uncharacterized protein LOC108151498 [Drosophila
>gi|1036784879|ref|XP_017135631.1| PREDICTED: ras-related protein Rab-43 isoform X1 [Drosophila
>gi|1036784898|ref|XP_017135632.1| PREDICTED: ras-related protein Rab-43 isoform X2 [Drosophila
>gi|1036784919|ref|XP_017135633.1| PREDICTED: ADP-ribosylation factor-like protein 5A [Drosophi
>gi|1036784772|ref|XP_017135624.1| PREDICTED: uncharacterized protein LOC108151495 [Drosophila
>gi|1036784823|ref|XP_017135627.1| PREDICTED: selenocysteine insertion sequence-binding protei
>gi|1036761902|ref|XP_017156985.1| PREDICTED: gustatory receptor for bitter taste 66a [Drosophi
>gi|1036784842|ref|XP_017135629.1| PREDICTED: bax inhibitor 1 isoform X1 [Drosophila miranda]
>gi|1036784861|ref|XP_017135630.1| PREDICTED: bax inhibitor 1 isoform X2 [Drosophila miranda]
>gi|1036823396|ref|XP_017137776.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036823377|ref|XP_017137775.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036823358|ref|XP_017137774.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036823337|ref|XP_017137773.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036848334|ref|XP_017139164.1| PREDICTED: alkaline phosphatase [Drosophila miranda]
>gi|1036838391|ref|XP_017138626.1| PREDICTED: syntaxin-7 [Drosophila miranda]
>gi|1036846454|ref|XP_017139058.1| PREDICTED: fork head domain-containing protein crocodile [D
>gi|1036847123|ref|XP_017139097.1| PREDICTED: forkhead box protein biniou [Drosophila miranda]
>gi|1036761920|ref|XP_017156986.1| PREDICTED: keratin-associated protein 5-1-like [Drosophila n
>gi|1036764298|ref|XP_017157113.1| PREDICTED: sperm-specific protein Don juan [Drosophila miran
>gi|1036764318|ref|XP_017157114.1| PREDICTED: sperm-specific protein Don juan [Drosophila miran
>gi|1036809371|ref|XP_017137004.1| PREDICTED: G protein alpha i subunit [Drosophila miranda]
>gi|1036809390|ref|XP_017137005.1| PREDICTED: uncharacterized protein LOC108152294 [Drosophila
>gi|1036809352|ref|XP_017137003.1| PREDICTED: putative serine protease F56F10.1 [Drosophila mi
>gi|1036809335|ref|XP_017137002.1| PREDICTED: cilia- and flagella-associated protein 52 [Drosop
>gi|1036784699|ref|XP_017135620.1| PREDICTED: UDP-glucose 6-dehydrogenase [Drosophila miranda]
>gi|1036784717|ref|XP_017135621.1| PREDICTED: SURF1-like protein [Drosophila miranda]
>gi|1036784736|ref|XP_017135622.1| PREDICTED: ubiquinol-cytochrome-c reductase complex assembly
>gi|1036784753|ref|XP_017135623.1| PREDICTED: transcription factor Adf-1 [Drosophila miranda]
>gi|1036784680|ref|XP_017135619.1| PREDICTED: CLIP-associating protein [Drosophila miranda]
>gi|1036825221|ref|XP_017137877.1| PREDICTED: LOW QUALITY PROTEIN: asparagine synthetase B [glu
>gi|1036825240|ref|XP_017137878.1| PREDICTED: asparagine synthetase [glutamine-hydrolyzing] 2-1
>gi|1036829537|ref|XP_017138121.1| PREDICTED: uncharacterized protein LOC108152938 [Drosophila
>gi|1036829476|ref|XP_017138118.1| PREDICTED: adenylate cyclase type 8 isoform X1 [Drosophila n
>gi|1036829497|ref|XP_017138119.1| PREDICTED: adenylate cyclase type 8 isoform X2 [Drosophila n
>gi|1036829516|ref|XP_017138120.1| PREDICTED: adenylate cyclase type 8 isoform X3 [Drosophila n
>gi|1036805107|ref|XP_017136773.1| PREDICTED: GTPase-activating protein isoform X1 [Drosophila
>gi|1036805127|ref|XP_017136774.1| PREDICTED: GTPase-activating protein isoform X2 [Drosophila
>gi|1036786635|ref|XP_017135730.1| PREDICTED: uncharacterized protein LOC108151570 isoform X1
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>gi|1036786656|ref|XP_017135731.1| PREDICTED: myotubularin-related protein 9-like isoform X2 [D
>gi|1036786675|ref|XP_017135732.1| PREDICTED: myotubularin-related protein 9-like isoform X3 [D
>gi|1036786694|ref|XP_017135733.1| PREDICTED: ATP-binding cassette sub-family E member 1 [Drosc
>gi|1036786713|ref|XP_017135734.1| PREDICTED: uncharacterized protein LOC108151573 [Drosophila
>gi|1036786791|ref|XP_017135739.1| PREDICTED: uncharacterized protein LOC108151575 [Drosophila
>gi|1036786810|ref|XP_017135740.1| PREDICTED: uncharacterized protein LOC108151575 [Drosophila
>gi|1036786734|ref|XP_017135735.1| PREDICTED: N-acetylgalactosamine kinase [Drosophila miranda]
>gi|1036786753|ref|XP_017135736.1| PREDICTED: N-acetylgalactosamine kinase [Drosophila miranda]
>gi|1036786772|ref|XP_017135738.1| PREDICTED: N-acetylgalactosamine kinase [Drosophila miranda]
>gi|1036800456|ref|XP_017136510.1| PREDICTED: vesicle transport protein USE1 isoform X1 [Drosop
>gi|1036800474|ref|XP_017136511.1| PREDICTED: vesicle transport protein USE1 isoform X2 [Drosop
>gi|1036800492|ref|XP_017136512.1| PREDICTED: dihydropteridine reductase [Drosophila miranda]
>gi|1036800271|ref|XP_017136500.1| PREDICTED: protein boule isoform X1 [Drosophila miranda]
>gi|1036800290|ref|XP_017136501.1| PREDICTED: protein boule isoform X1 [Drosophila miranda]
>gi|1036800309|ref|XP_017136502.1| PREDICTED: protein boule isoform X2 [Drosophila miranda]
>gi|1036800327|ref|XP_017136503.1| PREDICTED: protein boule isoform X3 [Drosophila miranda]
>gi|1036800344|ref|XP_017136504.1| PREDICTED: protein boule isoform X4 [Drosophila miranda]
>gi|1036800362|ref|XP_017136505.1| PREDICTED: protein boule isoform X5 [Drosophila miranda]
>gi|1036800382|ref|XP_017136506.1| PREDICTED: protein boule isoform X6 [Drosophila miranda]
>gi|1036800401|ref|XP_017136507.1| PREDICTED: protein boule isoform X7 [Drosophila miranda]
>gi|1036800418|ref|XP_017136508.1| PREDICTED: protein boule isoform X8 [Drosophila miranda]
>gi|1036800437|ref|XP_017136509.1| PREDICTED: protein boule isoform X9 [Drosophila miranda]
>gi|1036848370|ref|XP_017139166.1| PREDICTED: uncharacterized protein LOC108153596 [Drosophila
>gi|1036850361|ref|XP_017139284.1| PREDICTED: uncharacterized protein LOC108153678 [Drosophila
>gi|1036818582|ref|XP_017137514.1| PREDICTED: uncharacterized protein LOC108152581 isoform X4
>gi|1036818563|ref|XP_017137513.1| PREDICTED: uncharacterized protein LOC108152581 isoform X3
>gi|1036818525|ref|XP_017137511.1| PREDICTED: uncharacterized protein LOC108152581 isoform X1
>gi|1036818544|ref|XP_017137512.1| PREDICTED: uncharacterized protein LOC108152581 isoform X2
>gi|1036818603|ref|XP_017137515.1| PREDICTED: uncharacterized protein LOC108152581 isoform X5
>gi|1036818622|ref|XP_017137516.1| PREDICTED: mitochondrial inner membrane protein COX18 [Drosc
>gi|1036786319|ref|XP_017135714.1| PREDICTED: F-box only protein 39 [Drosophila miranda]
>gi|1036786266|ref|XP_017135710.1| PREDICTED: arginine kinase isoform X1 [Drosophila miranda]
>gi|1036786285|ref|XP_017135711.1| PREDICTED: arginine kinase isoform X2 [Drosophila miranda]
>gi|1036786302|ref|XP_017135712.1| PREDICTED: arginine kinase isoform X3 [Drosophila miranda]
>gi|1036786338|ref|XP_017135715.1| PREDICTED: arginine kinase-like [Drosophila miranda]
>gi|1036841632|ref|XP_017138811.1| PREDICTED: T-box transcription factor TBX6 [Drosophila miran
>gi|1036838591|ref|XP_017138638.1| PREDICTED: T-box protein 2 [Drosophila miranda]
>gi|1036845704|ref|XP_017139016.1| PREDICTED: T-box transcription factor TBX6 [Drosophila miran
>gi|1036828014|ref|XP_017138036.1| PREDICTED: ubiquitin-protein ligase E3B [Drosophila miranda]
>gi|1036810809|ref|XP_017137085.1| PREDICTED: protein Smaug isoform X2 [Drosophila miranda]
>gi|1036810828|ref|XP_017137086.1| PREDICTED: protein Smaug isoform X3 [Drosophila miranda]
>gi|1036810771|ref|XP_017137083.1| PREDICTED: protein Smaug isoform X1 [Drosophila miranda]
>gi|1036810789|ref|XP_017137084.1| PREDICTED: protein Smaug isoform X1 [Drosophila miranda]
>gi|1036835814|ref|XP_017138472.1| PREDICTED: UPF0193 protein EVG1 homolog [Drosophila miranda]
>gi|1036835795|ref|XP_017138471.1| PREDICTED: protein phosphatase methylesterase 1 [Drosophila
>gi|1036843002|ref|XP_017138879.1| PREDICTED: uncharacterized protein LOC108153400 isoform X1
>gi|1036843021|ref|XP_017138880.1| PREDICTED: uncharacterized protein LOC108153400 isoform X1
>gi|1036843041|ref|XP_017138881.1| PREDICTED: uncharacterized protein LOC108153400 isoform X2
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>gi|1036822080|ref|XP_017137700.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X1
>gi|1036822098|ref|XP_017137702.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X1
>gi|1036822116|ref|XP_017137703.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X1
>gi|1036822134|ref|XP_017137704.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X2
>gi|1036822174|ref|XP_017137706.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X4
>gi|1036822193|ref|XP_017137707.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X5
>gi|1036822153|ref|XP_017137705.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X3
>gi|1036822212|ref|XP_017137708.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X6
>gi|1036822231|ref|XP_017137709.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X7
>gi|1036828491|ref|XP_017138064.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828414|ref|XP_017138059.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828726|ref|XP_017138076.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828510|ref|XP_017138065.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828432|ref|XP_017138060.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828609|ref|XP_017138070.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828569|ref|XP_017138068.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828550|ref|XP_017138067.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828451|ref|XP_017138062.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828357|ref|XP_017138056.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828376|ref|XP_017138057.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828688|ref|XP_017138074.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828590|ref|XP_017138069.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828470|ref|XP_017138063.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828395|ref|XP_017138058.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828531|ref|XP_017138066.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828630|ref|XP_017138071.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828648|ref|XP_017138072.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828669|ref|XP_017138073.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828707|ref|XP_017138075.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036834176|ref|XP_017138379.1| PREDICTED: proton-associated sugar transporter A [Drosophila
>gi|1036834195|ref|XP_017138380.1| PREDICTED: proton-associated sugar transporter A [Drosophila
>gi|1036835275|ref|XP_017138440.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036847722|ref|XP_017139130.1| PREDICTED: glutamate receptor 1 [Drosophila miranda]
>gi|1036850691|ref|XP_017139304.1| PREDICTED: peptidoglycan-recognition protein LA isoform X1
>gi|1036850729|ref|XP_017139306.1| PREDICTED: peptidoglycan-recognition protein LA isoform X3
>gi|1036850710|ref|XP_017139305.1| PREDICTED: peptidoglycan-recognition protein LA isoform X2
>gi|1036803078|ref|XP_017136658.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803097|ref|XP_017136660.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803116|ref|XP_017136661.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803134|ref|XP_017136662.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803152|ref|XP_017136663.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803190|ref|XP_017136665.1| PREDICTED: peptidoglycan-recognition protein LC isoform X3
>gi|1036803228|ref|XP_017136667.1| PREDICTED: peptidoglycan-recognition protein LC isoform X5
>gi|1036803209|ref|XP_017136666.1| PREDICTED: peptidoglycan-recognition protein LC isoform X4
>gi|1036803171|ref|XP_017136664.1| PREDICTED: peptidoglycan-recognition protein LC isoform X2
>gi|1036803246|ref|XP_017136668.1| PREDICTED: peptidoglycan-recognition protein LC isoform X6
>gi|1036803265|ref|XP_017136669.1| PREDICTED: peptidoglycan-recognition protein LC isoform X7
>gi|1036803355|ref|XP_017136675.1| PREDICTED: peptidoglycan-recognition protein LF isoform X1
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>gi|1036803371|ref|XP_017136676.1| PREDICTED: peptidoglycan-recognition protein LF isoform X2
>gi|1036803338|ref|XP_017136674.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803322|ref|XP_017136673.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803303|ref|XP_017136672.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803284|ref|XP_017136671.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803388|ref|XP_017136677.1| PREDICTED: uncharacterized protein LOC108152112 [Drosophila
>gi|1036804590|ref|XP_017136744.1| PREDICTED: small VCP/p97-interacting protein [Drosophila mi
>gi|1036804572|ref|XP_017136743.1| PREDICTED: autophagy-related protein 2 homolog A [Drosophila
>gi|1036821275|ref|XP_017137656.1| PREDICTED: uncharacterized protein LOC108152670 [Drosophila
>gi|1036821256|ref|XP_017137655.1| PREDICTED: putative adenosylhomocysteinase 2 [Drosophila mi
>gi|1036836040|ref|XP_017138486.1| PREDICTED: upstream activation factor subunit spp27 [Drosop
>gi|1036802282|ref|XP_017136617.1| PREDICTED: histone-lysine N-methyltransferase 2D [Drosophila
>gi|1036837979|ref|XP_017138602.1| PREDICTED: regulating synaptic membrane exocytosis protein
>gi|1036837998|ref|XP_017138603.1| PREDICTED: regulating synaptic membrane exocytosis protein
>gi|1036838017|ref|XP_017138604.1| PREDICTED: regulating synaptic membrane exocytosis protein
>gi|1036838069|ref|XP_017138608.1| PREDICTED: synaptic vesicle glycoprotein 2B-like [Drosophila
>gi|1036838051|ref|XP_017138607.1| PREDICTED: putative transporter svop-1 [Drosophila miranda]
>gi|1036838034|ref|XP_017138605.1| PREDICTED: putative transporter SVOPL [Drosophila miranda]
>gi|1036764224|ref|XP_017157109.1| PREDICTED: uncharacterized protein LOC108165545 [Drosophila
>gi|1037099942|ref|XP_017144749.1| PREDICTED: extended synaptotagmin-2 isoform X2 [Drosophila
>gi|1037099932|ref|XP_017144747.1| PREDICTED: extended synaptotagmin-2 isoform X1 [Drosophila
>gi|1037099960|ref|XP_017144751.1| PREDICTED: extended synaptotagmin-1 isoform X4 [Drosophila
>gi|1037099950|ref|XP_017144750.1| PREDICTED: extended synaptotagmin-2 isoform X3 [Drosophila
>gi|1037079607|ref|XP_017143172.1| PREDICTED: protein CREBRF homolog [Drosophila miranda]
>gi|1037079621|ref|XP_017143173.1| PREDICTED: protein CREBRF homolog [Drosophila miranda]
>gi|1037079635|ref|XP_017143174.1| PREDICTED: BUD13 homolog [Drosophila miranda]
>gi|1037105933|ref|XP_017145427.1| PREDICTED: uncharacterized protein LOC108157757 [Drosophila
>gi|1037034935|ref|XP_017140187.1| PREDICTED: intraflagellar transport protein 56 [Drosophila
>gi|1037085350|ref|XP_017143610.1| PREDICTED: uncharacterized protein LOC108156584 [Drosophila
>gi|1037033899|ref|XP_017140125.1| PREDICTED: mitochondrial import receptor subunit TOM40 homol
>gi|1037034305|ref|XP_017140151.1| PREDICTED: uncharacterized protein LOC108154396 [Drosophila
>gi|1037034760|ref|XP_017140178.1| PREDICTED: tetratricopeptide repeat protein 12 [Drosophila
>gi|1037022592|ref|XP_017139431.1| PREDICTED: uncharacterized protein LOC108153843 [Drosophila
>gi|1037037994|ref|XP_017140356.1| PREDICTED: tetratricopeptide repeat protein 12 [Drosophila
>gi|1037105894|ref|XP_017145421.1| PREDICTED: sodium-independent sulfate anion transporter [Dro
>gi|1037103816|ref|XP_017145131.1| PREDICTED: uncharacterized protein LOC108157539 [Drosophila
>gi|1037034339|ref|XP_017140153.1| PREDICTED: uncharacterized protein LOC108154398 [Drosophila
>gi|1037090337|ref|XP_017143965.1| PREDICTED: uncharacterized protein LOC108156797 [Drosophila
>gi|1037093993|ref|XP_017144226.1| PREDICTED: LOW QUALITY PROTEIN: xanthine dehydrogenase [Dros
>gi|1037022607|ref|XP_017139432.1| PREDICTED: xanthine dehydrogenase/oxidase-like [Drosophila
>gi|1037093979|ref|XP_017144225.1| PREDICTED: indole-3-acetaldehyde oxidase [Drosophila miranda
>gi|1037092203|ref|XP_017144105.1| PREDICTED: LOW QUALITY PROTEIN: indole-3-acetaldehyde oxidas
>gi|1037092217|ref|XP_017144107.1| PREDICTED: xanthine dehydrogenase-like [Drosophila miranda]
>gi|1037037448|ref|XP_017140328.1| PREDICTED: uncharacterized protein LOC108154536 [Drosophila
>gi|1037033882|ref|XP_017140124.1| PREDICTED: uncharacterized protein LOC108154370 [Drosophila
>gi|1037105540|ref|XP_017145373.1| PREDICTED: uncharacterized protein LOC108157713 [Drosophila
>gi|1037105114|ref|XP_017145311.1| PREDICTED: uncharacterized protein LOC108157663 [Drosophila
>gi|1037103621|ref|XP_017145105.1| PREDICTED: dynein beta chain, ciliary isoform X2 [Drosophila
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>gi|1037103613|ref|XP_017145104.1| PREDICTED: dynein beta chain, ciliary isoform X1 [Drosophila
>gi|1037022621|ref|XP_017139433.1| PREDICTED: uncharacterized protein LOC108153845 [Drosophila
>gi|1037035151|ref|XP_017140200.1| PREDICTED: uncharacterized protein LOC108154438 [Drosophila
>gi|1037034409|ref|XP_017140157.1| PREDICTED: uncharacterized protein LOC108154401 isoform X1
>gi|1037034427|ref|XP_017140159.1| PREDICTED: uncharacterized protein LOC108154401 isoform X2
>gi|1037034447|ref|XP_017140160.1| PREDICTED: uncharacterized protein LOC108154401 isoform X3
>gi|1037032758|ref|XP_017140057.1| PREDICTED: circadian clock-controlled protein [Drosophila mi
>gi|1037034813|ref|XP_017140182.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037091349|ref|XP_017144042.1| PREDICTED: protein Malvolio isoform X2 [Drosophila miranda]
>gi|1037091335|ref|XP_017144041.1| PREDICTED: protein Malvolio isoform X1 [Drosophila miranda]
>gi|1037093883|ref|XP_017144218.1| PREDICTED: hematopoietic lineage cell-specific protein [Dros
>gi|1037093897|ref|XP_017144219.1| PREDICTED: hematopoietic lineage cell-specific protein [Dros
>gi|1037082949|ref|XP_017143427.1| PREDICTED: annexin B9 isoform X3 [Drosophila miranda]
>gi|1037082937|ref|XP_017143426.1| PREDICTED: annexin B9 isoform X2 [Drosophila miranda]
>gi|1037082910|ref|XP_017143424.1| PREDICTED: annexin B9 isoform X1 [Drosophila miranda]
>gi|1037082924|ref|XP_017143425.1| PREDICTED: annexin B9 isoform X1 [Drosophila miranda]
>gi|1037082896|ref|XP_017143422.1| PREDICTED: uridine 5'-monophosphate synthase [Drosophila mi
>gi|1037034221|ref|XP_017140144.1| PREDICTED: doublesex- and mab-3-related transcription facto
>gi|1037033178|ref|XP_017140083.1| PREDICTED: hematopoietically-expressed homeobox protein Hhe
>gi|1037067328|ref|XP_017142329.1| PREDICTED: hornerin [Drosophila miranda]
>gi|1037067344|ref|XP_017142330.1| PREDICTED: uncharacterized protein LOC108155781 [Drosophila
>gi|1037085504|ref|XP_017143620.1| PREDICTED: hematological and neurological expressed 1-like
>gi|1037085490|ref|XP_017143619.1| PREDICTED: THO complex subunit 4 [Drosophila miranda]
>gi|1037092245|ref|XP_017144109.1| PREDICTED: dephospho-CoA kinase [Drosophila miranda]
>gi|1037092231|ref|XP_017144108.1| PREDICTED: UTP:RNA uridylyltransferase 1 [Drosophila mirand
>gi|1037085365|ref|XP_017143611.1| PREDICTED: serine/threonine-protein kinase/endoribonuclease
>gi|1037085989|ref|XP_017143655.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085933|ref|XP_017143651.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037085921|ref|XP_017143650.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037086000|ref|XP_017143656.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085961|ref|XP_017143653.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085947|ref|XP_017143652.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085975|ref|XP_017143654.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085907|ref|XP_017143649.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037085893|ref|XP_017143647.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037086014|ref|XP_017143657.1| PREDICTED: uncharacterized protein LOC108156613 [Drosophila
>gi|1037085865|ref|XP_017143645.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dr
>gi|1037085879|ref|XP_017143646.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dr
>gi|1037106464|ref|XP_017145500.1| PREDICTED: 4-coumarate--CoA ligase-like 7 [Drosophila miran
>gi|1037106181|ref|XP_017145461.1| PREDICTED: 4-coumarate--CoA ligase-like [Drosophila miranda
>gi|1037033750|ref|XP_017140115.1| PREDICTED: luciferin 4-monooxygenase [Drosophila miranda]
>gi|1037105779|ref|XP_017145406.1| PREDICTED: luciferin 4-monooxygenase-like [Drosophila miran
>gi|1037096331|ref|XP_017144415.1| PREDICTED: glutamate-gated chloride channel isoform X8 [Dros
>gi|1037096303|ref|XP_017144413.1| PREDICTED: glutamate-gated chloride channel isoform X6 [Dros
>gi|1037096252|ref|XP_017144409.1| PREDICTED: glutamate-gated chloride channel isoform X2 [Dros
>gi|1037096343|ref|XP_017144416.1| PREDICTED: glutamate-gated chloride channel isoform X9 [Dros
>gi|1037096291|ref|XP_017144412.1| PREDICTED: glutamate-gated chloride channel isoform X5 [Dros
>gi|1037096281|ref|XP_017144411.1| PREDICTED: glutamate-gated chloride channel isoform X4 [Dros
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>gi|1037096226|ref|XP_017144406.1| PREDICTED: glutamate-gated chloride channel isoform X1 [Dros
>gi|1037096238|ref|XP_017144407.1| PREDICTED: glutamate-gated chloride channel isoform X1 [Dros
>gi|1037096315|ref|XP_017144414.1| PREDICTED: glutamate-gated chloride channel isoform X7 [Dros
>gi|1037096355|ref|XP_017144417.1| PREDICTED: glutamate-gated chloride channel isoform X10 [Dro
>gi|1037096270|ref|XP_017144410.1| PREDICTED: glutamate-gated chloride channel isoform X3 [Dros
>gi|1037038914|ref|XP_017140416.1| PREDICTED: uncharacterized protein LOC108154611 [Drosophila
>gi|1037096990|ref|XP_017144474.1| PREDICTED: pyrazinamidase/nicotinamidase [Drosophila miranda
>gi|1037097002|ref|XP_017144475.1| PREDICTED: pyrazinamidase/nicotinamidase [Drosophila miranda
>gi|1037096978|ref|XP_017144473.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1037077638|ref|XP_017143040.1| PREDICTED: glycine receptor subunit alpha-4 [Drosophila mira
>gi|1037077612|ref|XP_017143037.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037077649|ref|XP_017143041.1| PREDICTED: developmentally-regulated GTP-binding protein 2
>gi|1037077625|ref|XP_017143039.1| PREDICTED: probable nucleoporin Nup58 [Drosophila miranda]
>gi|1037102983|ref|XP_017145032.1| PREDICTED: V-type proton ATPase subunit G [Drosophila mirand
>gi|1037077531|ref|XP_017143032.1| PREDICTED: anoctamin-1 isoform X1 [Drosophila miranda]
>gi|1037077547|ref|XP_017143033.1| PREDICTED: anoctamin-1 isoform X2 [Drosophila miranda]
>gi|1037077559|ref|XP_017143034.1| PREDICTED: anoctamin-1 isoform X3 [Drosophila miranda]
>gi|1037077585|ref|XP_017143036.1| PREDICTED: uncharacterized protein LOC108156209 [Drosophila
>gi|1037077571|ref|XP_017143035.1| PREDICTED: organic cation transporter protein [Drosophila mi
>gi|1037022641|ref|XP_017139434.1| PREDICTED: uncharacterized protein LOC108153847 [Drosophila
>gi|1037100890|ref|XP_017144829.1| PREDICTED: organic cation transporter-like protein [Drosophi
>gi|1037106073|ref|XP_017145446.1| PREDICTED: organic cation transporter-like protein isoform
>gi|1037106065|ref|XP_017145445.1| PREDICTED: organic cation transporter-like protein isoform
>gi|1037106079|ref|XP_017145447.1| PREDICTED: uncharacterized protein LOC108157773 [Drosophila
>gi|1037047793|ref|XP_017141010.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1037047731|ref|XP_017141005.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X16 [Dros
>gi|1037047715|ref|XP_017141004.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X15 [Dros
>gi|1037047624|ref|XP_017140997.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X9 [Dros
>gi|1037047608|ref|XP_017140996.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X8 [Dros
>gi|1037047592|ref|XP_017140995.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X7 [Dros
>gi|1037047682|ref|XP_017141002.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X13 [Dros
>gi|1037047576|ref|XP_017140994.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X6 [Dros
>gi|1037047667|ref|XP_017141001.1| PREDICTED: dystrophin isoform X12 [Drosophila miranda]
>gi|1037047637|ref|XP_017140999.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X10 [Dros
>gi|1037047493|ref|XP_017140990.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X2 [Dros
>gi|1037047697|ref|XP_017141003.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X14 [Dros
>gi|1037047652|ref|XP_017141000.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X11 [Dros
>gi|1037047540|ref|XP_017140992.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X4 [Dros
>gi|1037047508|ref|XP_017140991.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X3 [Dros
>gi|1037047479|ref|XP_017140989.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X1 [Dros
>gi|1037047556|ref|XP_017140993.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X5 [Dros
>gi|1037047821|ref|XP_017141012.1| PREDICTED: uncharacterized protein LOC108154984 [Drosophila
>gi|1037047747|ref|XP_017141006.1| PREDICTED: dystrophin-like [Drosophila miranda]
>gi|1037047763|ref|XP_017141007.1| PREDICTED: uncharacterized protein LOC108154980 [Drosophila
>gi|1037047807|ref|XP_017141011.1| PREDICTED: IDLSRF-like peptide [Drosophila miranda]
>gi|1037047779|ref|XP_017141009.1| PREDICTED: uncharacterized protein LOC108154981 [Drosophila
>gi|1037055229|ref|XP_017141518.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037055247|ref|XP_017141519.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
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>gi|1037022659|ref|XP_017139435.1| PREDICTED: LOW QUALITY PROTEIN: aprataxin-like protein [Dros
>gi|1037055351|ref|XP_017141527.1| PREDICTED: LOW QUALITY PROTEIN: putative DNA helicase Ino80
>gi|1037055295|ref|XP_017141523.1| PREDICTED: uncharacterized protein LOC108155315 [Drosophila
>gi|1037055215|ref|XP_017141517.1| PREDICTED: LOW QUALITY PROTEIN: thymus-specific serine prote
>gi|1037055263|ref|XP_017141520.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037055277|ref|XP_017141521.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037055336|ref|XP_017141526.1| PREDICTED: pollen-specific leucine-rich repeat extensin-like
>gi|1037055309|ref|XP_017141524.1| PREDICTED: uncharacterized protein LOC108155316 [Drosophila
>gi|1037055321|ref|XP_017141525.1| PREDICTED: uncharacterized protein LOC108155317 [Drosophila
>gi|1037033126|ref|XP_017140080.1| PREDICTED: uncharacterized protein LOC108154329 [Drosophila
>gi|1037079796|ref|XP_017143186.1| PREDICTED: neurogenic locus protein delta [Drosophila miran
>gi|1037103661|ref|XP_017145111.1| PREDICTED: zinc finger CCCH domain-containing protein 18-li
>gi|1037103653|ref|XP_017145110.1| PREDICTED: uncharacterized protein LOC108157518 [Drosophila
>gi|1037085560|ref|XP_017143625.1| PREDICTED: uncharacterized protein LOC108156595 [Drosophila
>gi|1037085533|ref|XP_017143623.1| PREDICTED: protein unc-79 homolog isoform X2 [Drosophila mi
>gi|1037085520|ref|XP_017143622.1| PREDICTED: protein unc-79 homolog isoform X1 [Drosophila mi
>gi|1037085547|ref|XP_017143624.1| PREDICTED: uncharacterized protein LOC108156594 [Drosophila
>gi|1037105868|ref|XP_017145418.1| PREDICTED: uncharacterized protein LOC108157748 [Drosophila
>gi|1037105860|ref|XP_017145417.1| PREDICTED: neurofilament medium polypeptide [Drosophila mira
>gi|1037096899|ref|XP_017144465.1| PREDICTED: phosphatidylinositol transfer protein alpha isof
>gi|1037096873|ref|XP_017144463.1| PREDICTED: phosphatidylinositol transfer protein alpha isof
>gi|1037096887|ref|XP_017144464.1| PREDICTED: phosphatidylinositol transfer protein alpha isof
>gi|1037105243|ref|XP_017145329.1| PREDICTED: cryptochrome-1 [Drosophila miranda]
>gi|1037094670|ref|XP_017144278.1| PREDICTED: 45 kDa calcium-binding protein [Drosophila miran
>gi|1037094634|ref|XP_017144276.1| PREDICTED: BRCA1-associated protein [Drosophila miranda]
>gi|1037094680|ref|XP_017144279.1| PREDICTED: uncharacterized protein LOC108156996 [Drosophila
>gi|1037094746|ref|XP_017144283.1| PREDICTED: zinc finger protein squeeze [Drosophila miranda]
>gi|1037094760|ref|XP_017144284.1| PREDICTED: zinc finger protein squeeze [Drosophila miranda]
>gi|1037094771|ref|XP_017144285.1| PREDICTED: zinc finger protein squeeze [Drosophila miranda]
>gi|1037077224|ref|XP_017143009.1| PREDICTED: C-1-tetrahydrofolate synthase, cytoplasmic [Dros
>gi|1037077178|ref|XP_017143005.1| PREDICTED: probable methyltransferase-like protein 15 homolo
>gi|1037077208|ref|XP_017143008.1| PREDICTED: uncharacterized protein LOC108156188 [Drosophila
>gi|1037077194|ref|XP_017143006.1| PREDICTED: peptide deformylase, mitochondrial [Drosophila mi
>gi|1037101196|ref|XP_017144859.1| PREDICTED: peptide deformylase, mitochondrial [Drosophila mi
>gi|1037101134|ref|XP_017144853.1| PREDICTED: uncharacterized protein LOC108157344 [Drosophila
>gi|1037101156|ref|XP_017144855.1| PREDICTED: uncharacterized protein LOC108157346 [Drosophila
>gi|1037101144|ref|XP_017144854.1| PREDICTED: uncharacterized protein LOC108157345 [Drosophila
>gi|1037101176|ref|XP_017144857.1| PREDICTED: uncharacterized protein LOC108157348 isoform X1
>gi|1037101186|ref|XP_017144858.1| PREDICTED: uncharacterized protein LOC108157348 isoform X2
>gi|1037101166|ref|XP_017144856.1| PREDICTED: very low-density lipoprotein receptor-like [Dros
>gi|1037022676|ref|XP_017139436.1| PREDICTED: bifunctional protein FolD-like [Drosophila miran
>gi|1037103567|ref|XP_017145098.1| PREDICTED: mitochondrial import receptor subunit TOM20 homol
>gi|1037103559|ref|XP_017145097.1| PREDICTED: tachykinin-like peptides receptor 86C isoform X2
>gi|1037103551|ref|XP_017145096.1| PREDICTED: tachykinin-like peptides receptor 86C isoform X1
>gi|1037106448|ref|XP_017145498.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X1 [Dros
>gi|1037106456|ref|XP_017145499.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X2 [Dros
>gi|1037093004|ref|XP_017144158.1| PREDICTED: synaptic vesicle glycoprotein 2B [Drosophila mira
>gi|1037093018|ref|XP_017144159.1| PREDICTED: mediator of RNA polymerase II transcription subu
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>gi|1037077903|ref|XP_017143060.1| PREDICTED: uncharacterized protein LOC108156231 isoform X3
>gi|1037077875|ref|XP_017143058.1| PREDICTED: uncharacterized protein LOC108156231 isoform X1
>gi|1037077919|ref|XP_017143061.1| PREDICTED: uncharacterized protein LOC108156231 isoform X4
>gi|1037077889|ref|XP_017143059.1| PREDICTED: uncharacterized protein LOC108156231 isoform X2
>gi|1037077933|ref|XP_017143063.1| PREDICTED: general transcription factor IIF subunit 2 [Dros
>gi|1037078583|ref|XP_017143106.1| PREDICTED: alpha-1,2-mannosyltransferase ALG9 [Drosophila mi
>gi|1037078569|ref|XP_017143105.1| PREDICTED: protocadherin Fat 1 isoform X4 [Drosophila mirand
>gi|1037078539|ref|XP_017143103.1| PREDICTED: protocadherin Fat 1 isoform X2 [Drosophila mirand
>gi|1037078552|ref|XP_017143104.1| PREDICTED: protocadherin Fat 1 isoform X3 [Drosophila mirand
>gi|1037078512|ref|XP_017143102.1| PREDICTED: protocadherin Fat 1 isoform X1 [Drosophila mirand
>gi|1037078597|ref|XP_017143107.1| PREDICTED: nucleotide exchange factor SIL1 isoform X1 [Dros
>gi|1037078611|ref|XP_017143108.1| PREDICTED: nucleotide exchange factor SIL1 isoform X2 [Dros
>gi|1037078625|ref|XP_017143109.1| PREDICTED: nucleotide exchange factor SIL1 isoform X3 [Dros
>gi|1037103249|ref|XP_017145063.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037103239|ref|XP_017145061.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037101047|ref|XP_017144844.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037101037|ref|XP_017144843.1| PREDICTED: protein bag-of-marbles [Drosophila miranda]
>gi|1037093952|ref|XP_017144224.1| PREDICTED: 40S ribosomal protein S27 [Drosophila miranda]
>gi|1037093938|ref|XP_017144223.1| PREDICTED: nucleoporin Nup37 [Drosophila miranda]
>gi|1037047148|ref|XP_017140966.1| PREDICTED: E3 SUMO-protein ligase RanBP2 [Drosophila miranda
>gi|1037047194|ref|XP_017140969.1| PREDICTED: LOW QUALITY PROTEIN: transcription factor Ouib-li
>gi|1037047180|ref|XP_017140968.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037047210|ref|XP_017140970.1| PREDICTED: HIG1 domain family member 1A, mitochondrial [Dros
>gi|1037047164|ref|XP_017140967.1| PREDICTED: dihydrolipoyllysine-residue succinyltransferase o
>gi|1037067809|ref|XP_017142362.1| PREDICTED: serine/threonine-protein kinase MARK2 [Drosophila
>gi|1037067824|ref|XP_017142363.1| PREDICTED: serine/threonine-protein kinase MARK2 [Drosophila
>gi|1037067707|ref|XP_017142355.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1037067723|ref|XP_017142356.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1037067738|ref|XP_017142357.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1037067753|ref|XP_017142358.1| PREDICTED: homeobox protein prospero isoform X2 [Drosophila
>gi|1037067768|ref|XP_017142359.1| PREDICTED: homeobox protein prospero isoform X3 [Drosophila
>gi|1037067783|ref|XP_017142360.1| PREDICTED: homeobox protein prospero isoform X4 [Drosophila
>gi|1037067795|ref|XP_017142361.1| PREDICTED: homeobox protein prospero isoform X5 [Drosophila
>gi|1037057958|ref|XP_017141697.1| PREDICTED: 39S ribosomal protein L40, mitochondrial [Drosopl
>gi|1037057942|ref|XP_017141696.1| PREDICTED: prefoldin subunit 3 [Drosophila miranda]
>gi|1037057892|ref|XP_017141692.1| PREDICTED: ATP-dependent DNA helicase 2 subunit 1 [Drosophil
>gi|1037057876|ref|XP_017141691.1| PREDICTED: importin-9 [Drosophila miranda]
>gi|1037057926|ref|XP_017141694.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1037057910|ref|XP_017141693.1| PREDICTED: neither inactivation nor afterpotential protein C
>gi|1037058003|ref|XP_017141700.1| PREDICTED: D-tyrosyl-tRNA(Tyr) deacylase 1 isoform X2 [Dros
>gi|1037057987|ref|XP_017141699.1| PREDICTED: D-tyrosyl-tRNA(Tyr) deacylase 1 isoform X1 [Dros
>gi|1037057972|ref|XP_017141698.1| PREDICTED: probable ribosome biogenesis protein RLP24 [Dros
>gi|1037058789|ref|XP_017141755.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 8 [Drosophi
>gi|1037058839|ref|XP_017141758.1| PREDICTED: putative tRNA (cytidine(32)/guanosine(34)-2'-O)-n
>gi|1037058823|ref|XP_017141757.1| PREDICTED: uncharacterized protein LOC108155452 [Drosophila
>gi|1037058853|ref|XP_017141759.1| PREDICTED: ribosome production factor 2 homolog [Drosophila
>gi|1037058884|ref|XP_017141761.1| PREDICTED: uncharacterized protein LOC108155456 isoform X2
>gi|1037058869|ref|XP_017141760.1| PREDICTED: uncharacterized protein LOC108155456 isoform X1
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>gi|1037058900|ref|XP_017141762.1| PREDICTED: uncharacterized protein LOC108155456 isoform X3
>gi|1037058807|ref|XP_017141756.1| PREDICTED: ralA-binding protein 1 [Drosophila miranda]
>gi|1037022693|ref|XP_017139438.1| PREDICTED: LOW QUALITY PROTEIN: malate dehydrogenase, mitocl
>gi|1037068859|ref|XP_017142436.1| PREDICTED: retinal homeobox protein Rx [Drosophila miranda]
>gi|1037068829|ref|XP_017142434.1| PREDICTED: ribosomal protein S6 kinase delta-1 [Drosophila m
>gi|1037068845|ref|XP_017142435.1| PREDICTED: little elongation complex subunit 2 [Drosophila m
>gi|1037068904|ref|XP_017142439.1| PREDICTED: SRR1-like protein [Drosophila miranda]
>gi|1037068873|ref|XP_017142437.1| PREDICTED: coiled-coil domain-containing protein 174 [Drosop
>gi|1037068889|ref|XP_017142438.1| PREDICTED: uncharacterized protein LOC108155838 [Drosophila
>gi|1037104463|ref|XP_017145217.1| PREDICTED: hexosaminidase D [Drosophila miranda]
>gi|1037082963|ref|XP_017143428.1| PREDICTED: single-stranded DNA-binding protein 2 [Drosophila
>gi|1037062830|ref|XP_017142035.1| PREDICTED: uncharacterized protein LOC108155617 [Drosophila
>gi|1037062786|ref|XP_017142032.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1037062816|ref|XP_017142034.1| PREDICTED: proteasome assembly chaperone 2 [Drosophila miran
>gi|1037062770|ref|XP_017142031.1| PREDICTED: actin-related protein 5 [Drosophila miranda]
>gi|1037062801|ref|XP_017142033.1| PREDICTED: uncharacterized protein LOC108155615 [Drosophila
>gi|1037062754|ref|XP_017142030.1| PREDICTED: importin-13 [Drosophila miranda]
>gi|1037067872|ref|XP_017142366.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037067856|ref|XP_017142365.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037067840|ref|XP_017142364.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037067920|ref|XP_017142370.1| PREDICTED: ubiquitin-like protein 4A [Drosophila miranda]
>gi|1037067888|ref|XP_017142367.1| PREDICTED: peroxiredoxin-5, mitochondrial [Drosophila mirand
>gi|1037067904|ref|XP_017142369.1| PREDICTED: nuclear cap-binding protein subunit 2 [Drosophila
>gi|1037094430|ref|XP_017144260.1| PREDICTED: protein TAPT1 homolog [Drosophila miranda]
>gi|1037103985|ref|XP_017145153.1| PREDICTED: uncharacterized protein LOC108157558 [Drosophila
>gi|1037103969|ref|XP_017145151.1| PREDICTED: uncharacterized protein LOC108157557 isoform X2
>gi|1037103977|ref|XP_017145152.1| PREDICTED: uncharacterized protein LOC108157557 isoform X3
>gi|1037103961|ref|XP_017145150.1| PREDICTED: uncharacterized protein LOC108157557 isoform X1
>gi|1037033543|ref|XP_017140103.1| PREDICTED: uncharacterized protein LOC108154348 [Drosophila
>gi|1037098435|ref|XP_017144615.1| PREDICTED: hormone receptor 4 isoform X2 [Drosophila miranda
>gi|1037098425|ref|XP_017144614.1| PREDICTED: hormone receptor 4 isoform X1 [Drosophila miranda
>gi|1037098447|ref|XP_017144616.1| PREDICTED: alpha-protein kinase 1 isoform X3 [Drosophila mir
>gi|1037099902|ref|XP_017144744.1| PREDICTED: uncharacterized protein LOC108157267 [Drosophila
>gi|1037086058|ref|XP_017143660.1| PREDICTED: fibroblast growth factor receptor homolog 1 [Dros
>gi|1037086072|ref|XP_017143661.1| PREDICTED: fibroblast growth factor receptor homolog 1 [Dros
>gi|1037046824|ref|XP_017140944.1| PREDICTED: tubulin alpha-3 chain [Drosophila miranda]
>gi|1037046854|ref|XP_017140947.1| PREDICTED: SPRY domain-containing protein 7 [Drosophila mira
>gi|1037046840|ref|XP_017140945.1| PREDICTED: plancitoxin-1 [Drosophila miranda]
>gi|1037022711|ref|XP_017139439.1| PREDICTED: LOW QUALITY PROTEIN: protein couch potato, partia
>gi|1037046729|ref|XP_017140938.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046808|ref|XP_017140943.1| PREDICTED: uncharacterized protein LOC108154948 isoform X9
>gi|1037046713|ref|XP_017140936.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046745|ref|XP_017140939.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046761|ref|XP_017140940.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046681|ref|XP_017140934.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046697|ref|XP_017140935.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046776|ref|XP_017140941.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046792|ref|XP_017140942.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
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>gi|1037046868|ref|XP_017140948.1| PREDICTED: uncharacterized protein LOC108154952 [Drosophila
>gi|1037030467|ref|XP_017139919.1| PREDICTED: protein tincar isoform X1 [Drosophila miranda]
>gi|1037030483|ref|XP_017139920.1| PREDICTED: protein tincar isoform X2 [Drosophila miranda]
>gi|1037030517|ref|XP_017139923.1| PREDICTED: chitobiosyldiphosphodolichol beta-mannosyltransfe
>gi|1037030743|ref|XP_017139938.1| PREDICTED: uncharacterized protein LOC108154222 [Drosophila
>gi|1037030653|ref|XP_017139932.1| PREDICTED: arfaptin-2 [Drosophila miranda]
>gi|1037030781|ref|XP_017139940.1| PREDICTED: DPH3 homolog [Drosophila miranda]
>gi|1037030686|ref|XP_017139934.1| PREDICTED: dnaJ homolog subfamily C member 17 [Drosophila mi
>gi|1037030501|ref|XP_017139921.1| PREDICTED: GRIP and coiled-coil domain-containing protein 1
>gi|1037030533|ref|XP_017139924.1| PREDICTED: NADH dehydrogenase [ubiquinone] complex I, assemb
>gi|1037030707|ref|XP_017139935.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1037030725|ref|XP_017139936.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1037030548|ref|XP_017139925.1| PREDICTED: 60S ribosomal protein L3 isoform X1 [Drosophila m
>gi|1037030566|ref|XP_017139926.1| PREDICTED: 60S ribosomal protein L3 isoform X2 [Drosophila m
>gi|1037030599|ref|XP_017139928.1| PREDICTED: zinc finger protein 544-like [Drosophila miranda]
>gi|1037030671|ref|XP_017139933.1| PREDICTED: J domain-containing protein CG6693 [Drosophila mi
>gi|1037030617|ref|XP_017139929.1| PREDICTED: uncharacterized protein LOC108154215 [Drosophila
>gi|1037030582|ref|XP_017139927.1| PREDICTED: uncharacterized protein LOC108154213 [Drosophila
>gi|1037022729|ref|XP_017139440.1| PREDICTED: uncharacterized protein LOC108153853 [Drosophila
>gi|1037030638|ref|XP_017139931.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037030763|ref|XP_017139939.1| PREDICTED: 40S ribosomal protein S25 [Drosophila miranda]
>gi|1037094813|ref|XP_017144288.1| PREDICTED: zinc transporter 7 isoform X1 [Drosophila miranda
>gi|1037094827|ref|XP_017144289.1| PREDICTED: zinc transporter 7 isoform X2 [Drosophila miranda
>gi|1037094841|ref|XP_017144291.1| PREDICTED: zinc transporter 7 isoform X2 [Drosophila miranda
>gi|1037096743|ref|XP_017144452.1| PREDICTED: translationally-controlled tumor protein homolog
>gi|1037089986|ref|XP_017143937.1| PREDICTED: succinate dehydrogenase cytochrome b560 subunit,
>gi|1037089972|ref|XP_017143935.1| PREDICTED: nocturnin isoform X5 [Drosophila miranda]
>gi|1037089918|ref|XP_017143931.1| PREDICTED: nocturnin isoform X1 [Drosophila miranda]
>gi|1037089958|ref|XP_017143934.1| PREDICTED: nocturnin isoform X4 [Drosophila miranda]
>gi|1037089931|ref|XP_017143932.1| PREDICTED: nocturnin isoform X2 [Drosophila miranda]
>gi|1037089944|ref|XP_017143933.1| PREDICTED: nocturnin isoform X3 [Drosophila miranda]
>gi|1037090000|ref|XP_017143938.1| PREDICTED: uncharacterized protein LOC108156777 [Drosophila
>gi|1037071025|ref|XP_017142586.1| PREDICTED: UDP-glucuronosyltransferase 2C1 isoform X2 [Droso
>gi|1037071009|ref|XP_017142585.1| PREDICTED: UDP-glucuronosyltransferase 1-9 isoform X1 [Droso
>gi|1037071059|ref|XP_017142589.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1037070957|ref|XP_017142582.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1037071043|ref|XP_017142587.1| PREDICTED: UDP-glucuronosyltransferase 2A3 [Drosophila miran
>gi|1037070993|ref|XP_017142584.1| PREDICTED: UDP-glucuronosyltransferase 2A3-like [Drosophila
>gi|1037070941|ref|XP_017142581.1| PREDICTED: venom carboxylesterase-6 [Drosophila miranda]
>gi|1037070975|ref|XP_017142583.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1037102241|ref|XP_017144957.1| PREDICTED: UDP-glucuronosyltransferase 2B15-like [Drosophila
>gi|1037106057|ref|XP_017145444.1| PREDICTED: LOW QUALITY PROTEIN: UDP-glucuronosyltransferase
>gi|1037105360|ref|XP_017145347.1| PREDICTED: UDP-glucuronosyltransferase 2B31 [Drosophila mira
>gi|1037103921|ref|XP_017145145.1| PREDICTED: probable aconitate hydratase, mitochondrial [Dros
>gi|1037105260|ref|XP_017145331.1| PREDICTED: UDP-glucuronosyltransferase 2B15 [Drosophila mira
>gi|1037033526|ref|XP_017140102.1| PREDICTED: uncharacterized protein LOC108154347 [Drosophila
>gi|1037085126|ref|XP_017143595.1| PREDICTED: nephrin [Drosophila miranda]
>gi|1037085179|ref|XP_017143599.1| PREDICTED: uncharacterized protein LOC108156576 [Drosophila
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>gi|1037085153|ref|XP_017143597.1| PREDICTED: uncharacterized protein LOC108156575 isoform X1
>gi|1037085167|ref|XP_017143598.1| PREDICTED: uncharacterized protein LOC108156575 isoform X2
>gi|1037085207|ref|XP_017143602.1| PREDICTED: uncharacterized protein LOC108156578 [Drosophila
>gi|1037085193|ref|XP_017143601.1| PREDICTED: uncharacterized protein LOC108156577 [Drosophila
>gi|1037085140|ref|XP_017143596.1| PREDICTED: proteasome subunit alpha type-6-like [Drosophila
>gi|1037074191|ref|XP_017142804.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037074175|ref|XP_017142803.1| PREDICTED: tetratricopeptide repeat protein 14 homolog [Dros
>gi|1037074206|ref|XP_017142805.1| PREDICTED: GTP:AMP phosphotransferase AK3, mitochondrial [D
>gi|1037074220|ref|XP_017142806.1| PREDICTED: GTP:AMP phosphotransferase AK3, mitochondrial [D
>gi|1037095949|ref|XP_017144378.1| PREDICTED: arrestin domain-containing protein 1 [Drosophila
>gi|1037095961|ref|XP_017144379.1| PREDICTED: mitochondrial thiamine pyrophosphate carrier [Dro
>gi|1037099638|ref|XP_017144720.1| PREDICTED: uncharacterized protein LOC108157253 [Drosophila
>gi|1037099626|ref|XP_017144719.1| PREDICTED: sorbitol dehydrogenase isoform X3 [Drosophila mi
>gi|1037099614|ref|XP_017144718.1| PREDICTED: sorbitol dehydrogenase isoform X2 [Drosophila mi
>gi|1037099602|ref|XP_017144717.1| PREDICTED: sorbitol dehydrogenase isoform X1 [Drosophila mi
>gi|1037105340|ref|XP_017145343.1| PREDICTED: uncharacterized protein LOC108157687 [Drosophila
>gi|1037062657|ref|XP_017142023.1| PREDICTED: alcohol dehydrogenase class-3 [Drosophila miranda
>gi|1037062673|ref|XP_017142024.1| PREDICTED: tetraspanin-33 [Drosophila miranda]
>gi|1037062687|ref|XP_017142025.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X1 [
>gi|1037062705|ref|XP_017142026.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X2 [
>gi|1037062721|ref|XP_017142028.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X3 [
>gi|1037062736|ref|XP_017142029.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X4 [
>gi|1037062596|ref|XP_017142019.1| PREDICTED: thiamine transporter 1 isoform X1 [Drosophila mi
>gi|1037062612|ref|XP_017142020.1| PREDICTED: thiamine transporter 1 isoform X2 [Drosophila mi
>gi|1037062627|ref|XP_017142021.1| PREDICTED: thiamine transporter 2 isoform X1 [Drosophila mi
>gi|1037062641|ref|XP_017142022.1| PREDICTED: thiamine transporter 2 isoform X2 [Drosophila mi
>gi|1037074797|ref|XP_017142845.1| PREDICTED: uncharacterized protein LOC108156072 [Drosophila
>gi|1037074781|ref|XP_017142844.1| PREDICTED: protein AF-9 [Drosophila miranda]
>gi|1037074839|ref|XP_017142849.1| PREDICTED: uncharacterized protein LOC108156074 [Drosophila
>gi|1037074823|ref|XP_017142846.1| PREDICTED: nuclear transcription factor Y subunit beta [Dros
>gi|1037103175|ref|XP_017145055.1| PREDICTED: tropomyosin-2 isoform X1 [Drosophila miranda]
>gi|1037103185|ref|XP_017145056.1| PREDICTED: tropomyosin-2 isoform X2 [Drosophila miranda]
>gi|1037055526|ref|XP_017141539.1| PREDICTED: golgin-84 isoform X11 [Drosophila miranda]
>gi|1037055509|ref|XP_017141538.1| PREDICTED: golgin-84 isoform X10 [Drosophila miranda]
>gi|1037055649|ref|XP_017141546.1| PREDICTED: tropomyosin-1, isoforms 9A/A/B isoform X18 [Droso
>gi|1037055665|ref|XP_017141547.1| PREDICTED: tropomyosin-1, isoforms 9A/A/B isoform X19 [Droso
>gi|1037055584|ref|XP_017141542.1| PREDICTED: tropomyosin-2 isoform X14 [Drosophila miranda]
>gi|1037055600|ref|XP_017141543.1| PREDICTED: tropomyosin-2 isoform X15 [Drosophila miranda]
>gi|1037055465|ref|XP_017141534.1| PREDICTED: golgin-84 isoform X7 [Drosophila miranda]
>gi|1037055451|ref|XP_017141533.1| PREDICTED: golgin-84 isoform X6 [Drosophila miranda]
>gi|1037055383|ref|XP_017141529.1| PREDICTED: formin-binding protein 4 isoform X2 [Drosophila m
>gi|1037055367|ref|XP_017141528.1| PREDICTED: formin-binding protein 4 isoform X1 [Drosophila m
>gi|1037055399|ref|XP_017141530.1| PREDICTED: formin-binding protein 4 isoform X3 [Drosophila m
>gi|1037055568|ref|XP_017141541.1| PREDICTED: tropomyosin-1, isoforms 33/34 isoform X13 [Drosop
>gi|1037055543|ref|XP_017141540.1| PREDICTED: tropomyosin-2 isoform X12 [Drosophila miranda]
>gi|1037055415|ref|XP_017141531.1| PREDICTED: actin cytoskeleton-regulatory complex protein pa
>gi|1037055435|ref|XP_017141532.1| PREDICTED: actin cytoskeleton-regulatory complex protein pa
>gi|1037055614|ref|XP_017141544.1| PREDICTED: tropomyosin-1, isoforms 33/34 isoform X16 [Drosop
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>gi|1037055492|ref|XP_017141537.1| PREDICTED: golgin-84 isoform X9 [Drosophila miranda]
>gi|1037055476|ref|XP_017141536.1| PREDICTED: golgin-84 isoform X8 [Drosophila miranda]
>gi|1037055633|ref|XP_017141545.1| PREDICTED: tropomyosin-1, isoforms 9A/A/B isoform X17 [Drosc
>gi|1037043237|ref|XP_017140706.1| PREDICTED: cytochrome c oxidase assembly protein COX16 homol
>gi|1037043147|ref|XP_017140700.1| PREDICTED: nuA4 complex subunit EAF3 homolog [Drosophila mir
>gi|1037043212|ref|XP_017140704.1| PREDICTED: ribosome biogenesis protein TSR3 homolog [Drosopl
>gi|1037043055|ref|XP_017140693.1| PREDICTED: centrosome-associated zinc finger protein CP190
>gi|1037043069|ref|XP_017140694.1| PREDICTED: centrosome-associated zinc finger protein CP190
>gi|1037043085|ref|XP_017140695.1| PREDICTED: centrosome-associated zinc finger protein CP190
>gi|1037043180|ref|XP_017140702.1| PREDICTED: protein zntD isoform X2 [Drosophila miranda]
>gi|1037043164|ref|XP_017140701.1| PREDICTED: protein zntD isoform X1 [Drosophila miranda]
>gi|1037043101|ref|XP_017140696.1| PREDICTED: uncharacterized protein LOC108154808 [Drosophila
>gi|1037043224|ref|XP_017140705.1| PREDICTED: ATP synthase subunit O, mitochondrial [Drosophila
>gi|1037043196|ref|XP_017140703.1| PREDICTED: protein SET [Drosophila miranda]
>gi|1037043132|ref|XP_017140698.1| PREDICTED: eukaryotic translation initiation factor 2 subuni
>gi|1037043117|ref|XP_017140697.1| PREDICTED: histone-lysine N-methyltransferase Su(var)3-9 iso
>gi|1037043252|ref|XP_017140707.1| PREDICTED: AF4/FMR2 family member 4 [Drosophila miranda]
>gi|1037043264|ref|XP_017140708.1| PREDICTED: uncharacterized protein LOC108154819 [Drosophila
>gi|1037081134|ref|XP_017143291.1| PREDICTED: putative ammonium transporter 2 [Drosophila mirar
>gi|1037081120|ref|XP_017143289.1| PREDICTED: heat shock 70 kDa protein cognate 4 [Drosophila r
>gi|1037052045|ref|XP_017141299.1| PREDICTED: serine/threonine-protein kinase ATM [Drosophila r
>gi|1037052138|ref|XP_017141304.1| PREDICTED: uncharacterized protein LOC108155155 [Drosophila
>gi|1037052154|ref|XP_017141306.1| PREDICTED: 28S ribosomal protein S10, mitochondrial isoform
>gi|1037052170|ref|XP_017141307.1| PREDICTED: 28S ribosomal protein S10, mitochondrial isoform
>gi|1037052074|ref|XP_017141300.1| PREDICTED: protein arginine N-methyltransferase 1 isoform X
>gi|1037052090|ref|XP_017141301.1| PREDICTED: protein arginine N-methyltransferase 1 isoform X
>gi|1037052106|ref|XP_017141302.1| PREDICTED: probable histone-binding protein Caf1 [Drosophila
>gi|1037052122|ref|XP_017141303.1| PREDICTED: uncharacterized protein LOC108155154 [Drosophila
>gi|1037089715|ref|XP_017143914.1| PREDICTED: rho GTPase-activating protein 92B [Drosophila mir
>gi|1037106167|ref|XP_017145459.1| PREDICTED: uncharacterized protein LOC108157781 [Drosophila
>gi|1037106159|ref|XP_017145458.1| PREDICTED: uncharacterized protein LOC108157779 [Drosophila
>gi|1037034204|ref|XP_017140143.1| PREDICTED: solute carrier family 22 member 3 [Drosophila mir
>gi|1037098529|ref|XP_017144623.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037098505|ref|XP_017144621.1| PREDICTED: solute carrier family 22 member 13 [Drosophila mi
>gi|1037098517|ref|XP_017144622.1| PREDICTED: solute carrier family 22 member 13 [Drosophila mi
>gi|1037098495|ref|XP_017144620.1| PREDICTED: solute carrier family 22 member 1 [Drosophila mir
>gi|1037071205|ref|XP_017142599.1| PREDICTED: heparan-sulfate 6-O-sulfotransferase 2 [Drosophil
>gi|1037071189|ref|XP_017142598.1| PREDICTED: FYVE and coiled-coil domain-containing protein 1
>gi|1037071171|ref|XP_017142597.1| PREDICTED: E3 ubiquitin-protein ligase BRE1 isoform X1 [Dros
>gi|1037071236|ref|XP_017142600.1| PREDICTED: uncharacterized protein LOC108155946 [Drosophila
>gi|1037071268|ref|XP_017142602.1| PREDICTED: uncharacterized protein LOC108155949 [Drosophila
>gi|1037071252|ref|XP_017142601.1| PREDICTED: uncharacterized protein LOC108155948 [Drosophila
>gi|1037033234|ref|XP_017140086.1| PREDICTED: uncharacterized protein LOC108154336 [Drosophila
>gi|1037060596|ref|XP_017141883.1| PREDICTED: uncharacterized protein LOC108155531 isoform X2
>gi|1037060582|ref|XP_017141882.1| PREDICTED: uncharacterized protein LOC108155531 isoform X1
>gi|1037060610|ref|XP_017141884.1| PREDICTED: uncharacterized protein LOC108155531 isoform X3
>gi|1037060626|ref|XP_017141886.1| PREDICTED: uncharacterized protein LOC108155531 isoform X4
>gi|1037060642|ref|XP_017141887.1| PREDICTED: uncharacterized protein LOC108155531 isoform X5
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>gi|1037060706|ref|XP_017141891.1| PREDICTED: N-acetylglucosaminyl-phosphatidylinositol de-N-ac
>gi|1037060658|ref|XP_017141888.1| PREDICTED: phagocyte signaling-impaired protein isoform X1
>gi|1037060672|ref|XP_017141889.1| PREDICTED: phagocyte signaling-impaired protein isoform X2
>gi|1037060736|ref|XP_017141894.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037060752|ref|XP_017141895.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037060720|ref|XP_017141892.1| PREDICTED: zinc finger protein 287 [Drosophila miranda]
>gi|1037060690|ref|XP_017141890.1| PREDICTED: zinc finger protein 62 homolog [Drosophila mirand
>gi|1037090709|ref|XP_017143995.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1037090695|ref|XP_017143994.1| PREDICTED: probable ribonuclease ZC3H12C [Drosophila miranda
>gi|1037103772|ref|XP_017145126.1| PREDICTED: uncharacterized protein LOC108157534 [Drosophila
>gi|1037103748|ref|XP_017145122.1| PREDICTED: protein I'm not dead yet 2-like [Drosophila miran
>gi|1037103756|ref|XP_017145123.1| PREDICTED: protein I'm not dead yet 2-like [Drosophila miran
>gi|1037103764|ref|XP_017145125.1| PREDICTED: uncharacterized protein LOC108157532 [Drosophila
>gi|1037095312|ref|XP_017144328.1| PREDICTED: S-formylglutathione hydrolase [Drosophila miranda
>gi|1037095326|ref|XP_017144329.1| PREDICTED: S-formylglutathione hydrolase [Drosophila miranda
>gi|1037095341|ref|XP_017144330.1| PREDICTED: S-formylglutathione hydrolase [Drosophila miranda
>gi|1037095298|ref|XP_017144327.1| PREDICTED: RING finger protein 113A [Drosophila miranda]
>gi|1037104656|ref|XP_017145244.1| PREDICTED: opsin Rh3 [Drosophila miranda]
>gi|1037097219|ref|XP_017144495.1| PREDICTED: heat shock 70 kDa protein cognate 4-like [Drosoph
>gi|1037105251|ref|XP_017145330.1| PREDICTED: muscle-specific protein 20 [Drosophila miranda]
>gi|1037103359|ref|XP_017145075.1| PREDICTED: ureidoglycolate dehydrogenase (NAD(+)) [Drosophil
>gi|1037103339|ref|XP_017145073.1| PREDICTED: neuroligin 4-like [Drosophila miranda]
>gi|1037103349|ref|XP_017145074.1| PREDICTED: neuroligin 4-like [Drosophila miranda]
>gi|1037066448|ref|XP_017142272.1| PREDICTED: putative transcription factor capicua isoform X1
>gi|1037066496|ref|XP_017142275.1| PREDICTED: putative transcription factor capicua isoform X4
>gi|1037066464|ref|XP_017142273.1| PREDICTED: putative transcription factor capicua isoform X2
>gi|1037066511|ref|XP_017142276.1| PREDICTED: putative transcription factor capicua isoform X5
>gi|1037066480|ref|XP_017142274.1| PREDICTED: putative transcription factor capicua isoform X3
>gi|1037066529|ref|XP_017142277.1| PREDICTED: putative transcription factor capicua isoform X6
>gi|1037066561|ref|XP_017142279.1| PREDICTED: putative transcription factor capicua isoform X8
>gi|1037066546|ref|XP_017142278.1| PREDICTED: putative transcription factor capicua isoform X7
>gi|1037037294|ref|XP_017140318.1| PREDICTED: uncharacterized protein LOC108154526 [Drosophila
>gi|1037099154|ref|XP_017144674.1| PREDICTED: uncharacterized protein LOC108157228 [Drosophila
>gi|1037099087|ref|XP_017144672.1| PREDICTED: titin homolog [Drosophila miranda]
>gi|1037099105|ref|XP_017144673.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 [D
>gi|1037049314|ref|XP_017141109.1| PREDICTED: furin-like protease 1 isoform X2 [Drosophila mira
>gi|1037049284|ref|XP_017141107.1| PREDICTED: furin-like protease 1 isoform X1 [Drosophila mira
>gi|1037049300|ref|XP_017141108.1| PREDICTED: furin-like protease 1 isoform X1 [Drosophila mira
>gi|1037049328|ref|XP_017141110.1| PREDICTED: NADH-quinone oxidoreductase subunit D [Drosophila
>gi|1037049347|ref|XP_017141112.1| PREDICTED: carboxypeptidase N subunit 2 [Drosophila miranda]
>gi|1037049361|ref|XP_017141113.1| PREDICTED: collagen alpha-1(V) chain [Drosophila miranda]
>gi|1037049374|ref|XP_017141114.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1037094349|ref|XP_017144253.1| PREDICTED: vacuolar protein sorting-associated protein 33B
>gi|1037094363|ref|XP_017144254.1| PREDICTED: vacuolar protein sorting-associated protein 33B
>gi|1037094376|ref|XP_017144255.1| PREDICTED: uncharacterized protein LOC108156975 [Drosophila
>gi|1037031400|ref|XP_017139977.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1037106628|ref|XP_017145523.1| PREDICTED: regulator of chromosome condensation-like [Drosop
>gi|1037057418|ref|XP_017141659.1| PREDICTED: transcriptional regulator ATRX homolog [Drosophil
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>gi|1037057402|ref|XP_017141658.1| PREDICTED: nuclear factor related to kappa-B-binding protei
>gi|1037094855|ref|XP_017144292.1| PREDICTED: zinc finger protein 423 [Drosophila miranda]
>gi|1037091031|ref|XP_017144019.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1037090949|ref|XP_017144012.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037090963|ref|XP_017144013.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037090976|ref|XP_017144014.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037090990|ref|XP_017144015.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037091004|ref|XP_017144016.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037091018|ref|XP_017144017.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037091047|ref|XP_017144020.1| PREDICTED: venom allergen 3 [Drosophila miranda]
>gi|1037053318|ref|XP_017141385.1| PREDICTED: UPF0160 protein [Drosophila miranda]
>gi|1037053356|ref|XP_017141386.1| PREDICTED: proteasome subunit beta type-3 [Drosophila miranc
>gi|1037053304|ref|XP_017141384.1| PREDICTED: E3 ubiquitin-protein ligase RNF126-B [Drosophila
>gi|1037053276|ref|XP_017141382.1| PREDICTED: uncharacterized protein LOC108155209 [Drosophila
>gi|1037053250|ref|XP_017141380.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X2 [Dro
>gi|1037053264|ref|XP_017141381.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X3 [Dro
>gi|1037053202|ref|XP_017141377.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1037053218|ref|XP_017141378.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1037053234|ref|XP_017141379.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1037053288|ref|XP_017141383.1| PREDICTED: embryonic stem cell-specific 5-hydroxymethylcytos
>gi|1037053176|ref|XP_017141374.1| PREDICTED: aryl hydrocarbon receptor nuclear translocator ho
>gi|1037053196|ref|XP_017141376.1| PREDICTED: aryl hydrocarbon receptor nuclear translocator ho
>gi|1037076200|ref|XP_017142943.1| PREDICTED: protein neuralized isoform X2 [Drosophila mirand
>gi|1037076184|ref|XP_017142942.1| PREDICTED: protein neuralized isoform X1 [Drosophila mirand
>gi|1037065545|ref|XP_017142211.1| PREDICTED: parafibromin [Drosophila miranda]
>gi|1037065577|ref|XP_017142213.1| PREDICTED: bifunctional methylenetetrahydrofolate dehydroger
>gi|1037065561|ref|XP_017142212.1| PREDICTED: bifunctional methylenetetrahydrofolate dehydroger
>gi|1037065593|ref|XP_017142214.1| PREDICTED: bifunctional methylenetetrahydrofolate dehydroger
>gi|1037065527|ref|XP_017142210.1| PREDICTED: nuclear factor NF-kappa-B p110 subunit [Drosophil
>gi|1037065609|ref|XP_017142215.1| PREDICTED: uncharacterized protein LOC108155722 [Drosophila
>gi|1037077240|ref|XP_017143010.1| PREDICTED: putative protein TPRXL [Drosophila miranda]
>gi|1037072769|ref|XP_017142705.1| PREDICTED: uncharacterized protein LOC108156011 [Drosophila
>gi|1037072753|ref|XP_017142703.1| PREDICTED: uncharacterized protein LOC108156010 [Drosophila
>gi|1037072737|ref|XP_017142702.1| PREDICTED: T-complex protein 1 subunit eta [Drosophila mira
>gi|1037072722|ref|XP_017142701.1| PREDICTED: uncharacterized protein LOC108156008 [Drosophila
>gi|1037070676|ref|XP_017142562.1| PREDICTED: cytospin-A isoform X1 [Drosophila miranda]
>gi|1037070690|ref|XP_017142563.1| PREDICTED: cytospin-A isoform X2 [Drosophila miranda]
>gi|1037070765|ref|XP_017142569.1| PREDICTED: uncharacterized protein LOC108155927 [Drosophila
>gi|1037070705|ref|XP_017142564.1| PREDICTED: intracellular protein transport protein USO1 isor
>gi|1037070733|ref|XP_017142566.1| PREDICTED: uncharacterized protein LOC108155925 [Drosophila
>gi|1037070719|ref|XP_017142565.1| PREDICTED: uncharacterized protein LOC108155924 [Drosophila
>gi|1037022747|ref|XP_017139441.1| PREDICTED: uncharacterized protein LOC108153854 [Drosophila
>gi|1037070749|ref|XP_017142568.1| PREDICTED: uncharacterized protein LOC108155926 [Drosophila
>gi|1037105227|ref|XP_017145327.1| PREDICTED: chymotrypsinogen A isoform X1 [Drosophila miranda
>gi|1037105235|ref|XP_017145328.1| PREDICTED: chymotrypsinogen A isoform X2 [Drosophila miranda
>gi|1037105219|ref|XP_017145326.1| PREDICTED: uncharacterized protein LOC108157673 [Drosophila
>gi|1037090668|ref|XP_017143991.1| PREDICTED: uncharacterized protein LOC108156809 [Drosophila
>gi|1037090682|ref|XP_017143992.1| PREDICTED: protein NASP homolog [Drosophila miranda]
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>gi|1037048525|ref|XP_017141055.1| PREDICTED: PP2C-like domain-containing protein CG9801 [Dros
>gi|1037048509|ref|XP_017141054.1| PREDICTED: UPF0505 protein CG8202 isoform X2 [Drosophila mi
>gi|1037048496|ref|XP_017141053.1| PREDICTED: UPF0505 protein CG8202 isoform X1 [Drosophila mi
>gi|1037048573|ref|XP_017141058.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037048589|ref|XP_017141059.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037048667|ref|XP_017141065.1| PREDICTED: transcription factor Ouib-like [Drosophila miran
>gi|1037048651|ref|XP_017141063.1| PREDICTED: transcription factor Ouib isoform X4 [Drosophila
>gi|1037048619|ref|XP_017141061.1| PREDICTED: transcription factor Ouib isoform X2 [Drosophila
>gi|1037048635|ref|XP_017141062.1| PREDICTED: transcription factor Ouib isoform X3 [Drosophila
>gi|1037048605|ref|XP_017141060.1| PREDICTED: transcription factor Ouib isoform X1 [Drosophila
>gi|1037048557|ref|XP_017141057.1| PREDICTED: uncharacterized protein LOC108155016 [Drosophila
>gi|1037048683|ref|XP_017141066.1| PREDICTED: transmembrane protein 216 [Drosophila miranda]
>gi|1037048699|ref|XP_017141067.1| PREDICTED: uncharacterized protein LOC108155023 [Drosophila
>gi|1037048715|ref|XP_017141068.1| PREDICTED: cyclin-dependent kinases regulatory subunit [Dros
>gi|1037048541|ref|XP_017141056.1| PREDICTED: sterol O-acyltransferase 1 [Drosophila miranda]
>gi|1037088751|ref|XP_017143837.1| PREDICTED: protein hunchback [Drosophila miranda]
>gi|1037088761|ref|XP_017143838.1| PREDICTED: protein hunchback [Drosophila miranda]
>gi|1037044068|ref|XP_017140757.1| PREDICTED: nuclear RNA export factor 2-like isoform X2 [Dros
>gi|1037044053|ref|XP_017140756.1| PREDICTED: nuclear RNA export factor 2-like isoform X1 [Dros
>gi|1037044084|ref|XP_017140758.1| PREDICTED: nuclear RNA export factor 2-like isoform X3 [Dros
>gi|1037044207|ref|XP_017140766.1| PREDICTED: protein unc-50 homolog [Drosophila miranda]
>gi|1037044173|ref|XP_017140764.1| PREDICTED: 39S ribosomal protein L19, mitochondrial [Drosopl
>gi|1037044191|ref|XP_017140765.1| PREDICTED: 39S ribosomal protein L19, mitochondrial [Drosopl
>gi|1037044109|ref|XP_017140760.1| PREDICTED: zinc finger protein 2 homolog isoform X2 [Drosopl
>gi|1037044125|ref|XP_017140761.1| PREDICTED: zinc finger protein 2 homolog isoform X2 [Drosopl
>gi|1037044098|ref|XP_017140759.1| PREDICTED: oocyte zinc finger protein XlCOF6-like isoform X1
>gi|1037044223|ref|XP_017140767.1| PREDICTED: putative riboflavin kinase [Drosophila miranda]
>gi|1037044141|ref|XP_017140762.1| PREDICTED: amidophosphoribosyltransferase [Drosophila mirand
>gi|1037044157|ref|XP_017140763.1| PREDICTED: transcription initiation factor TFIID subunit 7
>gi|1037044039|ref|XP_017140754.1| PREDICTED: ER membrane protein complex subunit 1 [Drosophila
>gi|1037068662|ref|XP_017142422.1| PREDICTED: 28S ribosomal protein S9, mitochondrial [Drosophi
>gi|1037068406|ref|XP_017142405.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068376|ref|XP_017142402.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068391|ref|XP_017142403.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068437|ref|XP_017142407.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068422|ref|XP_017142406.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068451|ref|XP_017142408.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068646|ref|XP_017142421.1| PREDICTED: probable arginine--tRNA ligase, mitochondrial [D:
>gi|1037068467|ref|XP_017142409.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X:
>gi|1037068483|ref|XP_017142410.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X:
>gi|1037068581|ref|XP_017142416.1| PREDICTED: uncharacterized protein LOC108155825 isoform X7
>gi|1037068515|ref|XP_017142412.1| PREDICTED: formin-J isoform X3 [Drosophila miranda]
>gi|1037068598|ref|XP_017142417.1| PREDICTED: uncharacterized protein LOC108155825 isoform X8
>gi|1037068549|ref|XP_017142414.1| PREDICTED: chromatin modification-related protein EAF1 isofo
>gi|1037068498|ref|XP_017142411.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X2
>gi|1037068565|ref|XP_017142415.1| PREDICTED: chromatin modification-related protein EAF1 isofo
>gi|1037068531|ref|XP_017142413.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X4
>gi|1037068614|ref|XP_017142418.1| PREDICTED: uncharacterized protein LOC108155825 isoform X9
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>gi|1037068630|ref|XP_017142419.1| PREDICTED: uncharacterized protein LOC108155825 isoform X10
>gi|1037101801|ref|XP_017144910.1| PREDICTED: diacylglycerol kinase eta isoform X6 [Drosophila
>gi|1037101781|ref|XP_017144908.1| PREDICTED: diacylglycerol kinase eta isoform X4 [Drosophila
>gi|1037101761|ref|XP_017144906.1| PREDICTED: diacylglycerol kinase eta isoform X2 [Drosophila
>gi|1037101811|ref|XP_017144911.1| PREDICTED: diacylglycerol kinase eta isoform X7 [Drosophila
>gi|1037101771|ref|XP_017144907.1| PREDICTED: diacylglycerol kinase eta isoform X3 [Drosophila
>gi|1037101751|ref|XP_017144905.1| PREDICTED: diacylglycerol kinase eta isoform X1 [Drosophila
>gi|1037101791|ref|XP_017144909.1| PREDICTED: diacylglycerol kinase eta isoform X5 [Drosophila
>gi|1037100400|ref|XP_017144779.1| PREDICTED: pyridoxine/pyridoxamine 5'-phosphate oxidase [Dro
>gi|1037100384|ref|XP_017144778.1| PREDICTED: pyridoxine-5'-phosphate oxidase [Drosophila miran
>gi|1037090225|ref|XP_017143956.1| PREDICTED: arrestin domain-containing protein 1 [Drosophila
>gi|1037104913|ref|XP_017145280.1| PREDICTED: uncharacterized protein LOC108157644 [Drosophila
>gi|1037104905|ref|XP_017145279.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037100488|ref|XP_017144787.1| PREDICTED: arrestin domain-containing protein 5 [Drosophila
>gi|1037100476|ref|XP_017144786.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037100460|ref|XP_017144785.1| PREDICTED: arrestin domain-containing protein 2 [Drosophila
>gi|1037033073|ref|XP_017140076.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037033090|ref|XP_017140077.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037106440|ref|XP_017145497.1| PREDICTED: arrestin domain-containing protein 15 [Drosophila
>gi|1037106175|ref|XP_017145460.1| PREDICTED: arrestin domain-containing protein 2 [Drosophila
>gi|1037106015|ref|XP_017145439.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037035323|ref|XP_017140207.1| PREDICTED: uncharacterized protein LOC108154447 [Drosophila
>gi|1037105720|ref|XP_017145397.1| PREDICTED: serine protease easter isoform X1 [Drosophila mi
>gi|1037105726|ref|XP_017145398.1| PREDICTED: serine protease easter isoform X2 [Drosophila mi
>gi|1037034038|ref|XP_017140133.1| PREDICTED: uncharacterized protein LOC108154380 [Drosophila
>gi|1037034018|ref|XP_017140132.1| PREDICTED: uncharacterized protein LOC108154379 [Drosophila
>gi|1037037416|ref|XP_017140326.1| PREDICTED: uncharacterized protein LOC108154534 [Drosophila
>gi|1037033196|ref|XP_017140084.1| PREDICTED: solute carrier family 25 member 40 [Drosophila mi
>gi|1037099720|ref|XP_017144728.1| PREDICTED: uncharacterized protein LOC108157261 [Drosophila
>gi|1037099734|ref|XP_017144729.1| PREDICTED: uncharacterized protein LOC108157261 [Drosophila
>gi|1037103536|ref|XP_017145095.1| PREDICTED: cytochrome c oxidase subunit 4 isoform 1, mitoch
>gi|1037022765|ref|XP_017139443.1| PREDICTED: neuroligin-4, Y-linked [Drosophila miranda]
>gi|1037022781|ref|XP_017139444.1| PREDICTED: uncharacterized protein LOC108153856 [Drosophila
>gi|1037064400|ref|XP_017142143.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037064414|ref|XP_017142144.1| PREDICTED: esterase B1 isoform X1 [Drosophila miranda]
>gi|1037064430|ref|XP_017142145.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1037064446|ref|XP_017142146.1| PREDICTED: esterase B1-like isoform X1 [Drosophila miranda]
>gi|1037064460|ref|XP_017142148.1| PREDICTED: esterase B1-like isoform X2 [Drosophila miranda]
>gi|1037022805|ref|XP_017139445.1| PREDICTED: uncharacterized protein LOC108153857 [Drosophila
>gi|1037064476|ref|XP_017142149.1| PREDICTED: reticulon-3 [Drosophila miranda]
>gi|1037097936|ref|XP_017144562.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037097947|ref|XP_017144565.1| PREDICTED: ubiquitin-conjugating enzyme E2-18 kDa [Drosophil
>gi|1037088826|ref|XP_017143844.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037088840|ref|XP_017143845.1| PREDICTED: esterase B1 isoform X1 [Drosophila miranda]
>gi|1037088880|ref|XP_017143848.1| PREDICTED: growth hormone-inducible transmembrane protein [
>gi|1037088854|ref|XP_017143846.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1037088867|ref|XP_017143847.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037036647|ref|XP_017140281.1| PREDICTED: uncharacterized protein LOC108154492 [Drosophila
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>gi|1037035357|ref|XP_017140209.1| PREDICTED: neural cell adhesion molecule 1-B isoform X1 [Dro
>gi|1037035375|ref|XP_017140210.1| PREDICTED: neural cell adhesion molecule 1-B isoform X2 [Dro
>gi|1037106477|ref|XP_017145503.1| PREDICTED: uncharacterized protein LOC108157815 [Drosophila
>gi|1037022823|ref|XP_017139446.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1037022840|ref|XP_017139447.1| PREDICTED: glutamate receptor U1 [Drosophila miranda]
>gi|1037035824|ref|XP_017140231.1| PREDICTED: uncharacterized protein LOC108154471 isoform X1
>gi|1037035840|ref|XP_017140232.1| PREDICTED: uncharacterized protein LOC108154471 isoform X2
>gi|1037106378|ref|XP_017145488.1| PREDICTED: uncharacterized protein LOC108157805 [Drosophila
>gi|1037105594|ref|XP_017145380.1| PREDICTED: uncharacterized protein LOC108157722 [Drosophila
>gi|1037105586|ref|XP_017145378.1| PREDICTED: classical arabinogalactan protein 9 [Drosophila
>gi|1037022884|ref|XP_017139448.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037106815|ref|XP_017145550.1| PREDICTED: endoplasmic reticulum resident protein 44 [Drosop
>gi|1037095102|ref|XP_017144312.1| PREDICTED: LOW QUALITY PROTEIN: zinc finger protein rotund
>gi|1037095186|ref|XP_017144319.1| PREDICTED: GTPase-activating protein RacGAP84C-like [Drosop
>gi|1037022481|ref|XP_017139423.1| PREDICTED: LOW QUALITY PROTEIN: GTPase-activating protein Ra
>gi|1037095172|ref|XP_017144318.1| PREDICTED: transcription activator MSS11-like [Drosophila mi
>gi|1037095158|ref|XP_017144317.1| PREDICTED: very low-density lipoprotein receptor [Drosophila
>gi|1037095116|ref|XP_017144313.1| PREDICTED: synaptotagmin-4 [Drosophila miranda]
>gi|1037095130|ref|XP_017144314.1| PREDICTED: synaptotagmin-4 [Drosophila miranda]
>gi|1037095144|ref|XP_017144315.1| PREDICTED: synaptotagmin-4 [Drosophila miranda]
>gi|1037066760|ref|XP_017142289.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1037066776|ref|XP_017142290.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1037066901|ref|XP_017142299.1| PREDICTED: general odorant-binding protein 84a isoform X2 [
>gi|1037066886|ref|XP_017142298.1| PREDICTED: general odorant-binding protein 84a isoform X1 [
>gi|1037066917|ref|XP_017142300.1| PREDICTED: general odorant-binding protein 84a isoform X3 [
>gi|1037066870|ref|XP_017142297.1| PREDICTED: DTW domain-containing protein 2 [Drosophila mira
>gi|1037066822|ref|XP_017142294.1| PREDICTED: serine/threonine-protein kinase 16 [Drosophila mi
>gi|1037066854|ref|XP_017142296.1| PREDICTED: G patch domain-containing protein 11 isoform X2
>gi|1037066838|ref|XP_017142295.1| PREDICTED: G patch domain-containing protein 11 isoform X1
>gi|1037066744|ref|XP_017142288.1| PREDICTED: nucleolar complex protein 3 homolog [Drosophila
>gi|1037066808|ref|XP_017142292.1| PREDICTED: uncharacterized protein LOC108155760 isoform X2
>gi|1037066792|ref|XP_017142291.1| PREDICTED: uncharacterized protein LOC108155760 isoform X1
>gi|1037066700|ref|XP_017142285.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037066714|ref|XP_017142286.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037066729|ref|XP_017142287.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037105553|ref|XP_017145374.1| PREDICTED: facilitated trehalose transporter Tret1-like [Dro
>gi|1037106033|ref|XP_017145441.1| PREDICTED: uncharacterized protein LOC108157769 isoform X1
>gi|1037106041|ref|XP_017145442.1| PREDICTED: uncharacterized protein LOC108157769 isoform X2
>gi|1037106049|ref|XP_017145443.1| PREDICTED: uncharacterized protein LOC108157769 isoform X3
>gi|1037105999|ref|XP_017145436.1| PREDICTED: uncharacterized protein LOC108157765 [Drosophila
>gi|1037084425|ref|XP_017143541.1| PREDICTED: phosphatidylinositol-binding clathrin assembly pr
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>gi|1037084374|ref|XP_017143537.1| PREDICTED: phosphatidylinositol-binding clathrin assembly pr
>gi|1037084387|ref|XP_017143538.1| PREDICTED: phosphatidylinositol-binding clathrin assembly pr
>gi|1037084360|ref|XP_017143536.1| PREDICTED: phosphatidylinositol-binding clathrin assembly pr
>gi|1037084480|ref|XP_017143546.1| PREDICTED: phosphatidylinositol-binding clathrin assembly pr
>gi|1037084439|ref|XP_017143543.1| PREDICTED: phosphatidylinositol-binding clathrin assembly pr
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>gi|1037084401|ref|XP_017143539.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p:
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>gi|1037084453|ref|XP_017143544.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p:
>gi|1037096367|ref|XP_017144418.1| PREDICTED: protein SHQ1 homolog [Drosophila miranda]
>gi|1037072709|ref|XP_017142700.1| PREDICTED: putative epidermal cell surface receptor isoform
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>gi|1037072680|ref|XP_017142698.1| PREDICTED: putative epidermal cell surface receptor isoform
>gi|1037066134|ref|XP_017142250.1| PREDICTED: uncharacterized protein LOC108155743 [Drosophila
>gi|1037066150|ref|XP_017142251.1| PREDICTED: uncharacterized protein LOC108155743 [Drosophila
>gi|1037066165|ref|XP_017142252.1| PREDICTED: uncharacterized protein LOC108155743 [Drosophila
>gi|1037066106|ref|XP_017142248.1| PREDICTED: centromere protein J [Drosophila miranda]
>gi|1037066120|ref|XP_017142249.1| PREDICTED: GPN-loop GTPase 3 [Drosophila miranda]
>gi|1037066075|ref|XP_017142246.1| PREDICTED: uncharacterized protein LOC108155740 isoform X1
>gi|1037066089|ref|XP_017142247.1| PREDICTED: glutathione S-transferase 1-1 isoform X2 [Drosopl
>gi|1037104477|ref|XP_017145219.1| PREDICTED: uncharacterized protein LOC108157601 isoform X2
>gi|1037104469|ref|XP_017145218.1| PREDICTED: protein twist isoform X1 [Drosophila miranda]
>gi|1037099187|ref|XP_017144676.1| PREDICTED: uncharacterized protein LOC108157230 [Drosophila
>gi|1037099166|ref|XP_017144675.1| PREDICTED: uncharacterized protein LOC108157229 [Drosophila
>gi|1037050467|ref|XP_017141189.1| PREDICTED: ELMO domain-containing protein 2 [Drosophila mira
>gi|1037050452|ref|XP_017141188.1| PREDICTED: secernin-3 [Drosophila miranda]
>gi|1037050273|ref|XP_017141176.1| PREDICTED: protein AF-17 isoform X1 [Drosophila miranda]
>gi|1037050289|ref|XP_017141177.1| PREDICTED: protein AF-17 isoform X2 [Drosophila miranda]
>gi|1037050309|ref|XP_017141178.1| PREDICTED: protein AF-17 isoform X3 [Drosophila miranda]
>gi|1037050323|ref|XP_017141179.1| PREDICTED: protein AF-10 isoform X4 [Drosophila miranda]
>gi|1037050339|ref|XP_017141180.1| PREDICTED: protein AF-10 isoform X5 [Drosophila miranda]
>gi|1037050356|ref|XP_017141182.1| PREDICTED: protein AF-17 isoform X6 [Drosophila miranda]
>gi|1037050370|ref|XP_017141183.1| PREDICTED: protein AF-10 isoform X7 [Drosophila miranda]
>gi|1037050420|ref|XP_017141186.1| PREDICTED: muscle LIM protein Mlp84B [Drosophila miranda]
>gi|1037050436|ref|XP_017141187.1| PREDICTED: muscle LIM protein Mlp84B [Drosophila miranda]
>gi|1037050388|ref|XP_017141184.1| PREDICTED: mucin-5AC isoform X8 [Drosophila miranda]
>gi|1037050404|ref|XP_017141185.1| PREDICTED: mucin-19 isoform X9 [Drosophila miranda]
>gi|1037050481|ref|XP_017141191.1| PREDICTED: enhancer of yellow 2b transcription factor [Droso
>gi|1037052639|ref|XP_017141339.1| PREDICTED: tubulin alpha-1 chain [Drosophila miranda]
>gi|1037052705|ref|XP_017141344.1| PREDICTED: rRNA methyltransferase 2, mitochondrial [Drosophi
>gi|1037052625|ref|XP_017141338.1| PREDICTED: venom serine carboxypeptidase [Drosophila miranda
>gi|1037052721|ref|XP_017141345.1| PREDICTED: transmembrane protein 256 homolog [Drosophila mir
>gi|1037052607|ref|XP_017141337.1| PREDICTED: LOW QUALITY PROTEIN: probable multidrug resistanc
>gi|1037052672|ref|XP_017141342.1| PREDICTED: zinc finger homeobox protein 3 [Drosophila miranc
>gi|1037052656|ref|XP_017141341.1| PREDICTED: short-chain specific acyl-CoA dehydrogenase, mitc
>gi|1037052687|ref|XP_017141343.1| PREDICTED: zinc finger protein 420-like [Drosophila miranda]
>gi|1037070024|ref|XP_017142518.1| PREDICTED: mitogen-activated protein kinase kinase kinase 1!
>gi|1037070068|ref|XP_017142522.1| PREDICTED: opsin Rh1 [Drosophila miranda]
>gi|1037070038|ref|XP_017142520.1| PREDICTED: serine/threonine protein phosphatase 2A regulato:
>gi|1037070054|ref|XP_017142521.1| PREDICTED: serine/threonine protein phosphatase 2A regulato:
>gi|1037042973|ref|XP_017140688.1| PREDICTED: ATP-dependent Clp protease ATP-binding subunit c
>gi|1037043007|ref|XP_017140690.1| PREDICTED: surfeit locus protein 6 homolog [Drosophila mirar
>gi|1037043023|ref|XP_017140691.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB7 [Dro:
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>gi|1037042955|ref|XP_017140686.1| PREDICTED: small G protein signaling modulator 3 homolog [Di
>gi|1037042874|ref|XP_017140681.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042892|ref|XP_017140682.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042908|ref|XP_017140683.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042922|ref|XP_017140684.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042938|ref|XP_017140685.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037043039|ref|XP_017140692.1| PREDICTED: diamine acetyltransferase 2 [Drosophila miranda]
>gi|1037042827|ref|XP_017140678.1| PREDICTED: box A-binding factor isoform X2 [Drosophila miran
>gi|1037042813|ref|XP_017140677.1| PREDICTED: box A-binding factor isoform X1 [Drosophila miran
>gi|1037042843|ref|XP_017140679.1| PREDICTED: box A-binding factor isoform X3 [Drosophila miran
>gi|1037042859|ref|XP_017140680.1| PREDICTED: box A-binding factor isoform X4 [Drosophila miran
>gi|1037042991|ref|XP_017140689.1| PREDICTED: box A-binding factor, partial [Drosophila miranda
>gi|1037076802|ref|XP_017142981.1| PREDICTED: GATA-binding factor A isoform X1 [Drosophila mira
>gi|1037076818|ref|XP_017142982.1| PREDICTED: GATA-binding factor A isoform X2 [Drosophila mira
>gi|1037076834|ref|XP_017142983.1| PREDICTED: GATA-binding factor A isoform X3 [Drosophila mira
>gi|1037076868|ref|XP_017142986.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037076883|ref|XP_017142987.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037076850|ref|XP_017142984.1| PREDICTED: inhibitor of nuclear factor kappa-B kinase subuni
>gi|1037076899|ref|XP_017142988.1| PREDICTED: methyltransferase-like protein 23 [Drosophila mi
>gi|1037060005|ref|XP_017141843.1| PREDICTED: LOW QUALITY PROTEIN: cytoskeleton-associated prot
>gi|1037059987|ref|XP_017141842.1| PREDICTED: LOW QUALITY PROTEIN: NACHT and WD repeat domain-c
>gi|1037062249|ref|XP_017141995.1| PREDICTED: unconventional myosin-XVIIIa isoform X1 [Drosoph
>gi|1037062279|ref|XP_017141997.1| PREDICTED: unconventional myosin-XVIIIa isoform X3 [Drosoph
>gi|1037062263|ref|XP_017141996.1| PREDICTED: unconventional myosin-XVIIIa isoform X2 [Drosoph
>gi|1037062295|ref|XP_017141998.1| PREDICTED: unconventional myosin-XVIIIa isoform X4 [Drosoph
>gi|1037062311|ref|XP_017141999.1| PREDICTED: unconventional myosin-XVIIIa isoform X5 [Drosoph
>gi|1037062327|ref|XP_017142000.1| PREDICTED: uncharacterized protein LOC108155593 [Drosophila
>gi|1037062341|ref|XP_017142001.1| PREDICTED: lipase member H [Drosophila miranda]
>gi|1037091445|ref|XP_017144050.1| PREDICTED: RAC serine/threonine-protein kinase [Drosophila
>gi|1037097366|ref|XP_017144508.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
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>gi|1037097390|ref|XP_017144510.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1037097402|ref|XP_017144511.1| PREDICTED: serine proteinase stubble isoform X2 [Drosophila
>gi|1037097413|ref|XP_017144513.1| PREDICTED: uncharacterized protein LOC108157128 [Drosophila
>gi|1037094063|ref|XP_017144232.1| PREDICTED: growth hormone-regulated TBC protein 1-A [Drosop
>gi|1037094077|ref|XP_017144233.1| PREDICTED: growth hormone-regulated TBC protein 1-A [Drosop
>gi|1037094121|ref|XP_017144236.1| PREDICTED: uncharacterized protein LOC108156965 isoform X3
>gi|1037094107|ref|XP_017144235.1| PREDICTED: circumsporozoite protein isoform X2 [Drosophila
>gi|1037094091|ref|XP_017144234.1| PREDICTED: circumsporozoite protein isoform X1 [Drosophila
>gi|1037097460|ref|XP_017144517.1| PREDICTED: organic cation transporter protein isoform X1 [Di
>gi|1037097471|ref|XP_017144518.1| PREDICTED: organic cation transporter protein isoform X2 [Di
>gi|1037097482|ref|XP_017144519.1| PREDICTED: organic cation transporter protein isoform X2 [Di
>gi|1037097449|ref|XP_017144516.1| PREDICTED: organic cation transporter protein isoform X3 [Di
>gi|1037097425|ref|XP_017144514.1| PREDICTED: organic cation transporter protein isoform X1 [Di
>gi|1037097437|ref|XP_017144515.1| PREDICTED: organic cation transporter protein isoform X2 [Di
>gi|1037100870|ref|XP_017144827.1| PREDICTED: caprin homolog [Drosophila miranda]
>gi|1037100880|ref|XP_017144828.1| PREDICTED: histone deacetylase complex subunit SAP18 [Drosop
>gi|1037091678|ref|XP_017144066.1| PREDICTED: protein sarah [Drosophila miranda]
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>gi|1037091664|ref|XP_017144065.1| PREDICTED: organic cation transporter protein [Drosophila m
>gi|1037076111|ref|XP_017142936.1| PREDICTED: uncharacterized protein LOC108156135 [Drosophila
>gi|1037076127|ref|XP_017142937.1| PREDICTED: single-stranded DNA-binding protein, mitochondria
>gi|1037076171|ref|XP_017142941.1| PREDICTED: GPI ethanolamine phosphate transferase 1 [Drosoph
>gi|1037076142|ref|XP_017142939.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037076156|ref|XP_017142940.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037022900|ref|XP_017139449.1| PREDICTED: GPI ethanolamine phosphate transferase 1 [Drosoph
>gi|1037080982|ref|XP_017143279.1| PREDICTED: TBC1 domain family member 10A isoform X1 [Drosoph
>gi|1037080995|ref|XP_017143280.1| PREDICTED: TBC1 domain family member 10A isoform X2 [Drosoph
>gi|1037080968|ref|XP_017143277.1| PREDICTED: zinc finger protein 850 [Drosophila miranda]
>gi|1037078498|ref|XP_017143100.1| PREDICTED: microtubule-associated protein Jupiter isoform X6
>gi|1037078466|ref|XP_017143098.1| PREDICTED: microtubule-associated protein Jupiter isoform X4
>gi|1037078333|ref|XP_017143096.1| PREDICTED: microtubule-associated protein Jupiter isoform X2
>gi|1037078484|ref|XP_017143099.1| PREDICTED: microtubule-associated protein Jupiter isoform X5
>gi|1037078452|ref|XP_017143097.1| PREDICTED: microtubule-associated protein Jupiter isoform X3
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>gi|1037095425|ref|XP_017144336.1| PREDICTED: inhibitor of growth protein 2 [Drosophila miranda
>gi|1037095441|ref|XP_017144337.1| PREDICTED: uncharacterized protein LOC108157036 [Drosophila
>gi|1037093434|ref|XP_017144188.1| PREDICTED: uncharacterized protein LOC108156938 isoform X1
>gi|1037093450|ref|XP_017144190.1| PREDICTED: rho guanine nucleotide exchange factor 17 isoform
>gi|1037098889|ref|XP_017144656.1| PREDICTED: alpha-1D adrenergic receptor isoform X2 [Drosophi
>gi|1037098901|ref|XP_017144657.1| PREDICTED: alpha-1D adrenergic receptor isoform X3 [Drosophi
>gi|1037098858|ref|XP_017144654.1| PREDICTED: tyramine receptor 1 isoform X1 [Drosophila mirand
>gi|1037098871|ref|XP_017144655.1| PREDICTED: tyramine receptor 1 isoform X1 [Drosophila mirand
>gi|1037098913|ref|XP_017144659.1| PREDICTED: 5-hydroxytryptamine receptor 1A isoform X1 [Droso
>gi|1037098925|ref|XP_017144660.1| PREDICTED: 5-hydroxytryptamine receptor 1A isoform X1 [Droso
>gi|1037098937|ref|XP_017144661.1| PREDICTED: uncharacterized protein LOC108157216 isoform X2
>gi|1037098949|ref|XP_017144662.1| PREDICTED: uncharacterized protein LOC108157216 isoform X2
>gi|1037102468|ref|XP_017144980.1| PREDICTED: lutropin-choriogonadotropic hormone receptor [Dro
>gi|1037102478|ref|XP_017144981.1| PREDICTED: gamma-aminobutyric acid receptor-associated prote
>gi|1037063268|ref|XP_017142066.1| PREDICTED: trithorax group protein osa isoform X2 [Drosophil
>gi|1037063298|ref|XP_017142068.1| PREDICTED: trithorax group protein osa isoform X4 [Drosophil
>gi|1037063254|ref|XP_017142065.1| PREDICTED: trithorax group protein osa isoform X1 [Drosophil
>gi|1037063284|ref|XP_017142067.1| PREDICTED: trithorax group protein osa isoform X3 [Drosophil
>gi|1037083433|ref|XP_017143464.1| PREDICTED: protein YSC84 [Drosophila miranda]
>gi|1037083447|ref|XP_017143465.1| PREDICTED: protein YSC84 [Drosophila miranda]
>gi|1037083461|ref|XP_017143466.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083475|ref|XP_017143467.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083489|ref|XP_017143468.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083503|ref|XP_017143469.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083517|ref|XP_017143470.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037053454|ref|XP_017141394.1| PREDICTED: zinc finger protein 420 [Drosophila miranda]
>gi|1037053671|ref|XP_017141409.1| PREDICTED: transcription initiation factor TFIID subunit 12
>gi|1037053681|ref|XP_017141410.1| PREDICTED: transcription initiation factor TFIID subunit 12
>gi|1037053597|ref|XP_017141404.1| PREDICTED: heme oxygenase 1 [Drosophila miranda]
>gi|1037053470|ref|XP_017141395.1| PREDICTED: uncharacterized protein LOC108155222 [Drosophila
>gi|1037053549|ref|XP_017141401.1| PREDICTED: uncharacterized protein LOC108155225 [Drosophila
>gi|1037053581|ref|XP_017141403.1| PREDICTED: uncharacterized protein LOC108155227 [Drosophila
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>gi|1037053629|ref|XP_017141406.1| PREDICTED: protein Fer3 [Drosophila miranda]
>gi|1037053643|ref|XP_017141407.1| PREDICTED: protein Fer3 [Drosophila miranda]
>gi|1037053659|ref|XP_017141408.1| PREDICTED: protein Fer3 [Drosophila miranda]
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>gi|1037053533|ref|XP_017141399.1| PREDICTED: uncharacterized protein LOC108155224 [Drosophila
>gi|1037059111|ref|XP_017141778.1| PREDICTED: thiamin pyrophosphokinase 1 [Drosophila miranda]
>gi|1037059081|ref|XP_017141776.1| PREDICTED: Bloom syndrome protein homolog [Drosophila miran
>gi|1037059097|ref|XP_017141777.1| PREDICTED: WD repeat-containing protein 55 homolog [Drosophi
>gi|1037059067|ref|XP_017141774.1| PREDICTED: putative uncharacterized protein DDB_G0291608 [Dr
>gi|1037068272|ref|XP_017142395.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1037068284|ref|XP_017142396.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1037061788|ref|XP_017141963.1| PREDICTED: zinc finger protein 92 homolog [Drosophila miran
>gi|1037061736|ref|XP_017141959.1| PREDICTED: uncharacterized protein LOC108155570 [Drosophila
>gi|1037061754|ref|XP_017141960.1| PREDICTED: glycine receptor subunit alpha-4 isoform X1 [Dros
>gi|1037061770|ref|XP_017141962.1| PREDICTED: glycine receptor subunit alpha-4 isoform X2 [Dros
>gi|1037074159|ref|XP_017142802.1| PREDICTED: myotubularin-related protein 13 isoform X2 [Droso
>gi|1037074143|ref|XP_017142801.1| PREDICTED: myotubularin-related protein 13 isoform X1 [Droso
>gi|1037097317|ref|XP_017144504.1| PREDICTED: chloride channel protein 2 isoform X8 [Drosophila
>gi|1037097280|ref|XP_017144500.1| PREDICTED: chloride channel protein 2 isoform X5 [Drosophila
>gi|1037097292|ref|XP_017144501.1| PREDICTED: chloride channel protein 2 isoform X6 [Drosophila
>gi|1037097256|ref|XP_017144498.1| PREDICTED: chloride channel protein 2 isoform X3 [Drosophila
>gi|1037097245|ref|XP_017144497.1| PREDICTED: chloride channel protein 2 isoform X2 [Drosophila
>gi|1037097231|ref|XP_017144496.1| PREDICTED: chloride channel protein 2 isoform X1 [Drosophila
>gi|1037097306|ref|XP_017144503.1| PREDICTED: chloride channel protein 2 isoform X7 [Drosophila
>gi|1037097268|ref|XP_017144499.1| PREDICTED: chloride channel protein 2 isoform X4 [Drosophila
>gi|1037097329|ref|XP_017144505.1| PREDICTED: chloride channel protein 2 isoform X9 [Drosophila
>gi|1037089082|ref|XP_017143864.1| PREDICTED: heterogeneous nuclear ribonucleoprotein H2 isofor
>gi|1037089096|ref|XP_017143865.1| PREDICTED: heterogeneous nuclear ribonucleoprotein H2 isofor
>gi|1037085476|ref|XP_017143618.1| PREDICTED: uncharacterized protein LOC108156590 [Drosophila
>gi|1037085462|ref|XP_017143617.1| PREDICTED: LOW QUALITY PROTEIN: nuclear pore complex protei
>gi|1037096594|ref|XP_017144441.1| PREDICTED: protein vreteno-like [Drosophila miranda]
>gi|1037103575|ref|XP_017145099.1| PREDICTED: dynein heavy chain 5, axonemal isoform X1 [Drosop
>gi|1037103583|ref|XP_017145100.1| PREDICTED: dynein heavy chain 5, axonemal isoform X2 [Drosop
>gi|1037103591|ref|XP_017145101.1| PREDICTED: uncharacterized protein LOC108157511 [Drosophila
>gi|1037060274|ref|XP_017141861.1| PREDICTED: uncharacterized protein LOC108155523 isoform X2
>gi|1037060241|ref|XP_017141859.1| PREDICTED: uncharacterized protein LOC108155523 isoform X1
>gi|1037060259|ref|XP_017141860.1| PREDICTED: uncharacterized protein LOC108155523 isoform X1
>gi|1037060289|ref|XP_017141862.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1037060179|ref|XP_017141854.1| PREDICTED: polycomb protein Scm [Drosophila miranda]
>gi|1037060305|ref|XP_017141863.1| PREDICTED: probable fatty acid-binding protein [Drosophila m
>gi|1037060225|ref|XP_017141858.1| PREDICTED: putative tricarboxylate transport protein, mitoch
>gi|1037060194|ref|XP_017141855.1| PREDICTED: 8-oxo-dGDP phosphatase NUDT18 [Drosophila miranda
>gi|1037060209|ref|XP_017141856.1| PREDICTED: limbic system-associated membrane protein [Drosop
>gi|1037033057|ref|XP_017140075.1| PREDICTED: hemicentin-2 [Drosophila miranda]
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>gi|1037059627|ref|XP_017141813.1| PREDICTED: cilia- and flagella-associated protein 53 [Drosop
>gi|1037059641|ref|XP_017141815.1| PREDICTED: cilia- and flagella-associated protein 53 [Drosop
>gi|1037059693|ref|XP_017141819.1| PREDICTED: peptidoglycan-recognition protein LB isoform X1
>gi|1037059719|ref|XP_017141821.1| PREDICTED: peptidoglycan-recognition protein LB isoform X3
>gi|1037059705|ref|XP_017141820.1| PREDICTED: peptidoglycan-recognition protein LB isoform X2
>gi|1037059737|ref|XP_017141822.1| PREDICTED: peptidoglycan-recognition protein LB isoform X4
>gi|1037059601|ref|XP_017141811.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1037059562|ref|XP_017141808.1| PREDICTED: glutamic acid-rich protein isoform X1 [Drosophila
>gi|1037059576|ref|XP_017141809.1| PREDICTED: glutamic acid-rich protein isoform X1 [Drosophila
>gi|1037059587|ref|XP_017141810.1| PREDICTED: glutamic acid-rich protein isoform X1 [Drosophila
>gi|1037059667|ref|XP_017141817.1| PREDICTED: solute carrier family 35 member G1 [Drosophila m
>gi|1037059614|ref|XP_017141812.1| PREDICTED: exonuclease 3'-5' domain-containing protein 2 [Dr
>gi|1037059654|ref|XP_017141816.1| PREDICTED: soluble calcium-activated nucleotidase 1 [Drosopl
>gi|1037059679|ref|XP_017141818.1| PREDICTED: solute carrier family 35 member B1 homolog [Dros
>gi|1037056045|ref|XP_017141566.1| PREDICTED: uncharacterized protein LOC108155333 isoform X2
>gi|1037056171|ref|XP_017141574.1| PREDICTED: nephrocan [Drosophila miranda]
>gi|1037056187|ref|XP_017141575.1| PREDICTED: leucine-rich repeat-containing protein 15 [Drosop
>gi|1037056155|ref|XP_017141573.1| PREDICTED: sensory neuron membrane protein 1 [Drosophila mi
>gi|1037056030|ref|XP_017141565.1| PREDICTED: uncharacterized protein LOC108155333 isoform X1
>gi|1037056125|ref|XP_017141571.1| PREDICTED: uncharacterized protein LOC108155333 isoform X5
>gi|1037056062|ref|XP_017141567.1| PREDICTED: uncharacterized protein LOC108155333 isoform X3
>gi|1037056077|ref|XP_017141568.1| PREDICTED: uncharacterized protein LOC108155333 isoform X3
>gi|1037056090|ref|XP_017141569.1| PREDICTED: uncharacterized protein LOC108155333 isoform X3
>gi|1037056107|ref|XP_017141570.1| PREDICTED: uncharacterized protein LOC108155333 isoform X4
>gi|1037056141|ref|XP_017141572.1| PREDICTED: 5'-AMP-activated protein kinase subunit gamma-1
>gi|1037051148|ref|XP_017141239.1| PREDICTED: SET and MYND domain-containing protein 5 [Drosopl
>gi|1037051180|ref|XP_017141241.1| PREDICTED: transmembrane emp24 domain-containing protein ec
>gi|1037051197|ref|XP_017141242.1| PREDICTED: DDRGK domain-containing protein 1 [Drosophila mi
>gi|1037051211|ref|XP_017141243.1| PREDICTED: transmembrane emp24 domain-containing protein ec
>gi|1037051004|ref|XP_017141229.1| PREDICTED: solute carrier family 15 member 1 [Drosophila mi
>gi|1037051162|ref|XP_017141240.1| PREDICTED: nucleoside diphosphate kinase 7 [Drosophila mirar
>gi|1037051275|ref|XP_017141247.1| PREDICTED: uncharacterized protein CG16817 isoform X4 [Dros
>gi|1037051260|ref|XP_017141246.1| PREDICTED: uncharacterized protein CG16817 isoform X3 [Dros
>gi|1037051246|ref|XP_017141245.1| PREDICTED: uncharacterized protein CG16817 isoform X2 [Dros
>gi|1037051231|ref|XP_017141244.1| PREDICTED: uncharacterized protein CG16817 isoform X1 [Dros
>gi|1037051287|ref|XP_017141249.1| PREDICTED: cytochrome c oxidase subunit 5A, mitochondrial [[
>gi|1037051036|ref|XP_017141231.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform )
>gi|1037051020|ref|XP_017141230.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform )
>gi|1037051052|ref|XP_017141232.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform )
>gi|1037051068|ref|XP_017141233.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform )
>gi|1037051082|ref|XP_017141234.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform )
>gi|1037051098|ref|XP_017141235.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform )
>gi|1037051114|ref|XP_017141237.1| PREDICTED: trophoblast glycoprotein [Drosophila miranda]
>gi|1037051132|ref|XP_017141238.1| PREDICTED: trophoblast glycoprotein [Drosophila miranda]
>gi|1037091720|ref|XP_017144070.1| PREDICTED: cytochrome P450 315a1, mitochondrial isoform X1
>gi|1037091734|ref|XP_017144071.1| PREDICTED: cytochrome P450 315a1, mitochondrial isoform X2
>gi|1037091706|ref|XP_017144069.1| PREDICTED: sphingomyelin phosphodiesterase 4 [Drosophila mi
>gi|1037068299|ref|XP_017142397.1| PREDICTED: intron-binding protein aquarius isoform X1 [Dros
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>gi|1037068346|ref|XP_017142400.1| PREDICTED: probable G-protein coupled receptor Mth-like 5 is
>gi|1037068329|ref|XP_017142399.1| PREDICTED: probable G-protein coupled receptor Mth-like 5 is
>gi|1037068314|ref|XP_017142398.1| PREDICTED: intron-binding protein aquarius isoform X2 [Drosc
>gi|1037068361|ref|XP_017142401.1| PREDICTED: putative inner dynein arm light chain, axonemal
>gi|1037104540|ref|XP_017145228.1| PREDICTED: histone-like protein 18C [Drosophila miranda]
>gi|1037062089|ref|XP_017141984.1| PREDICTED: cadherin-87A [Drosophila miranda]
>gi|1037062108|ref|XP_017141985.1| PREDICTED: uncharacterized protein LOC108155587 isoform X1
>gi|1037062124|ref|XP_017141986.1| PREDICTED: uncharacterized protein LOC108155587 isoform X2
>gi|1037077061|ref|XP_017142999.1| PREDICTED: hormone receptor 4 isoform X1 [Drosophila miranda
>gi|1037077075|ref|XP_017143000.1| PREDICTED: hormone receptor 4 isoform X2 [Drosophila miranda
>gi|1037082883|ref|XP_017143421.1| PREDICTED: double-strand-break repair protein rad21-like pro
>gi|1037103848|ref|XP_017145135.1| PREDICTED: uncharacterized protein LOC108157543 [Drosophila
>gi|1037095355|ref|XP_017144331.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037093420|ref|XP_017144187.1| PREDICTED: piezo-type mechanosensitive ion channel component
>gi|1037100163|ref|XP_017144758.1| PREDICTED: homeobox protein Nkx-2.1 isoform X3 [Drosophila r
>gi|1037100151|ref|XP_017144756.1| PREDICTED: homeobox protein Nkx-2.1 isoform X2 [Drosophila r
>gi|1037100127|ref|XP_017144754.1| PREDICTED: homeobox protein Nkx-2.1 isoform X1 [Drosophila r
>gi|1037100139|ref|XP_017144755.1| PREDICTED: homeobox protein Nkx-2.1 isoform X1 [Drosophila r
>gi|1037083866|ref|XP_017143498.1| PREDICTED: poly [ADP-ribose] polymerase [Drosophila miranda]
>gi|1037099067|ref|XP_017144671.1| PREDICTED: programmed cell death protein 6 [Drosophila miran
>gi|1037096627|ref|XP_017144444.1| PREDICTED: ubiquitin-conjugating enzyme E2 G1 [Drosophila mi
>gi|1037102488|ref|XP_017144982.1| PREDICTED: uncharacterized protein LOC108157439 [Drosophila
>gi|1037092806|ref|XP_017144143.1| PREDICTED: calcium/calmodulin-dependent protein kinase kinas
>gi|1037092820|ref|XP_017144144.1| PREDICTED: calcium/calmodulin-dependent protein kinase kinas
>gi|1037092834|ref|XP_017144145.1| PREDICTED: uncharacterized protein LOC108156906 [Drosophila
>gi|1037037576|ref|XP_017140336.1| PREDICTED: uncharacterized protein LOC108154544 [Drosophila
>gi|1037047298|ref|XP_017140975.1| PREDICTED: rabankyrin-5 [Drosophila miranda]
>gi|1037047360|ref|XP_017140980.1| PREDICTED: uncharacterized protein LOC108154973 isoform X1
>gi|1037047376|ref|XP_017140981.1| PREDICTED: uncharacterized protein LOC108154973 isoform X2
>gi|1037047420|ref|XP_017140984.1| PREDICTED: cytochrome b-c1 complex subunit 6, mitochondrial
>gi|1037047284|ref|XP_017140974.1| PREDICTED: histone-lysine N-methyltransferase SETD1 [Drosopl
>gi|1037047408|ref|XP_017140983.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1037047392|ref|XP_017140982.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1037047312|ref|XP_017140976.1| PREDICTED: uncharacterized protein LOC108154972 isoform X1
>gi|1037047328|ref|XP_017140977.1| PREDICTED: uncharacterized protein LOC108154972 isoform X2
>gi|1037047344|ref|XP_017140978.1| PREDICTED: uncharacterized protein LOC108154972 isoform X2
>gi|1037047251|ref|XP_017140972.1| PREDICTED: fatty acid synthase isoform X1 [Drosophila miranc
>gi|1037047269|ref|XP_017140973.1| PREDICTED: fatty acid synthase isoform X2 [Drosophila miranc
>gi|1037106221|ref|XP_017145467.1| PREDICTED: uncharacterized protein LOC108157788 [Drosophila
>gi|1037106229|ref|XP_017145468.1| PREDICTED: uncharacterized protein LOC108157788 [Drosophila
>gi|1037106235|ref|XP_017145469.1| PREDICTED: uncharacterized protein LOC108157788 [Drosophila
>gi|1037106578|ref|XP_017145515.1| PREDICTED: tensin-2-like [Drosophila miranda]
>gi|1037106586|ref|XP_017145516.1| PREDICTED: tensin-2-like [Drosophila miranda]
>gi|1037102963|ref|XP_017145030.1| PREDICTED: chymotrypsin-like protease CTRL-1 [Drosophila mir
>gi|1037102973|ref|XP_017145031.1| PREDICTED: chymotrypsin-like protease CTRL-1 [Drosophila mir
>gi|1037095244|ref|XP_017144323.1| PREDICTED: uncharacterized protein LOC108157024 isoform X2
>gi|1037095216|ref|XP_017144321.1| PREDICTED: uncharacterized protein LOC108157024 isoform X1
>gi|1037095230|ref|XP_017144322.1| PREDICTED: uncharacterized protein LOC108157024 isoform X1
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>gi|1037102509|ref|XP_017144984.1| PREDICTED: uncharacterized protein LOC108157441 isoform X1
>gi|1037102521|ref|XP_017144985.1| PREDICTED: uncharacterized protein LOC108157441 isoform X1
>gi|1037102529|ref|XP_017144986.1| PREDICTED: uncharacterized protein LOC108157441 isoform X2
>gi|1037038977|ref|XP_017140420.1| PREDICTED: uncharacterized protein LOC108154614 [Drosophila
>gi|1037096789|ref|XP_017144456.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096824|ref|XP_017144459.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096800|ref|XP_017144457.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096836|ref|XP_017144460.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096812|ref|XP_017144458.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096848|ref|XP_017144461.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096862|ref|XP_017144462.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037038253|ref|XP_017140374.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1037038239|ref|XP_017140373.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037038269|ref|XP_017140375.1| PREDICTED: uncharacterized protein LOC108154582 isoform X3
>gi|1037106485|ref|XP_017145504.1| PREDICTED: alpha-N-acetylgalactosaminidase-like isoform X1
>gi|1037106493|ref|XP_017145505.1| PREDICTED: alpha-N-acetylgalactosaminidase-like isoform X2
>gi|1037078991|ref|XP_017143138.1| PREDICTED: uncharacterized protein LOC108156277 isoform X1
>gi|1037079002|ref|XP_017143139.1| PREDICTED: uncharacterized protein LOC108156277 isoform X2
>gi|1037079030|ref|XP_017143141.1| PREDICTED: drosulfakinins [Drosophila miranda]
>gi|1037079016|ref|XP_017143140.1| PREDICTED: beta-mannosidase-like [Drosophila miranda]
>gi|1037093464|ref|XP_017144191.1| PREDICTED: beta-mannosidase [Drosophila miranda]
>gi|1037093478|ref|XP_017144192.1| PREDICTED: uncharacterized protein LOC108156940 [Drosophila
>gi|1037093492|ref|XP_017144193.1| PREDICTED: uncharacterized protein LOC108156940 [Drosophila
>gi|1037106743|ref|XP_017145538.1| PREDICTED: uncharacterized protein LOC108157846 [Drosophila
>gi|1037068068|ref|XP_017142381.1| PREDICTED: cyclin-G-associated kinase [Drosophila miranda]
>gi|1037068083|ref|XP_017142382.1| PREDICTED: guanine deaminase [Drosophila miranda]
>gi|1037068099|ref|XP_017142383.1| PREDICTED: pre-mRNA-splicing factor RBM22 [Drosophila mirand
>gi|1037081173|ref|XP_017143294.1| PREDICTED: ATP-dependent RNA helicase abstrakt [Drosophila m
>gi|1037081160|ref|XP_017143293.1| PREDICTED: gelsolin isoform X2 [Drosophila miranda]
>gi|1037081148|ref|XP_017143292.1| PREDICTED: gelsolin isoform X1 [Drosophila miranda]
>gi|1037081187|ref|XP_017143295.1| PREDICTED: serine protease snake isoform X1 [Drosophila mira
>gi|1037081201|ref|XP_017143296.1| PREDICTED: uncharacterized protein LOC108156389 isoform X2
>gi|1037081215|ref|XP_017143297.1| PREDICTED: uncharacterized protein LOC108156390 [Drosophila
>gi|1037036907|ref|XP_017140295.1| PREDICTED: uncharacterized protein LOC108154506 [Drosophila
>gi|1037038787|ref|XP_017140408.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037035752|ref|XP_017140227.1| PREDICTED: adenylate cyclase, terminal-differentiation speci
>gi|1037038088|ref|XP_017140362.1| PREDICTED: uncharacterized protein LOC108154573 [Drosophila
>gi|1037035631|ref|XP_017140220.1| PREDICTED: odorant receptor 82a [Drosophila miranda]
>gi|1037033818|ref|XP_017140120.1| PREDICTED: omega-amidase NIT2-like [Drosophila miranda]
>gi|1037090527|ref|XP_017143981.1| PREDICTED: uncharacterized protein LOC108156805 [Drosophila
>gi|1037090543|ref|XP_017143982.1| PREDICTED: ankyrin repeat, PH and SEC7 domain containing pro
>gi|1037090557|ref|XP_017143983.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090571|ref|XP_017143985.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090585|ref|XP_017143986.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090599|ref|XP_017143987.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090613|ref|XP_017143988.1| PREDICTED: complexin isoform X2 [Drosophila miranda]
>gi|1037090627|ref|XP_017143989.1| PREDICTED: complexin isoform X3 [Drosophila miranda]
>gi|1037090641|ref|XP_017143990.1| PREDICTED: complexin isoform X4 [Drosophila miranda]
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>gi|1037090514|ref|XP_017143980.1| PREDICTED: uncharacterized protein LOC108156804 [Drosophila
>gi|1037096934|ref|XP_017144468.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1037096946|ref|XP_017144469.1| PREDICTED: charged multivesicular body protein 3 [Drosophila
>gi|1037092548|ref|XP_017144128.1| PREDICTED: ATP-dependent RNA helicase SUV3 homolog, mitochon
>gi|1037092562|ref|XP_017144129.1| PREDICTED: synaptotagmin-16 [Drosophila miranda]
>gi|1037071455|ref|XP_017142614.1| PREDICTED: uncharacterized protein LOC108155960 isoform X1
>gi|1037071471|ref|XP_017142615.1| PREDICTED: uncharacterized protein LOC108155960 isoform X1
>gi|1037071487|ref|XP_017142617.1| PREDICTED: uncharacterized protein LOC108155960 isoform X1
>gi|1037071501|ref|XP_017142618.1| PREDICTED: uncharacterized protein LOC108155960 isoform X2
>gi|1037071549|ref|XP_017142621.1| PREDICTED: AF4/FMR2 family member 4 isoform X5 [Drosophila m
>gi|1037071533|ref|XP_017142620.1| PREDICTED: GATA zinc finger domain-containing protein 6 iso
>gi|1037071567|ref|XP_017142622.1| PREDICTED: AF4/FMR2 family member 4 isoform X6 [Drosophila m
>gi|1037071517|ref|XP_017142619.1| PREDICTED: uncharacterized protein LOC108155960 isoform X3
>gi|1037071423|ref|XP_017142612.1| PREDICTED: zinc finger matrin-type protein CG9776 isoform X
>gi|1037071439|ref|XP_017142613.1| PREDICTED: zinc finger matrin-type protein CG9776 isoform X
>gi|1037076263|ref|XP_017142946.1| PREDICTED: uncharacterized protein LOC108156143 [Drosophila
>gi|1037076216|ref|XP_017142944.1| PREDICTED: S-phase kinase-associated protein 2 [Drosophila m
>gi|1037076233|ref|XP_017142945.1| PREDICTED: uncharacterized protein LOC108156141 [Drosophila
>gi|1037103629|ref|XP_017145107.1| PREDICTED: Krueppel-like factor 16 [Drosophila miranda]
>gi|1037100943|ref|XP_017144834.1| PREDICTED: probable sodium/potassium/calcium exchanger CG109
>gi|1037101911|ref|XP_017144922.1| PREDICTED: sideroflexin-1 [Drosophila miranda]
>gi|1037079115|ref|XP_017143148.1| PREDICTED: contactin [Drosophila miranda]
>gi|1037085089|ref|XP_017143592.1| PREDICTED: protein Tube [Drosophila miranda]
>gi|1037085101|ref|XP_017143593.1| PREDICTED: E3 ubiquitin-protein ligase TM129 [Drosophila mi
>gi|1037085115|ref|XP_017143594.1| PREDICTED: uncharacterized protein LOC108156571 [Drosophila
>gi|1037097506|ref|XP_017144521.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 [Drosophila m
>gi|1037097520|ref|XP_017144522.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 [Drosophila m
>gi|1037022918|ref|XP_017139450.1| PREDICTED: Golgi to ER traffic protein 4 homolog, partial [
>gi|1037093255|ref|XP_017144174.1| PREDICTED: Golgi to ER traffic protein 4 homolog [Drosophila
>gi|1037093269|ref|XP_017144175.1| PREDICTED: Golgi to ER traffic protein 4 homolog [Drosophila
>gi|1037093283|ref|XP_017144177.1| PREDICTED: BTB/POZ domain-containing protein KCTD9 isoform
>gi|1037093297|ref|XP_017144178.1| PREDICTED: BTB/POZ domain-containing protein KCTD9 isoform
>gi|1037080299|ref|XP_017143226.1| PREDICTED: methenyltetrahydrofolate synthase domain-containi
>gi|1037080265|ref|XP_017143224.1| PREDICTED: serine/threonine-protein kinase 32A isoform X1 [
>gi|1037080279|ref|XP_017143225.1| PREDICTED: serine/threonine-protein kinase 32A isoform X2 [
>gi|1037080331|ref|XP_017143228.1| PREDICTED: uncharacterized protein LOC108156340 [Drosophila
>gi|1037080313|ref|XP_017143227.1| PREDICTED: ubiquilin [Drosophila miranda]
>gi|1037060524|ref|XP_017141879.1| PREDICTED: uncharacterized protein LOC108155529 [Drosophila
>gi|1037060551|ref|XP_017141880.1| PREDICTED: uncharacterized protein LOC108155530 isoform X1
>gi|1037060565|ref|XP_017141881.1| PREDICTED: uncharacterized protein LOC108155530 isoform X2
>gi|1037060508|ref|XP_017141878.1| PREDICTED: eukaryotic translation initiation factor 3 subun
>gi|1037060066|ref|XP_017141846.1| PREDICTED: tRNA (guanine(10)-N2)-methyltransferase homolog
>gi|1037060163|ref|XP_017141853.1| PREDICTED: putative lipoyltransferase 2, mitochondrial [Dros
>gi|1037060034|ref|XP_017141844.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037060050|ref|XP_017141845.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037060084|ref|XP_017141847.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060102|ref|XP_017141849.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060118|ref|XP_017141850.1| PREDICTED: elongation of very long chain fatty acids protei
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>gi|1037060133|ref|XP_017141851.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060149|ref|XP_017141852.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078713|ref|XP_017143116.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078725|ref|XP_017143117.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078737|ref|XP_017143118.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078749|ref|XP_017143119.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037069646|ref|XP_017142493.1| PREDICTED: uncharacterized protein LOC108155878 isoform X2
>gi|1037069674|ref|XP_017142494.1| PREDICTED: uncharacterized protein LOC108155878 isoform X2
>gi|1037069690|ref|XP_017142495.1| PREDICTED: uncharacterized protein LOC108155879 [Drosophila
>gi|1037069706|ref|XP_017142496.1| PREDICTED: uncharacterized protein LOC108155879 [Drosophila
>gi|1037069614|ref|XP_017142490.1| PREDICTED: uncharacterized protein LOC108155878 isoform X1
>gi|1037069632|ref|XP_017142491.1| PREDICTED: uncharacterized protein LOC108155878 isoform X1
>gi|1037048108|ref|XP_017141026.1| PREDICTED: maternal protein pumilio isoform X1 [Drosophila m
>gi|1037048124|ref|XP_017141027.1| PREDICTED: maternal protein pumilio isoform X2 [Drosophila m
>gi|1037048189|ref|XP_017141032.1| PREDICTED: maternal protein pumilio isoform X6 [Drosophila m
>gi|1037048158|ref|XP_017141029.1| PREDICTED: maternal protein pumilio isoform X4 [Drosophila m
>gi|1037048174|ref|XP_017141031.1| PREDICTED: maternal protein pumilio isoform X5 [Drosophila m
>gi|1037048142|ref|XP_017141028.1| PREDICTED: maternal protein pumilio isoform X3 [Drosophila m
>gi|1037048205|ref|XP_017141033.1| PREDICTED: uncharacterized protein LOC108154995 [Drosophila
>gi|1037048221|ref|XP_017141034.1| PREDICTED: uncharacterized protein LOC108154996 [Drosophila
>gi|1037071687|ref|XP_017142632.1| PREDICTED: chromosomal protein D1 [Drosophila miranda]
>gi|1037071701|ref|XP_017142633.1| PREDICTED: chromosomal protein D1 [Drosophila miranda]
>gi|1037071673|ref|XP_017142631.1| PREDICTED: dipeptidyl peptidase 9 [Drosophila miranda]
>gi|1037071712|ref|XP_017142634.1| PREDICTED: acylphosphatase-2 [Drosophila miranda]
>gi|1037039892|ref|XP_017140482.1| PREDICTED: uncharacterized protein LOC108154656 [Drosophila
>gi|1037039878|ref|XP_017140481.1| PREDICTED: putative uncharacterized protein DDB_G0271606 [Dr
>gi|1037039864|ref|XP_017140480.1| PREDICTED: uncharacterized protein LOC108154654 [Drosophila
>gi|1037039756|ref|XP_017140471.1| PREDICTED: HEAT repeat-containing protein 5B isoform X2 [Dro
>gi|1037039740|ref|XP_017140470.1| PREDICTED: HEAT repeat-containing protein 5B isoform X1 [Dro
>gi|1037039772|ref|XP_017140472.1| PREDICTED: HEAT repeat-containing protein 5B isoform X3 [Dro
>gi|1037039914|ref|XP_017140483.1| PREDICTED: uncharacterized protein LOC108154657 [Drosophila
>gi|1037039944|ref|XP_017140485.1| PREDICTED: probable 18S rRNA (guanine-N(7))-methyltransferas
>gi|1037039958|ref|XP_017140486.1| PREDICTED: dnaJ homolog subfamily C member 30 [Drosophila mi
>gi|1037039848|ref|XP_017140479.1| PREDICTED: protein unc-45 homolog B [Drosophila miranda]
>gi|1037039928|ref|XP_017140484.1| PREDICTED: aquaporin-2 [Drosophila miranda]
>gi|1037022936|ref|XP_017139451.1| PREDICTED: uncharacterized protein LOC108153865 [Drosophila
>gi|1037022952|ref|XP_017139453.1| PREDICTED: uncharacterized protein LOC108153866 [Drosophila
>gi|1037039786|ref|XP_017140473.1| PREDICTED: transcription factor mef2A isoform X1 [Drosophila
>gi|1037039800|ref|XP_017140476.1| PREDICTED: transcription factor mef2A isoform X1 [Drosophila
>gi|1037039816|ref|XP_017140477.1| PREDICTED: transcription factor mef2A isoform X1 [Drosophila
>gi|1037039832|ref|XP_017140478.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1037022972|ref|XP_017139454.1| PREDICTED: uncharacterized protein LOC108153867 [Drosophila
>gi|1037022990|ref|XP_017139455.1| PREDICTED: uncharacterized protein LOC108153868 [Drosophila
>gi|1037098096|ref|XP_017144578.1| PREDICTED: uncharacterized protein LOC108157170 [Drosophila
>gi|1037076471|ref|XP_017142959.1| PREDICTED: transforming acidic coiled-coil-containing protei
>gi|1037076437|ref|XP_017142957.1| PREDICTED: transforming acidic coiled-coil-containing protei
>gi|1037076421|ref|XP_017142956.1| PREDICTED: transforming acidic coiled-coil-containing protei
>gi|1037076453|ref|XP_017142958.1| PREDICTED: transforming acidic coiled-coil-containing protei
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>gi|1037101075|ref|XP_017144847.1| PREDICTED: membrane metallo-endopeptidase-like 1 isoform X1
>gi|1037101083|ref|XP_017144848.1| PREDICTED: membrane metallo-endopeptidase-like 1 isoform X2
>gi|1037093749|ref|XP_017144207.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037093707|ref|XP_017144203.1| PREDICTED: kelch-like protein 7 isoform X1 [Drosophila mira
>gi|1037093721|ref|XP_017144205.1| PREDICTED: kelch-like protein 7 isoform X2 [Drosophila mira
>gi|1037093735|ref|XP_017144206.1| PREDICTED: uncharacterized protein LOC108156948 isoform X3
>gi|1037069800|ref|XP_017142503.1| PREDICTED: formin-J isoform X2 [Drosophila miranda]
>gi|1037069768|ref|XP_017142500.1| PREDICTED: formin-J isoform X1 [Drosophila miranda]
>gi|1037069784|ref|XP_017142502.1| PREDICTED: formin-J isoform X1 [Drosophila miranda]
>gi|1037069834|ref|XP_017142505.1| PREDICTED: dual specificity protein kinase splA isoform X4
>gi|1037069850|ref|XP_017142506.1| PREDICTED: dual specificity protein kinase splA isoform X5
>gi|1037069818|ref|XP_017142504.1| PREDICTED: formin-J isoform X3 [Drosophila miranda]
>gi|1037069862|ref|XP_017142507.1| PREDICTED: uncharacterized protein CG43427 isoform X6 [Drosc
>gi|1037023008|ref|XP_017139457.1| PREDICTED: uncharacterized protein LOC108153869 [Drosophila
>gi|1037023025|ref|XP_017139458.1| PREDICTED: uncharacterized protein LOC108153870 [Drosophila
>gi|1037023043|ref|XP_017139459.1| PREDICTED: uncharacterized protein LOC108153871 [Drosophila
>gi|1037104073|ref|XP_017145163.1| PREDICTED: zinc finger protein 878 isoform X1 [Drosophila m
>gi|1037104081|ref|XP_017145165.1| PREDICTED: oocyte zinc finger protein XlCOF22 isoform X2 [D
>gi|1037095369|ref|XP_017144332.1| PREDICTED: protein SON [Drosophila miranda]
>gi|1037023061|ref|XP_017139461.1| PREDICTED: uncharacterized protein LOC108153872 [Drosophila
>gi|1037091250|ref|XP_017144035.1| PREDICTED: importin subunit alpha-4 [Drosophila miranda]
>gi|1037102314|ref|XP_017144964.1| PREDICTED: mitochondrial pyruvate carrier 2 [Drosophila mira
>gi|1037106362|ref|XP_017145486.1| PREDICTED: mitochondrial pyruvate carrier 2-like [Drosophila
>gi|1037106370|ref|XP_017145487.1| PREDICTED: mitochondrial pyruvate carrier 2-like [Drosophila
>gi|1037058944|ref|XP_017141766.1| PREDICTED: vacuolar protein sorting-associated protein 45 [
>gi|1037058914|ref|XP_017141764.1| PREDICTED: IQ and AAA domain-containing protein 1 isoform X
>gi|1037058930|ref|XP_017141765.1| PREDICTED: IQ and AAA domain-containing protein 1 isoform X2
>gi|1037059051|ref|XP_017141773.1| PREDICTED: U6 snRNA phosphodiesterase [Drosophila miranda]
>gi|1037059021|ref|XP_017141771.1| PREDICTED: metaxin-1 homolog isoform X1 [Drosophila miranda]
>gi|1037059037|ref|XP_017141772.1| PREDICTED: metaxin-1 homolog isoform X2 [Drosophila miranda]
>gi|1037059006|ref|XP_017141770.1| PREDICTED: methyl-CpG-binding domain protein 2 isoform X2 [
>gi|1037058990|ref|XP_017141769.1| PREDICTED: methyl-CpG-binding domain protein 2 isoform X1 [
>gi|1037058976|ref|XP_017141768.1| PREDICTED: AP-1 complex subunit mu-1 [Drosophila miranda]
>gi|1037058960|ref|XP_017141767.1| PREDICTED: 2-oxoisovalerate dehydrogenase subunit alpha, mit
>gi|1037092948|ref|XP_017144153.1| PREDICTED: probable tRNA (uracil-O(2)-)-methyltransferase [
>gi|1037092990|ref|XP_017144157.1| PREDICTED: RNA-binding protein with serine-rich domain 1-B
>gi|1037092976|ref|XP_017144155.1| PREDICTED: RNA-binding protein with serine-rich domain 1-B
>gi|1037092962|ref|XP_017144154.1| PREDICTED: RNA-binding protein with serine-rich domain 1-B
>gi|1037102333|ref|XP_017144966.1| PREDICTED: uncharacterized protein LOC108157428 isoform X1
>gi|1037102343|ref|XP_017144967.1| PREDICTED: uncharacterized protein LOC108157428 isoform X2
>gi|1037081782|ref|XP_017143342.1| PREDICTED: DNA polymerase alpha catalytic subunit [Drosophil
>gi|1037081795|ref|XP_017143343.1| PREDICTED: uncharacterized protein LOC108156409 [Drosophila
>gi|1037063920|ref|XP_017142111.1| PREDICTED: uncharacterized protein LOC108155660 [Drosophila
>gi|1037063933|ref|XP_017142112.1| PREDICTED: protein archease-like [Drosophila miranda]
>gi|1037063858|ref|XP_017142107.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037063873|ref|XP_017142108.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037063889|ref|XP_017142109.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037063906|ref|XP_017142110.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
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>gi|1037023078|ref|XP_017139462.1| PREDICTED: GTPase-activating protein RacGAP84C-like, partial
>gi|1037097794|ref|XP_017144549.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037097784|ref|XP_017144548.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037097803|ref|XP_017144550.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037034374|ref|XP_017140155.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-1
>gi|1037034391|ref|XP_017140156.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-1
>gi|1037035081|ref|XP_017140196.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-1
>gi|1037062140|ref|XP_017141987.1| PREDICTED: insulin-like receptor [Drosophila miranda]
>gi|1037062156|ref|XP_017141988.1| PREDICTED: insulin-like receptor [Drosophila miranda]
>gi|1037023095|ref|XP_017139463.1| PREDICTED: acyl-CoA Delta(11) desaturase-like [Drosophila mi
>gi|1037096110|ref|XP_017144395.1| PREDICTED: adenosine 3'-phospho 5'-phosphosulfate transporte
>gi|1037096123|ref|XP_017144396.1| PREDICTED: uncharacterized protein LOC108157072 [Drosophila
>gi|1037038062|ref|XP_017140360.1| PREDICTED: pupal cuticle protein Edg-91 isoform X2 [Drosophi
>gi|1037038050|ref|XP_017140359.1| PREDICTED: pupal cuticle protein Edg-91 isoform X1 [Drosophi
>gi|1037038961|ref|XP_017140419.1| PREDICTED: keratin-associated protein 19-2 [Drosophila mira
>gi|1037037891|ref|XP_017140352.1| PREDICTED: pupal cuticle protein Edg-91 [Drosophila miranda]
>gi|1037040857|ref|XP_017140548.1| PREDICTED: acanthoscurrin-1 [Drosophila miranda]
>gi|1037023113|ref|XP_017139464.1| PREDICTED: translation initiation factor IF-2 [Drosophila mi
>gi|1037023131|ref|XP_017139465.1| PREDICTED: uncharacterized protein LOC108153876 [Drosophila
>gi|1037035649|ref|XP_017140221.1| PREDICTED: attacin-A [Drosophila miranda]
>gi|1037106689|ref|XP_017145531.1| PREDICTED: uncharacterized protein LOC108157839 [Drosophila
>gi|1037050510|ref|XP_017141192.1| PREDICTED: uncharacterized protein LOC108155085 isoform X1
>gi|1037050524|ref|XP_017141193.1| PREDICTED: uncharacterized protein LOC108155085 isoform X1
>gi|1037050540|ref|XP_017141194.1| PREDICTED: uncharacterized protein LOC108155085 isoform X2
>gi|1037050634|ref|XP_017141201.1| PREDICTED: bax inhibitor 1-like [Drosophila miranda]
>gi|1037050618|ref|XP_017141200.1| PREDICTED: transmembrane protein 192 [Drosophila miranda]
>gi|1037050602|ref|XP_017141198.1| PREDICTED: serine/threonine-protein kinase shk1/pak1 [Drosop
>gi|1037050586|ref|XP_017141197.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037050554|ref|XP_017141195.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037050570|ref|XP_017141196.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037052014|ref|XP_017141297.1| PREDICTED: splicing factor 3A subunit 3 [Drosophila miranda]
>gi|1037052029|ref|XP_017141298.1| PREDICTED: mitotic spindle assembly checkpoint protein MAD2
>gi|1037051884|ref|XP_017141288.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1037051900|ref|XP_017141289.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1037051936|ref|XP_017141291.1| PREDICTED: dihydropyrimidinase isoform X2 [Drosophila miran
>gi|1037051918|ref|XP_017141290.1| PREDICTED: dihydropyrimidinase isoform X1 [Drosophila miran
>gi|1037051952|ref|XP_017141292.1| PREDICTED: dihydropyrimidinase 2 isoform X3 [Drosophila mira
>gi|1037051998|ref|XP_017141296.1| PREDICTED: dihydropyrimidinase-related protein 3 isoform X6
>gi|1037051968|ref|XP_017141294.1| PREDICTED: dihydropyrimidinase isoform X4 [Drosophila miran
>gi|1037051984|ref|XP_017141295.1| PREDICTED: dihydropyrimidinase isoform X5 [Drosophila miran
>gi|1037032364|ref|XP_017140031.1| PREDICTED: uncharacterized protein LOC108154292 isoform X1
>gi|1037032382|ref|XP_017140032.1| PREDICTED: uncharacterized protein LOC108154292 isoform X2
>gi|1037093311|ref|XP_017144179.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037093325|ref|XP_017144180.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037093339|ref|XP_017144181.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037093353|ref|XP_017144182.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037076079|ref|XP_017142934.1| PREDICTED: uncharacterized protein LOC108156132 [Drosophila
>gi|1037076064|ref|XP_017142933.1| PREDICTED: LOW QUALITY PROTEIN: tankyrase [Drosophila miran
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>gi|1037089110|ref|XP_017143866.1| PREDICTED: probable serine/threonine-protein kinase cdc7 [D
>gi|1037089128|ref|XP_017143867.1| PREDICTED: probable serine/threonine-protein kinase cdc7 [D
>gi|1037089142|ref|XP_017143868.1| PREDICTED: probable serine/threonine-protein kinase cdc7 [D
>gi|1037080415|ref|XP_017143235.1| PREDICTED: relaxin receptor 2 [Drosophila miranda]
>gi|1037080471|ref|XP_017143239.1| PREDICTED: zinc finger protein OZF [Drosophila miranda]
>gi|1037080499|ref|XP_017143241.1| PREDICTED: uncharacterized protein LOC108156351 [Drosophila
>gi|1037080457|ref|XP_017143238.1| PREDICTED: S-adenosylmethionine mitochondrial carrier prote
>gi|1037080443|ref|XP_017143237.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit gamma, mit
>gi|1037080429|ref|XP_017143236.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit gamma, mit
>gi|1037080485|ref|XP_017143240.1| PREDICTED: 60S ribosomal protein L27 [Drosophila miranda]
>gi|1037098387|ref|XP_017144610.1| PREDICTED: probable cardiolipin synthase (CMP-forming) [Dros
>gi|1037071807|ref|XP_017142640.1| PREDICTED: low-density lipoprotein receptor isoform X3 [Dros
>gi|1037071838|ref|XP_017142641.1| PREDICTED: low-density lipoprotein receptor isoform X4 [Dros
>gi|1037071775|ref|XP_017142638.1| PREDICTED: low-density lipoprotein receptor isoform X1 [Dros
>gi|1037071854|ref|XP_017142642.1| PREDICTED: low-density lipoprotein receptor isoform X5 [Dros
>gi|1037071791|ref|XP_017142639.1| PREDICTED: low-density lipoprotein receptor isoform X2 [Dros
>gi|1037071868|ref|XP_017142643.1| PREDICTED: low-density lipoprotein receptor isoform X6 [Dros
>gi|1037071923|ref|XP_017142645.1| PREDICTED: low-density lipoprotein receptor isoform X8 [Dros
>gi|1037071892|ref|XP_017142644.1| PREDICTED: low-density lipoprotein receptor isoform X7 [Dros
>gi|1037071941|ref|XP_017142646.1| PREDICTED: LOW QUALITY PROTEIN: very low-density lipoprotei
>gi|1037096671|ref|XP_017144445.1| PREDICTED: myosin-9-like [Drosophila miranda]
>gi|1037032969|ref|XP_017140070.1| PREDICTED: probable palmitoyltransferase ZDHHC24 [Drosophila
>gi|1037105991|ref|XP_017145435.1| PREDICTED: uncharacterized protein LOC108157764 [Drosophila
>gi|1037036977|ref|XP_017140299.1| PREDICTED: uncharacterized protein LOC108154509 isoform X2
>gi|1037036960|ref|XP_017140298.1| PREDICTED: uncharacterized protein LOC108154509 isoform X1
>gi|1037106386|ref|XP_017145489.1| PREDICTED: uncharacterized protein LOC108157806 [Drosophila
>gi|1037104422|ref|XP_017145211.1| PREDICTED: calmodulin-like [Drosophila miranda]
>gi|1037104432|ref|XP_017145212.1| PREDICTED: calmodulin-like [Drosophila miranda]
>gi|1037104440|ref|XP_017145213.1| PREDICTED: calmodulin-related protein 97A-like [Drosophila m
>gi|1037104448|ref|XP_017145214.1| PREDICTED: calmodulin-related protein 97A-like [Drosophila m
>gi|1037103667|ref|XP_017145112.1| PREDICTED: protein TolA [Drosophila miranda]
>gi|1037023147|ref|XP_017139466.1| PREDICTED: uncharacterized protein LOC108153877 [Drosophila
>gi|1037094502|ref|XP_017144266.1| PREDICTED: uncharacterized protein LOC108156984 [Drosophila
>gi|1037094516|ref|XP_017144267.1| PREDICTED: uncharacterized protein LOC108156984 [Drosophila
>gi|1037094488|ref|XP_017144264.1| PREDICTED: beta-taxilin [Drosophila miranda]
>gi|1037104804|ref|XP_017145265.1| PREDICTED: Kv channel-interacting protein 1 [Drosophila mira
>gi|1037104812|ref|XP_017145267.1| PREDICTED: Kv channel-interacting protein 1 [Drosophila mira
>gi|1037084074|ref|XP_017143514.1| PREDICTED: CD81 antigen [Drosophila miranda]
>gi|1037064178|ref|XP_017142128.1| PREDICTED: V-type proton ATPase subunit E [Drosophila mirand
>gi|1037064132|ref|XP_017142124.1| PREDICTED: protein krasavietz [Drosophila miranda]
>gi|1037064148|ref|XP_017142125.1| PREDICTED: protein krasavietz [Drosophila miranda]
>gi|1037064164|ref|XP_017142127.1| PREDICTED: uncharacterized protein LOC108155674 [Drosophila
>gi|1037064194|ref|XP_017142129.1| PREDICTED: protein KRTCAP2 homolog [Drosophila miranda]
>gi|1037094708|ref|XP_017144281.1| PREDICTED: 28S ribosomal protein S18a, mitochondrial [Drosop
>gi|1037094692|ref|XP_017144280.1| PREDICTED: pachytene checkpoint protein 2 homolog [Drosophil
>gi|1037023163|ref|XP_017139467.1| PREDICTED: putative odorant receptor 85d [Drosophila miranda
>gi|1037037628|ref|XP_017140338.1| PREDICTED: uncharacterized protein LOC108154546 [Drosophila
>gi|1037106137|ref|XP_017145455.1| PREDICTED: transmembrane protein 135-like [Drosophila miranc
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>gi|1037096683|ref|XP_017144446.1| PREDICTED: transmembrane protein 135 [Drosophila miranda]
>gi|1037096697|ref|XP_017144447.1| PREDICTED: transmembrane protein 135 [Drosophila miranda]
>gi|1037037380|ref|XP_017140324.1| PREDICTED: odorant receptor 85c [Drosophila miranda]
>gi|1037106602|ref|XP_017145519.1| PREDICTED: odorant receptor 85b-like [Drosophila miranda]
>gi|1037046571|ref|XP_017140927.1| PREDICTED: irregular chiasm C-roughest protein isoform X3 [
>gi|1037046539|ref|XP_017140925.1| PREDICTED: uncharacterized protein LOC108154943 isoform X1
>gi|1037046555|ref|XP_017140926.1| PREDICTED: uncharacterized protein LOC108154943 isoform X2
>gi|1037046605|ref|XP_017140929.1| PREDICTED: hemicentin-2 isoform X5 [Drosophila miranda]
>gi|1037046587|ref|XP_017140928.1| PREDICTED: hemicentin-2 isoform X4 [Drosophila miranda]
>gi|1037022499|ref|XP_017139424.1| PREDICTED: uncharacterized protein LOC108153837 [Drosophila
>gi|1037046633|ref|XP_017140931.1| PREDICTED: zinc finger protein 383 isoform X1 [Drosophila m:
>gi|1037046649|ref|XP_017140932.1| PREDICTED: zinc finger protein 383 isoform X2 [Drosophila m:
>gi|1037046663|ref|XP_017140933.1| PREDICTED: adenosine deaminase-like protein [Drosophila mira
>gi|1037046619|ref|XP_017140930.1| PREDICTED: protein Red [Drosophila miranda]
>gi|1037046473|ref|XP_017140920.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037046489|ref|XP_017140921.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037046505|ref|XP_017140922.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037046521|ref|XP_017140924.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037102426|ref|XP_017144975.1| PREDICTED: vasotab [Drosophila miranda]
>gi|1037102414|ref|XP_017144974.1| PREDICTED: iron-sulfur cluster assembly scaffold protein Iso
>gi|1037038601|ref|XP_017140397.1| PREDICTED: UPF0668 protein C10orf76 [Drosophila miranda]
>gi|1037038396|ref|XP_017140383.1| PREDICTED: tight junction protein ZO-1 isoform X1 [Drosophil
>gi|1037038412|ref|XP_017140384.1| PREDICTED: tight junction protein ZO-1 isoform X2 [Drosophil
>gi|1037038430|ref|XP_017140385.1| PREDICTED: tight junction protein ZO-1 isoform X3 [Drosophil
>gi|1037038446|ref|XP_017140386.1| PREDICTED: tight junction protein ZO-1 isoform X4 [Drosophil
>gi|1037038462|ref|XP_017140387.1| PREDICTED: tight junction protein ZO-1 isoform X5 [Drosophil
>gi|1037038478|ref|XP_017140389.1| PREDICTED: uncharacterized protein LOC108154591 isoform X6
>gi|1037038490|ref|XP_017140390.1| PREDICTED: uncharacterized protein LOC108154591 isoform X7
>gi|1037038538|ref|XP_017140393.1| PREDICTED: tight junction protein ZO-1 isoform X10 [Drosoph:
>gi|1037038504|ref|XP_017140391.1| PREDICTED: tight junction protein ZO-2 isoform X8 [Drosophil
>gi|1037038520|ref|XP_017140392.1| PREDICTED: tight junction protein ZO-2 isoform X9 [Drosophil
>gi|1037038553|ref|XP_017140394.1| PREDICTED: tight junction protein ZO-2 isoform X11 [Drosoph:
>gi|1037038663|ref|XP_017140401.1| PREDICTED: uncharacterized protein LOC108154597 isoform X2
>gi|1037038651|ref|XP_017140400.1| PREDICTED: uncharacterized protein LOC108154597 isoform X1
>gi|1037038619|ref|XP_017140398.1| PREDICTED: maternal effect protein oskar isoform X1 [Drosop]
>gi|1037038636|ref|XP_017140399.1| PREDICTED: maternal effect protein oskar isoform X2 [Drosop]
>gi|1037038569|ref|XP_017140395.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1037038583|ref|XP_017140396.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1037038772|ref|XP_017140407.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, i
>gi|1037038724|ref|XP_017140404.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, i
>gi|1037038740|ref|XP_017140405.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, i
>gi|1037038756|ref|XP_017140406.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, i
>gi|1037038708|ref|XP_017140403.1| PREDICTED: beta-catenin-like protein 1 [Drosophila miranda]
>gi|1037103808|ref|XP_017145130.1| PREDICTED: uncharacterized protein LOC108157538 [Drosophila
>gi|1037078833|ref|XP_017143126.1| PREDICTED: uncharacterized protein LOC108156269 [Drosophila
>gi|1037078847|ref|XP_017143127.1| PREDICTED: uncharacterized protein LOC108156269 [Drosophila
>gi|1037078889|ref|XP_017143130.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037078861|ref|XP_017143128.1| PREDICTED: UPF0528 protein CG10038 isoform X1 [Drosophila m:
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>gi|1037078875|ref|XP_017143129.1| PREDICTED: UPF0528 protein CG10038 isoform X2 [Drosophila m
>gi|1037104388|ref|XP_017145207.1| PREDICTED: uncharacterized protein LOC108157591 [Drosophila
>gi|1037092345|ref|XP_017144112.1| PREDICTED: steroid receptor seven-up, isoforms B/C isoform
>gi|1037092271|ref|XP_017144111.1| PREDICTED: steroid receptor seven-up isoform X1 [Drosophila
>gi|1037035011|ref|XP_017140192.1| PREDICTED: uncharacterized protein F12A10.7 [Drosophila mir
>gi|1037105578|ref|XP_017145377.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [
>gi|1037033490|ref|XP_017140100.1| PREDICTED: probable cytochrome P450 313a3 [Drosophila miran
>gi|1037105620|ref|XP_017145383.1| PREDICTED: probable cytochrome P450 313a1 [Drosophila miran
>gi|1037105628|ref|XP_017145384.1| PREDICTED: phosphoglycerate mutase 2 [Drosophila miranda]
>gi|1037023179|ref|XP_017139469.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037023195|ref|XP_017139470.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037023213|ref|XP_017139471.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037106780|ref|XP_017145544.1| PREDICTED: LIM domain-binding protein 3 [Drosophila miranda]
>gi|1037106213|ref|XP_017145466.1| PREDICTED: short-chain specific acyl-CoA dehydrogenase, mit
>gi|1037090820|ref|XP_017144003.1| PREDICTED: conserved oligomeric Golgi complex subunit 1 [Dr
>gi|1037090836|ref|XP_017144004.1| PREDICTED: glycine-rich protein 5 [Drosophila miranda]
>gi|1037033984|ref|XP_017140130.1| PREDICTED: 4-coumarate--CoA ligase 1 [Drosophila miranda]
>gi|1037106671|ref|XP_017145529.1| PREDICTED: adenosine kinase [Drosophila miranda]
>gi|1037105844|ref|XP_017145414.1| PREDICTED: eukaryotic translation initiation factor 3 subun
>gi|1037105276|ref|XP_017145334.1| PREDICTED: calphotin [Drosophila miranda]
>gi|1037105852|ref|XP_017145416.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1037074127|ref|XP_017142800.1| PREDICTED: uncharacterized protein LOC108156041 isoform X4
>gi|1037074061|ref|XP_017142795.1| PREDICTED: uncharacterized protein LOC108156041 isoform X1
>gi|1037074077|ref|XP_017142796.1| PREDICTED: uncharacterized protein LOC108156041 isoform X1
>gi|1037074093|ref|XP_017142797.1| PREDICTED: uncharacterized protein LOC108156041 isoform X2
>gi|1037074109|ref|XP_017142798.1| PREDICTED: uncharacterized protein LOC108156041 isoform X3
>gi|1037074029|ref|XP_017142793.1| PREDICTED: uncharacterized protein LOC108156040 [Drosophila
>gi|1037074045|ref|XP_017142794.1| PREDICTED: uncharacterized protein LOC108156040 [Drosophila
>gi|1037073931|ref|XP_017142786.1| PREDICTED: phospholipid-transporting ATPase ID isoform X5 [
>gi|1037073947|ref|XP_017142787.1| PREDICTED: phospholipid-transporting ATPase ID isoform X6 [
>gi|1037073979|ref|XP_017142790.1| PREDICTED: phospholipid-transporting ATPase ID isoform X8 [
>gi|1037073995|ref|XP_017142791.1| PREDICTED: phospholipid-transporting ATPase ID isoform X9 [
>gi|1037074011|ref|XP_017142792.1| PREDICTED: phospholipid-transporting ATPase ID isoform X10
>gi|1037073839|ref|XP_017142780.1| PREDICTED: phospholipid-transporting ATPase ID isoform X1 [
>gi|1037073854|ref|XP_017142781.1| PREDICTED: phospholipid-transporting ATPase ID isoform X1 [
>gi|1037073868|ref|XP_017142782.1| PREDICTED: phospholipid-transporting ATPase ID isoform X1 [
>gi|1037073883|ref|XP_017142783.1| PREDICTED: phospholipid-transporting ATPase ID isoform X2 [
>gi|1037073915|ref|XP_017142785.1| PREDICTED: phospholipid-transporting ATPase ID isoform X4 [
>gi|1037073899|ref|XP_017142784.1| PREDICTED: phospholipid-transporting ATPase ID isoform X3 [
>gi|1037073963|ref|XP_017142788.1| PREDICTED: phospholipid-transporting ATPase ID isoform X7 [
>gi|1037105941|ref|XP_017145428.1| PREDICTED: probable citrate synthase, mitochondrial [Drosop
>gi|1037035169|ref|XP_017140201.1| PREDICTED: ubiquitin-conjugating enzyme E2 6 [Drosophila mi
>gi|1037102021|ref|XP_017144934.1| PREDICTED: band 7 protein AGAP004871 [Drosophila miranda]
>gi|1037101966|ref|XP_017144929.1| PREDICTED: uncharacterized protein LOC108157399 [Drosophila
>gi|1037101978|ref|XP_017144930.1| PREDICTED: uncharacterized protein LOC108157400 isoform X1
>gi|1037101991|ref|XP_017144931.1| PREDICTED: band 7 protein AGAP004871-like isoform X2 [Drosop
>gi|1037102002|ref|XP_017144932.1| PREDICTED: band 7 protein AGAP004871-like isoform X3 [Drosop
>gi|1037102011|ref|XP_017144933.1| PREDICTED: band 7 protein AGAP004871-like isoform X4 [Drosop
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>gi|1037102031|ref|XP_017144935.1| PREDICTED: uncharacterized protein LOC108157402 [Drosophila
>gi|1037100552|ref|XP_017144793.1| PREDICTED: transcription initiation protein SPT3 homolog [D:
>gi|1037076930|ref|XP_017142990.1| PREDICTED: DDB1- and CUL4-associated factor 12 [Drosophila i
>gi|1037076915|ref|XP_017142989.1| PREDICTED: GRIP and coiled-coil domain-containing protein 2
>gi|1037094722|ref|XP_017144282.1| PREDICTED: kelch-like protein 18 [Drosophila miranda]
>gi|1037058204|ref|XP_017141715.1| PREDICTED: beta-galactosidase [Drosophila miranda]
>gi|1037058251|ref|XP_017141719.1| PREDICTED: tachykinins isoform X1 [Drosophila miranda]
>gi|1037058265|ref|XP_017141720.1| PREDICTED: tachykinins isoform X2 [Drosophila miranda]
>gi|1037058147|ref|XP_017141711.1| PREDICTED: transforming growth factor-beta-induced protein :
>gi|1037058162|ref|XP_017141712.1| PREDICTED: transforming growth factor-beta-induced protein :
>gi|1037058116|ref|XP_017141709.1| PREDICTED: transforming growth factor-beta-induced protein :
>gi|1037058132|ref|XP_017141710.1| PREDICTED: transforming growth factor-beta-induced protein :
>gi|1037058177|ref|XP_017141713.1| PREDICTED: transforming growth factor-beta-induced protein :
>gi|1037058187|ref|XP_017141714.1| PREDICTED: transforming growth factor-beta-induced protein :
>gi|1037058236|ref|XP_017141718.1| PREDICTED: L-galactose dehydrogenase [Drosophila miranda]
>gi|1037058218|ref|XP_017141717.1| PREDICTED: L-galactose dehydrogenase [Drosophila miranda]
>gi|1037058279|ref|XP_017141721.1| PREDICTED: uncharacterized protein LOC108155425 [Drosophila
>gi|1037032056|ref|XP_017140013.1| PREDICTED: solute carrier family 35 member F6 [Drosophila m:
>gi|1037032119|ref|XP_017140018.1| PREDICTED: mitochondrial glutamate carrier 1 [Drosophila mi:
>gi|1037032136|ref|XP_017140019.1| PREDICTED: mitochondrial glutamate carrier 1-like [Drosophil
>gi|1037032092|ref|XP_017140015.1| PREDICTED: mitochondrial glutamate carrier 1-like [Drosophil
>gi|1037032106|ref|XP_017140017.1| PREDICTED: mitochondrial glutamate carrier 1-like [Drosophil
>gi|1037031807|ref|XP_017139998.1| PREDICTED: uncharacterized protein LOC108154271 isoform X1
>gi|1037031841|ref|XP_017140000.1| PREDICTED: protein IWS1 homolog isoform X3 [Drosophila mira:
>gi|1037031823|ref|XP_017139999.1| PREDICTED: uncharacterized protein LOC108154271 isoform X2
>gi|1037032037|ref|XP_017140012.1| PREDICTED: aurora kinase C [Drosophila miranda]
>gi|1037031773|ref|XP_017139996.1| PREDICTED: zinc finger protein 829 isoform X1 [Drosophila m:
>gi|1037031791|ref|XP_017139997.1| PREDICTED: uncharacterized protein LOC108154270 isoform X2
>gi|1037031612|ref|XP_017139991.1| PREDICTED: arginine/serine-rich protein PNISR [Drosophila m:
>gi|1037031648|ref|XP_017139994.1| PREDICTED: ATPase family AAA domain-containing protein 3 [D:
>gi|1037031757|ref|XP_017139995.1| PREDICTED: ATPase family AAA domain-containing protein 3 [D:
>gi|1037031629|ref|XP_017139992.1| PREDICTED: protein asunder [Drosophila miranda]
>gi|1037032074|ref|XP_017140014.1| PREDICTED: uncharacterized protein LOC108154276 [Drosophila
>gi|1037031596|ref|XP_017139990.1| PREDICTED: nucleolar protein 6 [Drosophila miranda]
>gi|1037031954|ref|XP_017140007.1| PREDICTED: casein kinase I isoform X6 [Drosophila miranda]
>gi|1037031970|ref|XP_017140008.1| PREDICTED: casein kinase I isoform X6 [Drosophila miranda]
>gi|1037031886|ref|XP_017140003.1| PREDICTED: casein kinase I isoform X2 [Drosophila miranda]
>gi|1037031936|ref|XP_017140006.1| PREDICTED: casein kinase I isoform X5 [Drosophila miranda]
>gi|1037031986|ref|XP_017140009.1| PREDICTED: casein kinase I isoform X7 [Drosophila miranda]
>gi|1037031869|ref|XP_017140001.1| PREDICTED: casein kinase I isoform X1 [Drosophila miranda]
>gi|1037031922|ref|XP_017140005.1| PREDICTED: casein kinase I isoform X4 [Drosophila miranda]
>gi|1037031904|ref|XP_017140004.1| PREDICTED: casein kinase I isoform X3 [Drosophila miranda]
>gi|1037032001|ref|XP_017140010.1| PREDICTED: casein kinase I isoform X8 [Drosophila miranda]
>gi|1037032019|ref|XP_017140011.1| PREDICTED: casein kinase I isoform X9 [Drosophila miranda]
>gi|1037101928|ref|XP_017144924.1| PREDICTED: zinc transporter ZIP1 [Drosophila miranda]
>gi|1037082760|ref|XP_017143412.1| PREDICTED: extracellular sulfatase SULF-1 homolog [Drosophil
>gi|1037082774|ref|XP_017143413.1| PREDICTED: extracellular sulfatase SULF-1 homolog [Drosophil
>gi|1037082801|ref|XP_017143415.1| PREDICTED: facilitated trehalose transporter Tret1-like [Dro
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>gi|1037082788|ref|XP_017143414.1| PREDICTED: probable plastidic glucose transporter 2 [Drosoph
>gi|1037073595|ref|XP_017142764.1| PREDICTED: serine/arginine-rich splicing factor 1B [Drosophi
>gi|1037073611|ref|XP_017142765.1| PREDICTED: transcription factor Sp4 [Drosophila miranda]
>gi|1037073661|ref|XP_017142768.1| PREDICTED: uncharacterized protein LOC108156027 isoform X3
>gi|1037073627|ref|XP_017142766.1| PREDICTED: uncharacterized protein LOC108156027 isoform X1
>gi|1037073643|ref|XP_017142767.1| PREDICTED: uncharacterized protein LOC108156027 isoform X2
>gi|1037097565|ref|XP_017144527.1| PREDICTED: mesencephalic astrocyte-derived neurotrophic fact
>gi|1037097575|ref|XP_017144528.1| PREDICTED: uncharacterized protein LOC108157138 [Drosophila
>gi|1037102456|ref|XP_017144979.1| PREDICTED: putative SERF-like protein [Drosophila miranda]
>gi|1037104768|ref|XP_017145260.1| PREDICTED: uncharacterized protein LOC108157631 [Drosophila
>gi|1037104760|ref|XP_017145259.1| PREDICTED: uncharacterized protein LOC108157630 [Drosophila
>gi|1037105570|ref|XP_017145376.1| PREDICTED: uncharacterized protein LOC108157719 [Drosophila
>gi|1037102165|ref|XP_017144948.1| PREDICTED: circadian locomoter output cycles protein kaput
>gi|1037102144|ref|XP_017144946.1| PREDICTED: single-minded homolog 2 isoform X1 [Drosophila m
>gi|1037102154|ref|XP_017144947.1| PREDICTED: single-minded homolog 2 isoform X1 [Drosophila m
>gi|1037033470|ref|XP_017140099.1| PREDICTED: uncharacterized protein LOC108154344 isoform X4
>gi|1037033414|ref|XP_017140095.1| PREDICTED: uncharacterized protein LOC108154344 isoform X1
>gi|1037033452|ref|XP_017140098.1| PREDICTED: uncharacterized protein LOC108154344 isoform X3
>gi|1037033434|ref|XP_017140097.1| PREDICTED: uncharacterized protein LOC108154344 isoform X2
>gi|1037050839|ref|XP_017141217.1| PREDICTED: protein farnesyltransferase subunit beta [Drosoph
>gi|1037050883|ref|XP_017141220.1| PREDICTED: uncharacterized protein LOC108155102 isoform X2
>gi|1037050871|ref|XP_017141219.1| PREDICTED: armadillo repeat-containing protein 7 isoform X1
>gi|1037050901|ref|XP_017141221.1| PREDICTED: 28S ribosomal protein S33, mitochondrial [Drosoph
>gi|1037050793|ref|XP_017141212.1| PREDICTED: protein CLEC16A isoform X2 [Drosophila miranda]
>gi|1037050777|ref|XP_017141211.1| PREDICTED: protein CLEC16A isoform X1 [Drosophila miranda]
>gi|1037050855|ref|XP_017141218.1| PREDICTED: tetratricopeptide repeat protein 1 [Drosophila m
>gi|1037050825|ref|XP_017141216.1| PREDICTED: methionine synthase reductase [Drosophila miranda
>gi|1037050809|ref|XP_017141215.1| PREDICTED: serine/threonine-protein kinase PAK 2 [Drosophila
>gi|1037054821|ref|XP_017141488.1| PREDICTED: techylectin-5A [Drosophila miranda]
>gi|1037054761|ref|XP_017141484.1| PREDICTED: uncharacterized protein LOC108155281 isoform X4
>gi|1037054746|ref|XP_017141483.1| PREDICTED: uncharacterized protein LOC108155281 isoform X3
>gi|1037054732|ref|XP_017141482.1| PREDICTED: uncharacterized protein LOC108155281 isoform X2
>gi|1037054697|ref|XP_017141479.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037054714|ref|XP_017141481.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037054789|ref|XP_017141486.1| PREDICTED: mucin-5AC isoform X6 [Drosophila miranda]
>gi|1037054777|ref|XP_017141485.1| PREDICTED: mucin-5AC isoform X5 [Drosophila miranda]
>gi|1037054852|ref|XP_017141491.1| PREDICTED: acylphosphatase-2 [Drosophila miranda]
>gi|1037054836|ref|XP_017141490.1| PREDICTED: uroporphyrinogen-III synthase [Drosophila miranda
>gi|1037054805|ref|XP_017141487.1| PREDICTED: protein FAM114A2 [Drosophila miranda]
>gi|1037036661|ref|XP_017140282.1| PREDICTED: uncharacterized protein LOC108154493 [Drosophila
>gi|1037026831|ref|XP_017139698.1| PREDICTED: uncharacterized protein LOC108154093 isoform X1
>gi|1037026849|ref|XP_017139699.1| PREDICTED: uncharacterized protein LOC108154093 isoform X2
>gi|1037027074|ref|XP_017139713.1| PREDICTED: uncharacterized protein LOC108154102 [Drosophila
>gi|1037027164|ref|XP_017139719.1| PREDICTED: syntaxin-18 [Drosophila miranda]
>gi|1037027281|ref|XP_017139727.1| PREDICTED: ras-like GTP-binding protein RhoL [Drosophila mi
>gi|1037027213|ref|XP_017139722.1| PREDICTED: uncharacterized protein LOC108154110 isoform X1
>gi|1037027230|ref|XP_017139724.1| PREDICTED: SAP domain-containing ribonucleoprotein isoform
>gi|1037027025|ref|XP_017139710.1| PREDICTED: heterogeneous nuclear ribonucleoprotein M isoform
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>gi|1037027040|ref|XP_017139711.1| PREDICTED: glycine-rich RNA-binding protein isoform X2 [Dros
>gi|1037027299|ref|XP_017139728.1| PREDICTED: ras-like protein 1 [Drosophila miranda]
>gi|1037027147|ref|XP_017139718.1| PREDICTED: serum response factor-binding protein 1 [Drosophi
>gi|1037027264|ref|XP_017139726.1| PREDICTED: 39S ribosomal protein L47, mitochondrial [Drosopl
>gi|1037026957|ref|XP_017139705.1| PREDICTED: lysine-specific demethylase 3B [Drosophila mirand
>gi|1037026975|ref|XP_017139706.1| PREDICTED: lysine-specific demethylase 3B [Drosophila mirand
>gi|1037026866|ref|XP_017139700.1| PREDICTED: F-BAR domain only protein 2 isoform X1 [Drosophil
>gi|1037026881|ref|XP_017139701.1| PREDICTED: F-BAR domain only protein 2 isoform X1 [Drosophil
>gi|1037026899|ref|XP_017139702.1| PREDICTED: F-BAR domain only protein 2 isoform X1 [Drosophil
>gi|1037026915|ref|XP_017139703.1| PREDICTED: F-BAR domain only protein 2 isoform X2 [Drosophil
>gi|1037027010|ref|XP_017139709.1| PREDICTED: tensin [Drosophila miranda]
>gi|1037026993|ref|XP_017139708.1| PREDICTED: solute carrier family 22 member 3 [Drosophila mi
>gi|1037027092|ref|XP_017139714.1| PREDICTED: RNA polymerase II-associated factor 1 homolog [D
>gi|1037027112|ref|XP_017139716.1| PREDICTED: zinc finger protein 628 [Drosophila miranda]
>gi|1037026933|ref|XP_017139704.1| PREDICTED: coiled-coil domain-containing protein 39 [Drosopl
>gi|1037026813|ref|XP_017139697.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X5 [D
>gi|1037026796|ref|XP_017139696.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X4 [D
>gi|1037026776|ref|XP_017139695.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X3 [D
>gi|1037026758|ref|XP_017139693.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X2 [D
>gi|1037026724|ref|XP_017139691.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X1 [D
>gi|1037026740|ref|XP_017139692.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X1 [D
>gi|1037027249|ref|XP_017139725.1| PREDICTED: checkpoint protein HUS1 [Drosophila miranda]
>gi|1037027195|ref|XP_017139721.1| PREDICTED: mannose-1-phosphate guanyltransferase beta [Dros
>gi|1037027180|ref|XP_017139720.1| PREDICTED: Bardet-Biedl syndrome 5 protein homolog [Drosophi
>gi|1037027130|ref|XP_017139717.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1037027317|ref|XP_017139729.1| PREDICTED: uncharacterized protein LOC108154115 [Drosophila
>gi|1037027335|ref|XP_017139730.1| PREDICTED: uncharacterized protein LOC108154116 [Drosophila
>gi|1037027057|ref|XP_017139712.1| PREDICTED: pair-rule protein odd-paired [Drosophila miranda]
>gi|1037105559|ref|XP_017145375.1| PREDICTED: uncharacterized protein LOC108157716 [Drosophila
>gi|1037040931|ref|XP_017140554.1| PREDICTED: LOW QUALITY PROTEIN: replication factor C subunit
>gi|1037040873|ref|XP_017140550.1| PREDICTED: FERM, RhoGEF and pleckstrin domain-containing pro
>gi|1037040887|ref|XP_017140551.1| PREDICTED: FERM, RhoGEF and pleckstrin domain-containing pro
>gi|1037040903|ref|XP_017140552.1| PREDICTED: microtubule-associated protein futsch isoform X3
>gi|1037040917|ref|XP_017140553.1| PREDICTED: ubiquitin-conjugating enzyme E2-17 kDa [Drosophil
>gi|1037023229|ref|XP_017139472.1| PREDICTED: uncharacterized protein LOC108153883, partial [Dr
>gi|1037023247|ref|XP_017139473.1| PREDICTED: lachesin, partial [Drosophila miranda]
>gi|1037105292|ref|XP_017145336.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037106432|ref|XP_017145496.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037105268|ref|XP_017145332.1| PREDICTED: uncharacterized protein LOC108157678 [Drosophila
>gi|1037106594|ref|XP_017145518.1| PREDICTED: uncharacterized protein LOC108157828 [Drosophila
>gi|1037104929|ref|XP_017145284.1| PREDICTED: uncharacterized protein LOC108157646 [Drosophila
>gi|1037104561|ref|XP_017145231.1| PREDICTED: mpv17-like protein [Drosophila miranda]
>gi|1037104576|ref|XP_017145233.1| PREDICTED: mpv17-like protein [Drosophila miranda]
>gi|1037081808|ref|XP_017143344.1| PREDICTED: centrosomal and chromosomal factor [Drosophila m
>gi|1037081819|ref|XP_017143345.1| PREDICTED: uncharacterized protein LOC108156411 [Drosophila
>gi|1037106711|ref|XP_017145534.1| PREDICTED: 4-coumarate--CoA ligase 1-like [Drosophila mirand
>gi|1037102251|ref|XP_017144958.1| PREDICTED: LOW QUALITY PROTEIN: luciferin 4-monooxygenase-li
>gi|1037102261|ref|XP_017144960.1| PREDICTED: luciferin 4-monooxygenase-like [Drosophila mirand
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>gi|1037073267|ref|XP_017142741.1| PREDICTED: LOW QUALITY PROTEIN: geranylgeranyl transferase t
>gi|1037073251|ref|XP_017142740.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037103413|ref|XP_017145079.1| PREDICTED: uncharacterized protein LOC108157494 isoform X1
>gi|1037103432|ref|XP_017145081.1| PREDICTED: uncharacterized protein LOC108157495 isoform X1
>gi|1037103444|ref|XP_017145082.1| PREDICTED: uncharacterized protein LOC108157495 isoform X2
>gi|1037103422|ref|XP_017145080.1| PREDICTED: uncharacterized protein LOC108157494 isoform X2
>gi|1037058322|ref|XP_017141724.1| PREDICTED: afadin isoform X2 [Drosophila miranda]
>gi|1037058308|ref|XP_017141722.1| PREDICTED: afadin isoform X1 [Drosophila miranda]
>gi|1037058338|ref|XP_017141725.1| PREDICTED: uncharacterized protein LOC108155427 isoform X3
>gi|1037101251|ref|XP_017144861.1| PREDICTED: RAB6A-GEF complex partner protein 2 [Drosophila m
>gi|1037054219|ref|XP_017141446.1| PREDICTED: vacuolar protein sorting-associated protein 37B
>gi|1037054170|ref|XP_017141443.1| PREDICTED: katanin p60 ATPase-containing subunit A-like 1 is
>gi|1037054188|ref|XP_017141444.1| PREDICTED: katanin p60 ATPase-containing subunit A-like 1 is
>gi|1037054123|ref|XP_017141439.1| PREDICTED: MMS19 nucleotide excision repair protein homolog
>gi|1037054203|ref|XP_017141445.1| PREDICTED: MORN repeat-containing protein 4 [Drosophila mira
>gi|1037054154|ref|XP_017141441.1| PREDICTED: putative cysteine proteinase CG12163 isoform X2
>gi|1037054139|ref|XP_017141440.1| PREDICTED: putative cysteine proteinase CG12163 isoform X1
>gi|1037054235|ref|XP_017141447.1| PREDICTED: uncharacterized protein LOC108155260 [Drosophila
>gi|1037100842|ref|XP_017144824.1| PREDICTED: uncharacterized protein LOC108157320 [Drosophila
>gi|1037100850|ref|XP_017144825.1| PREDICTED: uncharacterized protein LOC108157320 [Drosophila
>gi|1037100860|ref|XP_017144826.1| PREDICTED: enolase-phosphatase E1 [Drosophila miranda]
>gi|1037046300|ref|XP_017140908.1| PREDICTED: egl nine homolog 1 isoform X1 [Drosophila miranda
>gi|1037046330|ref|XP_017140910.1| PREDICTED: probable mitochondrial import inner membrane tra
>gi|1037046314|ref|XP_017140909.1| PREDICTED: egl nine homolog 1 isoform X2 [Drosophila miranda
>gi|1037046098|ref|XP_017140892.1| PREDICTED: oxidation resistance protein 1 isoform X9 [Drosop
>gi|1037046082|ref|XP_017140891.1| PREDICTED: oxidation resistance protein 1 isoform X8 [Drosop
>gi|1037046244|ref|XP_017140904.1| PREDICTED: oxidation resistance protein 1 isoform X20 [Dros
>gi|1037046064|ref|XP_017140890.1| PREDICTED: oxidation resistance protein 1 isoform X7 [Drosop
>gi|1037046271|ref|XP_017140906.1| PREDICTED: oxidation resistance protein 1 isoform X22 [Dros
>gi|1037046046|ref|XP_017140889.1| PREDICTED: oxidation resistance protein 1 isoform X6 [Drosop
>gi|1037046258|ref|XP_017140905.1| PREDICTED: oxidation resistance protein 1 isoform X21 [Dros
>gi|1037046003|ref|XP_017140886.1| PREDICTED: oxidation resistance protein 1 isoform X3 [Drosop
>gi|1037046033|ref|XP_017140888.1| PREDICTED: oxidation resistance protein 1 isoform X5 [Drosop
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>gi|1037046154|ref|XP_017140896.1| PREDICTED: oxidation resistance protein 1 isoform X13 [Dros
>gi|1037046128|ref|XP_017140894.1| PREDICTED: oxidation resistance protein 1 isoform X11 [Dros
>gi|1037046142|ref|XP_017140895.1| PREDICTED: oxidation resistance protein 1 isoform X12 [Dros
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>gi|1037046286|ref|XP_017140907.1| PREDICTED: oxidation resistance protein 1 isoform X23 [Dros
>gi|1037046164|ref|XP_017140897.1| PREDICTED: oxidation resistance protein 1 isoform X14 [Dros
>gi|1037046204|ref|XP_017140901.1| PREDICTED: TLD domain-containing protein 2 isoform X17 [Dros
>gi|1037046190|ref|XP_017140900.1| PREDICTED: TLD domain-containing protein 2 isoform X16 [Dros
>gi|1037046218|ref|XP_017140902.1| PREDICTED: TLD domain-containing protein 2 isoform X18 [Dros
>gi|1037046347|ref|XP_017140911.1| PREDICTED: probable mitochondrial import inner membrane tra
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>gi|1037046363|ref|XP_017140912.1| PREDICTED: probable mitochondrial import inner membrane tra
>gi|1037089816|ref|XP_017143922.1| PREDICTED: stromal cell-derived factor 2 [Drosophila mirand
>gi|1037089802|ref|XP_017143921.1| PREDICTED: uncharacterized protein CG1161 [Drosophila miran
>gi|1037089788|ref|XP_017143920.1| PREDICTED: proteasome subunit beta type-4 [Drosophila miran
>gi|1037096204|ref|XP_017144404.1| PREDICTED: ceramide kinase [Drosophila miranda]
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>gi|1037076786|ref|XP_017142980.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu
>gi|1037076726|ref|XP_017142976.1| PREDICTED: protein downstream neighbor of son homolog [Dros
>gi|1037076757|ref|XP_017142978.1| PREDICTED: zinc finger protein 17 [Drosophila miranda]
>gi|1037076770|ref|XP_017142979.1| PREDICTED: neuroendocrine protein 7B2 [Drosophila miranda]
>gi|1037076708|ref|XP_017142975.1| PREDICTED: uncharacterized protein LOC108156158 isoform X2
>gi|1037076662|ref|XP_017142971.1| PREDICTED: uncharacterized protein LOC108156158 isoform X1
>gi|1037076678|ref|XP_017142972.1| PREDICTED: uncharacterized protein LOC108156158 isoform X1
>gi|1037076694|ref|XP_017142974.1| PREDICTED: uncharacterized protein LOC108156158 isoform X1
>gi|1037076742|ref|XP_017142977.1| PREDICTED: uncharacterized protein LOC108156160 [Drosophila
>gi|1037027607|ref|XP_017139746.1| PREDICTED: homeobox protein 5 [Drosophila miranda]
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>gi|1037027817|ref|XP_017139761.1| PREDICTED: derlin-2 [Drosophila miranda]
>gi|1037027961|ref|XP_017139768.1| PREDICTED: succinate dehydrogenase assembly factor 3, mitoc
>gi|1037027943|ref|XP_017139767.1| PREDICTED: dihydrofolate reductase [Drosophila miranda]
>gi|1037027575|ref|XP_017139744.1| PREDICTED: soluble guanylate cyclase 89Db [Drosophila miran
>gi|1037027590|ref|XP_017139745.1| PREDICTED: soluble guanylate cyclase 89Db [Drosophila miran
>gi|1037027558|ref|XP_017139743.1| PREDICTED: soluble guanylate cyclase 89Da [Drosophila miran
>gi|1037027800|ref|XP_017139759.1| PREDICTED: COP9 signalosome complex subunit 5 [Drosophila m
>gi|1037027381|ref|XP_017139732.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1037027403|ref|XP_017139733.1| PREDICTED: microtubule-associated protein futsch isoform X2
>gi|1037027755|ref|XP_017139756.1| PREDICTED: glycerophosphodiester phosphodiesterase 1 [Droso
>gi|1037027769|ref|XP_017139757.1| PREDICTED: glycerophosphodiester phosphodiesterase 1 [Droso
>gi|1037027785|ref|XP_017139758.1| PREDICTED: glycerophosphodiester phosphodiesterase 1 [Droso
>gi|1037027835|ref|XP_017139762.1| PREDICTED: trypsin-4 [Drosophila miranda]
>gi|1037027421|ref|XP_017139734.1| PREDICTED: alpha-mannosidase 2x isoform X1 [Drosophila mira
>gi|1037027438|ref|XP_017139735.1| PREDICTED: alpha-mannosidase 2x isoform X2 [Drosophila mira
>gi|1037027456|ref|XP_017139736.1| PREDICTED: alpha-mannosidase 2x isoform X2 [Drosophila mira
>gi|1037027488|ref|XP_017139738.1| PREDICTED: DNA replication licensing factor REC isoform X1
>gi|1037027706|ref|XP_017139753.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophi
>gi|1037027506|ref|XP_017139740.1| PREDICTED: DNA replication licensing factor REC isoform X2
>gi|1037027524|ref|XP_017139741.1| PREDICTED: DNA replication licensing factor REC isoform X3
>gi|1037027851|ref|XP_017139763.1| PREDICTED: uncharacterized protein LOC108154138 [Drosophila
>gi|1037027641|ref|XP_017139749.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1037027473|ref|XP_017139737.1| PREDICTED: importin-5 [Drosophila miranda]
>gi|1037027658|ref|XP_017139750.1| PREDICTED: protein spartin [Drosophila miranda]
>gi|1037027673|ref|XP_017139751.1| PREDICTED: protein spartin [Drosophila miranda]
>gi|1037027690|ref|XP_017139752.1| PREDICTED: dnaJ homolog subfamily C member 3 [Drosophila mi
>gi|1037027738|ref|XP_017139755.1| PREDICTED: otefin [Drosophila miranda]
>gi|1037027541|ref|XP_017139742.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037027871|ref|XP_017139764.1| PREDICTED: GILT-like protein F37H8.5 isoform X1 [Drosophila
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>gi|1037027885|ref|XP_017139765.1| PREDICTED: GILT-like protein F37H8.5 isoform X2 [Drosophila
>gi|1037027925|ref|XP_017139766.1| PREDICTED: uncharacterized protein LOC108154140 [Drosophila
>gi|1037027721|ref|XP_017139754.1| PREDICTED: zinc finger protein 664 [Drosophila miranda]
>gi|1037096911|ref|XP_017144466.1| PREDICTED: ras-related GTP-binding protein A [Drosophila mi
>gi|1037096922|ref|XP_017144467.1| PREDICTED: beta carbonic anhydrase 1 [Drosophila miranda]
>gi|1037093506|ref|XP_017144194.1| PREDICTED: ubiquitin domain-containing protein 1 [Drosophila
>gi|1037093575|ref|XP_017144195.1| PREDICTED: ubiquitin domain-containing protein 1 [Drosophila
>gi|1037023263|ref|XP_017139475.1| PREDICTED: odorant receptor 83a [Drosophila miranda]
>gi|1037032414|ref|XP_017140035.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1037104858|ref|XP_017145273.1| PREDICTED: odorant receptor coreceptor [Drosophila miranda]
>gi|1037106800|ref|XP_017145547.1| PREDICTED: casein kinase I [Drosophila miranda]
>gi|1037075783|ref|XP_017142915.1| PREDICTED: LOW QUALITY PROTEIN: ras-like protein family memb
>gi|1037075749|ref|XP_017142913.1| PREDICTED: eukaryotic translation initiation factor 2-alpha
>gi|1037075767|ref|XP_017142914.1| PREDICTED: eukaryotic translation initiation factor 2-alpha
>gi|1037075799|ref|XP_017142916.1| PREDICTED: 60S ribosomal protein L35a [Drosophila miranda]
>gi|1037054356|ref|XP_017141457.1| PREDICTED: polymerase delta-interacting protein 2 [Drosophil
>gi|1037054383|ref|XP_017141459.1| PREDICTED: 39S ribosomal protein L44, mitochondrial [Drosoph
>gi|1037054368|ref|XP_017141458.1| PREDICTED: SWI/SNF-related matrix-associated actin-dependent
>gi|1037054340|ref|XP_017141456.1| PREDICTED: histone deacetylase 3 [Drosophila miranda]
>gi|1037054251|ref|XP_017141448.1| PREDICTED: biotin--protein ligase isoform X1 [Drosophila mi
>gi|1037054281|ref|XP_017141449.1| PREDICTED: biotin--protein ligase isoform X2 [Drosophila mi
>gi|1037054293|ref|XP_017141450.1| PREDICTED: biotin--protein ligase isoform X3 [Drosophila mi
>gi|1037054313|ref|XP_017141452.1| PREDICTED: biotin--protein ligase isoform X5 [Drosophila mi
>gi|1037054305|ref|XP_017141451.1| PREDICTED: uncharacterized protein LOC108155261 isoform X4
>gi|1037054325|ref|XP_017141453.1| PREDICTED: kinase suppressor of Ras 1 [Drosophila miranda]
>gi|1037083755|ref|XP_017143489.1| PREDICTED: SURP and G-patch domain-containing protein 1 iso
>gi|1037083769|ref|XP_017143490.1| PREDICTED: SURP and G-patch domain-containing protein 1 iso
>gi|1037083783|ref|XP_017143491.1| PREDICTED: formin-like protein 20 isoform X3 [Drosophila mi
>gi|1037083727|ref|XP_017143486.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1037083741|ref|XP_017143488.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1037059827|ref|XP_017141830.1| PREDICTED: glutamate [NMDA] receptor subunit 1 [Drosophila m
>gi|1037059860|ref|XP_017141832.1| PREDICTED: uncharacterized protein LOC108155504 isoform X2
>gi|1037059842|ref|XP_017141831.1| PREDICTED: uncharacterized protein LOC108155504 isoform X1
>gi|1037059751|ref|XP_017141824.1| PREDICTED: inositol 1,4,5-trisphosphate receptor isoform X1
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>gi|1037059779|ref|XP_017141826.1| PREDICTED: inositol 1,4,5-trisphosphate receptor isoform X2
>gi|1037059811|ref|XP_017141829.1| PREDICTED: uncharacterized protein LOC108155502 isoform X2
>gi|1037059795|ref|XP_017141827.1| PREDICTED: uncharacterized protein LOC108155502 isoform X1
>gi|1037071659|ref|XP_017142629.1| PREDICTED: protein-L-isoaspartate(D-aspartate) O-methyltrans
>gi|1037071581|ref|XP_017142623.1| PREDICTED: DNA cross-link repair 1A protein [Drosophila mira
>gi|1037071594|ref|XP_017142624.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037071607|ref|XP_017142625.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037071632|ref|XP_017142627.1| PREDICTED: phosphatidylinositol 4-kinase type 2-alpha isofor
>gi|1037071646|ref|XP_017142628.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037071620|ref|XP_017142626.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037089775|ref|XP_017143918.1| PREDICTED: uncharacterized protein LOC108156765 [Drosophila
>gi|1037089731|ref|XP_017143915.1| PREDICTED: phospholipid phosphatase 5 isoform X1 [Drosophila
>gi|1037089747|ref|XP_017143916.1| PREDICTED: phospholipid phosphatase 5 isoform X2 [Drosophila
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>gi|1037089761|ref|XP_017143917.1| PREDICTED: RNA-binding protein 42 [Drosophila miranda]
>gi|1037101065|ref|XP_017144846.1| PREDICTED: GTP-binding protein Rheb homolog [Drosophila mira
>gi|1037098227|ref|XP_017144594.1| PREDICTED: succinate-semialdehyde dehydrogenase [NADP(+)] Ga
>gi|1037049406|ref|XP_017141116.1| PREDICTED: nuclear protein localization protein 4 homolog is
>gi|1037049422|ref|XP_017141117.1| PREDICTED: nuclear protein localization protein 4 homolog is
>gi|1037049388|ref|XP_017141115.1| PREDICTED: nuclear protein localization protein 4 homolog is
>gi|1037049440|ref|XP_017141118.1| PREDICTED: uncharacterized protein LOC108155052 isoform X1
>gi|1037049472|ref|XP_017141121.1| PREDICTED: uncharacterized protein LOC108155052 isoform X3
>gi|1037049456|ref|XP_017141119.1| PREDICTED: uncharacterized protein LOC108155052 isoform X2
>gi|1037049632|ref|XP_017141131.1| PREDICTED: putative uncharacterized protein DDB_G0274435 [D:
>gi|1037049614|ref|XP_017141130.1| PREDICTED: uncharacterized protein LOC108155056 isoform X2
>gi|1037049582|ref|XP_017141128.1| PREDICTED: uncharacterized protein LOC108155056 isoform X1
>gi|1037049598|ref|XP_017141129.1| PREDICTED: uncharacterized protein LOC108155056 isoform X1
>gi|1037049534|ref|XP_017141125.1| PREDICTED: protein gar2 isoform X3 [Drosophila miranda]
>gi|1037049520|ref|XP_017141124.1| PREDICTED: CUGBP Elav-like family member 2 isoform X2 [Droso
>gi|1037049488|ref|XP_017141122.1| PREDICTED: CUGBP Elav-like family member 2 isoform X1 [Droso
>gi|1037049504|ref|XP_017141123.1| PREDICTED: CUGBP Elav-like family member 2 isoform X1 [Droso
>gi|1037049550|ref|XP_017141126.1| PREDICTED: lipoprotein lipase [Drosophila miranda]
>gi|1037049566|ref|XP_017141127.1| PREDICTED: uncharacterized protein LOC108155055 [Drosophila
>gi|1037104163|ref|XP_017145175.1| PREDICTED: protein regulator of cytokinesis 1-like isoform ]
>gi|1037104171|ref|XP_017145176.1| PREDICTED: protein regulator of cytokinesis 1-like isoform ]
>gi|1037104176|ref|XP_017145177.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1037036570|ref|XP_017140276.1| PREDICTED: uncharacterized protein LOC108154486 [Drosophila
>gi|1037100542|ref|XP_017144792.1| PREDICTED: fatty-acid amide hydrolase 2 [Drosophila miranda]
>gi|1037079511|ref|XP_017143164.1| PREDICTED: dihydroxyacetone phosphate acyltransferase [Droso
>gi|1037079524|ref|XP_017143165.1| PREDICTED: putative GTP-binding protein 6 [Drosophila miranc
>gi|1037079538|ref|XP_017143166.1| PREDICTED: transmembrane protein 141 [Drosophila miranda]
>gi|1037075515|ref|XP_017142896.1| PREDICTED: zinc finger protein 436 [Drosophila miranda]
>gi|1037075498|ref|XP_017142895.1| PREDICTED: MAU2 chromatid cohesion factor homolog [Drosophil
>gi|1037075531|ref|XP_017142897.1| PREDICTED: serpin B3 [Drosophila miranda]
>gi|1037075545|ref|XP_017142898.1| PREDICTED: serpin B3 [Drosophila miranda]
>gi|1037079498|ref|XP_017143163.1| PREDICTED: transmembrane protein 165 isoform X2 [Drosophila
>gi|1037079470|ref|XP_017143161.1| PREDICTED: transmembrane protein 165 isoform X1 [Drosophila
>gi|1037079482|ref|XP_017143162.1| PREDICTED: transmembrane protein 165 isoform X1 [Drosophila
>gi|1037079456|ref|XP_017143160.1| PREDICTED: E3 ubiquitin-protein ligase RNF123 [Drosophila m:
>gi|1037032776|ref|XP_017140058.1| PREDICTED: soluble guanylate cyclase 88E [Drosophila miranda
>gi|1037099815|ref|XP_017144735.1| PREDICTED: uncharacterized protein LOC108157264 isoform X2
>gi|1037099839|ref|XP_017144737.1| PREDICTED: uncharacterized protein LOC108157264 isoform X4
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>gi|1037099827|ref|XP_017144736.1| PREDICTED: uncharacterized protein LOC108157264 isoform X3
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>gi|1037099872|ref|XP_017144741.1| PREDICTED: uncharacterized protein LOC108157264 isoform X7
>gi|1037099882|ref|XP_017144742.1| PREDICTED: uncharacterized protein LOC108157264 isoform X8
>gi|1037088541|ref|XP_017143821.1| PREDICTED: glycogen [starch] synthase isoform X1 [Drosophila
>gi|1037088555|ref|XP_017143822.1| PREDICTED: glycogen [starch] synthase isoform X2 [Drosophila
>gi|1037088569|ref|XP_017143823.1| PREDICTED: uncharacterized protein LOC108156690 [Drosophila
>gi|1037105433|ref|XP_017145357.1| PREDICTED: vegetative cell wall protein gp1 [Drosophila mira
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>gi|1037105438|ref|XP_017145359.1| PREDICTED: uncharacterized protein LOC108157700 [Drosophila
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>gi|1037092078|ref|XP_017144096.1| PREDICTED: uncharacterized protein LOC108156877 [Drosophila
>gi|1037089359|ref|XP_017143885.1| PREDICTED: protein fem-1 homolog CG6966 [Drosophila miranda]
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>gi|1037104354|ref|XP_017145202.1| PREDICTED: uncharacterized protein LOC108157589 isoform X1
>gi|1037104362|ref|XP_017145203.1| PREDICTED: uncharacterized protein LOC108157589 isoform X1
>gi|1037104372|ref|XP_017145205.1| PREDICTED: uncharacterized protein LOC108157589 isoform X2
>gi|1037037786|ref|XP_017140347.1| PREDICTED: uncharacterized protein LOC108154555 [Drosophila
>gi|1037106727|ref|XP_017145536.1| PREDICTED: uncharacterized protein LOC108157843 isoform X2
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>gi|1037104554|ref|XP_017145230.1| PREDICTED: uncharacterized protein LOC108157612 [Drosophila
>gi|1037104547|ref|XP_017145229.1| PREDICTED: general transcription factor IIF subunit 1-like
>gi|1037092373|ref|XP_017144114.1| PREDICTED: insulin-like peptide receptor [Drosophila miranda
>gi|1037037208|ref|XP_017140312.1| PREDICTED: luciferin 4-monooxygenase-like [Drosophila mirand
>gi|1037023299|ref|XP_017139477.1| PREDICTED: succinate-semialdehyde dehydrogenase [NADP(+)] Ga
>gi|1037072161|ref|XP_017142662.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit
>gi|1037072177|ref|XP_017142663.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit
>gi|1037072116|ref|XP_017142658.1| PREDICTED: 40S ribosomal protein S5b [Drosophila miranda]
>gi|1037072131|ref|XP_017142660.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit
>gi|1037072147|ref|XP_017142661.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit
>gi|1037072068|ref|XP_017142655.1| PREDICTED: RRP15-like protein [Drosophila miranda]
>gi|1037072100|ref|XP_017142657.1| PREDICTED: UPF0568 protein C14orf166 homolog [Drosophila mi
>gi|1037072191|ref|XP_017142664.1| PREDICTED: uncharacterized protein LOC108155984 [Drosophila
>gi|1037072052|ref|XP_017142654.1| PREDICTED: nudC domain-containing protein 3 [Drosophila mira
>gi|1037072084|ref|XP_017142656.1| PREDICTED: nurim homolog [Drosophila miranda]
>gi|1037090445|ref|XP_017143974.1| PREDICTED: eukaryotic translation initiation factor 1A, X-ch
>gi|1037090459|ref|XP_017143975.1| PREDICTED: eukaryotic translation initiation factor 1A, X-ch
>gi|1037090473|ref|XP_017143977.1| PREDICTED: eukaryotic translation initiation factor 1A, X-ch
>gi|1037090486|ref|XP_017143978.1| PREDICTED: eukaryotic translation initiation factor 1A, X-ch
>gi|1037090431|ref|XP_017143973.1| PREDICTED: transmembrane protease serine 9 [Drosophila mira
>gi|1037090500|ref|XP_017143979.1| PREDICTED: uncharacterized protein LOC108156803 [Drosophila
>gi|1037087703|ref|XP_017143756.1| PREDICTED: WD repeat-containing protein 3 [Drosophila mirano
>gi|1037087717|ref|XP_017143757.1| PREDICTED: peptidyl-prolyl cis-trans isomerase FKBP2 [Drosop
>gi|1037068919|ref|XP_017142440.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037068935|ref|XP_017142441.1| PREDICTED: putative elongator complex protein 1 [Drosophila
>gi|1037064004|ref|XP_017142117.1| PREDICTED: uncharacterized protein LOC108155665 [Drosophila
>gi|1037063989|ref|XP_017142116.1| PREDICTED: nuclear pore complex protein Nup214 isoform X2 [I
>gi|1037063961|ref|XP_017142113.1| PREDICTED: nuclear pore complex protein Nup214 isoform X1 [I
>gi|1037063974|ref|XP_017142114.1| PREDICTED: nuclear pore complex protein Nup214 isoform X1 [I
>gi|1037040551|ref|XP_017140528.1| PREDICTED: zinc finger protein 10 [Drosophila miranda]
>gi|1037040624|ref|XP_017140533.1| PREDICTED: gastrula zinc finger protein XlCGF17.1 isoform X1
>gi|1037040638|ref|XP_017140534.1| PREDICTED: zinc finger protein 211 isoform X2 [Drosophila mi
>gi|1037040539|ref|XP_017140526.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037040597|ref|XP_017140531.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037040565|ref|XP_017140529.1| PREDICTED: zinc finger protein 20 [Drosophila miranda]
>gi|1037040382|ref|XP_017140514.1| PREDICTED: zinc finger protein 229 [Drosophila miranda]
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>gi|1037040654|ref|XP_017140535.1| PREDICTED: zinc finger protein 648-like [Drosophila miranda]
>gi|1037040446|ref|XP_017140520.1| PREDICTED: transcription factor Ouib-like isoform X5 [Drosop
>gi|1037040430|ref|XP_017140519.1| PREDICTED: transcription factor Ouib-like isoform X4 [Drosop
>gi|1037040470|ref|XP_017140521.1| PREDICTED: transcription factor Ouib-like isoform X6 [Drosop
>gi|1037040485|ref|XP_017140522.1| PREDICTED: transcription factor Ouib-like isoform X6 [Drosop
>gi|1037040497|ref|XP_017140523.1| PREDICTED: transcription factor Ouib-like isoform X7 [Drosop
>gi|1037040400|ref|XP_017140515.1| PREDICTED: transcription factor Ouib-like isoform X1 [Drosop
>gi|1037040420|ref|XP_017140518.1| PREDICTED: transcription factor Ouib-like isoform X3 [Drosop
>gi|1037040410|ref|XP_017140516.1| PREDICTED: transcription factor Ouib-like isoform X2 [Drosop
>gi|1037040581|ref|XP_017140530.1| PREDICTED: transcription factor Ouib-like [Drosophila miranc
>gi|1037040666|ref|XP_017140536.1| PREDICTED: protein RER1 [Drosophila miranda]
>gi|1037040608|ref|XP_017140532.1| PREDICTED: 3-ketodihydrosphingosine reductase [Drosophila m:
>gi|1037040680|ref|XP_017140537.1| PREDICTED: peptidyl-prolyl cis-trans isomerase NIMA-interact
>gi|1037040346|ref|XP_017140512.1| PREDICTED: probable glutamine--tRNA ligase [Drosophila mirar
>gi|1037040362|ref|XP_017140513.1| PREDICTED: serine/threonine-protein kinase RIO2 [Drosophila
>gi|1037040527|ref|XP_017140525.1| PREDICTED: uncharacterized protein LOC108154688 [Drosophila
>gi|1037040511|ref|XP_017140524.1| PREDICTED: uncharacterized protein LOC108154687 [Drosophila
>gi|1037104456|ref|XP_017145216.1| PREDICTED: uncharacterized protein LOC108157599 [Drosophila
>gi|1037102846|ref|XP_017145017.1| PREDICTED: uncharacterized protein LOC108157456 [Drosophila
>gi|1037102855|ref|XP_017145018.1| PREDICTED: uncharacterized protein LOC108157457 [Drosophila
>gi|1037092862|ref|XP_017144147.1| PREDICTED: uncharacterized protein LOC108156909 [Drosophila
>gi|1037092892|ref|XP_017144149.1| PREDICTED: uncharacterized protein LOC108156912 [Drosophila
>gi|1037092878|ref|XP_017144148.1| PREDICTED: uncharacterized protein LOC108156910 [Drosophila
>gi|1037106153|ref|XP_017145457.1| PREDICTED: uncharacterized protein LOC108157778 [Drosophila
>gi|1037106205|ref|XP_017145465.1| PREDICTED: uncharacterized protein LOC108157786 [Drosophila
>gi|1037105460|ref|XP_017145362.1| PREDICTED: uncharacterized protein LOC108157702 [Drosophila
>gi|1037032813|ref|XP_017140061.1| PREDICTED: uncharacterized protein LOC108154314 isoform X1
>gi|1037032831|ref|XP_017140062.1| PREDICTED: uncharacterized protein LOC108154314 isoform X2
>gi|1037104324|ref|XP_017145198.1| PREDICTED: uncharacterized protein LOC108157585 [Drosophila
>gi|1037104332|ref|XP_017145199.1| PREDICTED: uncharacterized protein LOC108157585 [Drosophila
>gi|1037033108|ref|XP_017140079.1| PREDICTED: uncharacterized protein LOC108154328 [Drosophila
>gi|1037106774|ref|XP_017145543.1| PREDICTED: uncharacterized protein LOC108157850 [Drosophila
>gi|1037106087|ref|XP_017145448.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037106095|ref|XP_017145449.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037106103|ref|XP_017145451.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037106111|ref|XP_017145452.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037023315|ref|XP_017139478.1| PREDICTED: uncharacterized protein LOC108153890 [Drosophila
>gi|1037032314|ref|XP_017140029.1| PREDICTED: uncharacterized protein LOC108154289 [Drosophila
>gi|1037032484|ref|XP_017140040.1| PREDICTED: uncharacterized protein LOC108154298 [Drosophila
>gi|1037093392|ref|XP_017144185.1| PREDICTED: uncharacterized protein LOC108156935 [Drosophila
>gi|1037093406|ref|XP_017144186.1| PREDICTED: uncharacterized protein LOC108156936 [Drosophila
>gi|1037093379|ref|XP_017144183.1| PREDICTED: zinc finger protein 62 homolog [Drosophila mirano
>gi|1037086961|ref|XP_017143730.1| PREDICTED: zinc finger protein 91 [Drosophila miranda]
>gi|1037079058|ref|XP_017143144.1| PREDICTED: TATA-binding protein-associated factor 172 [Droso
>gi|1037058017|ref|XP_017141701.1| PREDICTED: SWI/SNF complex subunit SMARCC1 [Drosophila miran
>gi|1037058031|ref|XP_017141702.1| PREDICTED: retinoblastoma family protein [Drosophila miranda
>gi|1037058106|ref|XP_017141708.1| PREDICTED: uncharacterized protein LOC108155419 [Drosophila
>gi|1037058077|ref|XP_017141707.1| PREDICTED: zinc finger CCHC domain-containing protein 10 [Di
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>gi|1037058047|ref|XP_017141703.1| PREDICTED: uncharacterized protein LOC108155417 [Drosophila
>gi|1037058063|ref|XP_017141704.1| PREDICTED: uncharacterized protein LOC108155417 [Drosophila
>gi|1037061293|ref|XP_017141932.1| PREDICTED: synapse-associated protein of 47 kDa isoform X9
>gi|1037061279|ref|XP_017141931.1| PREDICTED: synapse-associated protein of 47 kDa isoform X8
>gi|1037061264|ref|XP_017141930.1| PREDICTED: synapse-associated protein of 47 kDa isoform X7
>gi|1037061186|ref|XP_017141924.1| PREDICTED: synapse-associated protein of 47 kDa isoform X2
>gi|1037061202|ref|XP_017141925.1| PREDICTED: synapse-associated protein of 47 kDa isoform X3
>gi|1037061216|ref|XP_017141926.1| PREDICTED: synapse-associated protein of 47 kDa isoform X4
>gi|1037061155|ref|XP_017141922.1| PREDICTED: synapse-associated protein of 47 kDa isoform X1
>gi|1037061170|ref|XP_017141923.1| PREDICTED: synapse-associated protein of 47 kDa isoform X1
>gi|1037061309|ref|XP_017141933.1| PREDICTED: synapse-associated protein of 47 kDa isoform X10
>gi|1037061248|ref|XP_017141929.1| PREDICTED: synapse-associated protein of 47 kDa isoform X6
>gi|1037061232|ref|XP_017141928.1| PREDICTED: synapse-associated protein of 47 kDa isoform X5
>gi|1037061325|ref|XP_017141934.1| PREDICTED: synapse-associated protein of 47 kDa isoform X11
>gi|1037061371|ref|XP_017141935.1| PREDICTED: synapse-associated protein of 47 kDa isoform X12
>gi|1037061414|ref|XP_017141937.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037061398|ref|XP_017141936.1| PREDICTED: uncharacterized protein LOC108155556 [Drosophila
>gi|1037061137|ref|XP_017141921.1| PREDICTED: F-box/LRR-repeat protein 7 [Drosophila miranda]
>gi|1037061430|ref|XP_017141938.1| PREDICTED: PI-actitoxin-Afv2a [Drosophila miranda]
>gi|1037095049|ref|XP_017144307.1| PREDICTED: uncharacterized protein LOC108157013 isoform X3
>gi|1037095033|ref|XP_017144306.1| PREDICTED: serologically defined colon cancer antigen 8 hom
>gi|1037095019|ref|XP_017144305.1| PREDICTED: serologically defined colon cancer antigen 8 hom
>gi|1037095063|ref|XP_017144308.1| PREDICTED: poly(U)-specific endoribonuclease homolog [Droso
>gi|1037097697|ref|XP_017144540.1| PREDICTED: globin CTT-VI [Drosophila miranda]
>gi|1037097711|ref|XP_017144541.1| PREDICTED: globin CTT-VI [Drosophila miranda]
>gi|1037097836|ref|XP_017144554.1| PREDICTED: atrial natriuretic peptide receptor 3 [Drosophila
>gi|1037097823|ref|XP_017144553.1| PREDICTED: atrial natriuretic peptide receptor 1 [Drosophila
>gi|1037097846|ref|XP_017144555.1| PREDICTED: uncharacterized protein LOC108157157 [Drosophila
>gi|1037052769|ref|XP_017141348.1| PREDICTED: ATP-binding cassette sub-family B member 6, mito
>gi|1037052797|ref|XP_017141349.1| PREDICTED: ATP-binding cassette sub-family B member 6, mito
>gi|1037052737|ref|XP_017141346.1| PREDICTED: uncharacterized protein LOC108155186 [Drosophila
>gi|1037052813|ref|XP_017141350.1| PREDICTED: arginine kinase [Drosophila miranda]
>gi|1037052753|ref|XP_017141347.1| PREDICTED: probable ATP-dependent RNA helicase spindle-E [D
>gi|1037052827|ref|XP_017141351.1| PREDICTED: uncharacterized protein LOC108155191 [Drosophila
>gi|1037069046|ref|XP_017142449.1| PREDICTED: NADH-quinone oxidoreductase subunit I [Drosophila
>gi|1037068998|ref|XP_017142446.1| PREDICTED: structure-specific endonuclease subunit SLX1 hom
>gi|1037069012|ref|XP_017142447.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1037069028|ref|XP_017142448.1| PREDICTED: actin-related protein 2/3 complex subunit 3 [Dro
>gi|1037068966|ref|XP_017142443.1| PREDICTED: uncharacterized protein LOC108155844 isoform X1
>gi|1037068982|ref|XP_017142445.1| PREDICTED: uncharacterized protein LOC108155844 isoform X2
>gi|1037092846|ref|XP_017144146.1| PREDICTED: micronuclear linker histone polyprotein [Drosoph
>gi|1037103905|ref|XP_017145143.1| PREDICTED: 5-hydroxytryptamine receptor 2B [Drosophila mira
>gi|1037103913|ref|XP_017145144.1| PREDICTED: uncharacterized protein LOC108157551 [Drosophila
>gi|1037104673|ref|XP_017145246.1| PREDICTED: uncharacterized protein LOC108157621 [Drosophila
>gi|1037104683|ref|XP_017145247.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1037094620|ref|XP_017144275.1| PREDICTED: pre-mRNA 3'-end-processing factor FIP1 [Drosophil
>gi|1037032619|ref|XP_017140048.1| PREDICTED: nuclear receptor subfamily 2 group E member 1 iso
>gi|1037032656|ref|XP_017140050.1| PREDICTED: nuclear receptor subfamily 2 group E member 1 iso
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>gi|1037032636|ref|XP_017140049.1| PREDICTED: nuclear receptor subfamily 2 group E member 1 iso
>gi|1037088719|ref|XP_017143835.1| PREDICTED: serine/threonine-protein kinase PAK 1 [Drosophila
>gi|1037088737|ref|XP_017143836.1| PREDICTED: serine/threonine-protein kinase PAK 1 [Drosophila
>gi|1037056216|ref|XP_017141578.1| PREDICTED: transcription elongation regulator 1 isoform X2
>gi|1037056203|ref|XP_017141576.1| PREDICTED: transcription elongation regulator 1 isoform X1
>gi|1037056264|ref|XP_017141581.1| PREDICTED: transcription elongation regulator 1 isoform X5
>gi|1037056232|ref|XP_017141579.1| PREDICTED: transcription elongation regulator 1 isoform X3
>gi|1037056248|ref|XP_017141580.1| PREDICTED: transcription elongation regulator 1 isoform X4
>gi|1037056282|ref|XP_017141583.1| PREDICTED: HEAT repeat-containing protein 3 [Drosophila mira
>gi|1037023331|ref|XP_017139479.1| PREDICTED: uncharacterized protein LOC108153891 [Drosophila
>gi|1037056314|ref|XP_017141585.1| PREDICTED: uncharacterized protein LOC108155341 [Drosophila
>gi|1037056298|ref|XP_017141584.1| PREDICTED: general transcription factor IIF subunit 1 [Droso
>gi|1037094390|ref|XP_017144257.1| PREDICTED: E3 ubiquitin-protein ligase RNF13 [Drosophila mir
>gi|1037082465|ref|XP_017143392.1| PREDICTED: transmembrane and coiled-coil domains protein 2
>gi|1037060768|ref|XP_017141896.1| PREDICTED: histone-lysine N-methyltransferase, H3 lysine-79
>gi|1037060786|ref|XP_017141897.1| PREDICTED: histone-lysine N-methyltransferase, H3 lysine-79
>gi|1037089564|ref|XP_017143898.1| PREDICTED: uncharacterized protein LOC108156747 [Drosophila
>gi|1037097097|ref|XP_017144484.1| PREDICTED: zinc finger protein 616 [Drosophila miranda]
>gi|1037103062|ref|XP_017145041.1| PREDICTED: neurochondrin homolog [Drosophila miranda]
>gi|1037102740|ref|XP_017145006.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102780|ref|XP_017145011.1| PREDICTED: uncharacterized protein LOC108157454 isoform X5
>gi|1037102758|ref|XP_017145009.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102802|ref|XP_017145013.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102748|ref|XP_017145008.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102792|ref|XP_017145012.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102768|ref|XP_017145010.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102816|ref|XP_017145014.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102827|ref|XP_017145015.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037036995|ref|XP_017140300.1| PREDICTED: uncharacterized protein LOC108154510 [Drosophila
>gi|1037035889|ref|XP_017140236.1| PREDICTED: uncharacterized protein LOC108154473 isoform X2
>gi|1037035871|ref|XP_017140235.1| PREDICTED: uncharacterized protein LOC108154473 isoform X1
>gi|1037034002|ref|XP_017140131.1| PREDICTED: uncharacterized protein LOC108154378 [Drosophila
>gi|1037100520|ref|XP_017144790.1| PREDICTED: hemicentin-2 [Drosophila miranda]
>gi|1037100500|ref|XP_017144788.1| PREDICTED: probable histone-binding protein Caf1 [Drosophila
>gi|1037100510|ref|XP_017144789.1| PREDICTED: golgin subfamily A member 6-like protein 22 [Dros
>gi|1037100530|ref|XP_017144791.1| PREDICTED: uncharacterized protein LOC108157301 [Drosophila
>gi|1037033766|ref|XP_017140116.1| PREDICTED: uncharacterized protein LOC108154362 [Drosophila
>gi|1037106808|ref|XP_017145549.1| PREDICTED: cytoplasmic dynein 1 light intermediate chain 1-l
>gi|1037035613|ref|XP_017140219.1| PREDICTED: Golgi-associated plant pathogenesis-related prote
>gi|1037095800|ref|XP_017144367.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Droso
>gi|1037095823|ref|XP_017144369.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Droso
>gi|1037095835|ref|XP_017144370.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Droso
>gi|1037095846|ref|XP_017144371.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Droso
>gi|1037095858|ref|XP_017144372.1| PREDICTED: uncharacterized protein LOC108157054 isoform X2
>gi|1037095870|ref|XP_017144373.1| PREDICTED: uncharacterized protein LOC108157055 [Drosophila
>gi|1037095913|ref|XP_017144374.1| PREDICTED: uncharacterized protein LOC108157056 [Drosophila
>gi|1037092386|ref|XP_017144115.1| PREDICTED: argininosuccinate synthase isoform X1 [Drosophila
>gi|1037092400|ref|XP_017144116.1| PREDICTED: argininosuccinate synthase isoform X2 [Drosophila
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>gi|1037092414|ref|XP_017144117.1| PREDICTED: uncharacterized protein LOC108156891 isoform X3
>gi|1037092430|ref|XP_017144119.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037092444|ref|XP_017144120.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037092458|ref|XP_017144121.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037092472|ref|XP_017144122.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037062362|ref|XP_017142003.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062378|ref|XP_017142004.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062392|ref|XP_017142005.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062407|ref|XP_017142006.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062455|ref|XP_017142009.1| PREDICTED: probable peptidyl-tRNA hydrolase 2 [Drosophila mi
>gi|1037062439|ref|XP_017142008.1| PREDICTED: signal peptidase complex catalytic subunit SEC11A
>gi|1037062423|ref|XP_017142007.1| PREDICTED: regulatory protein ada [Drosophila miranda]
>gi|1037102136|ref|XP_017144945.1| PREDICTED: homeotic protein labial [Drosophila miranda]
>gi|1037023347|ref|XP_017139480.1| PREDICTED: pupal cuticle protein Edg-84A [Drosophila miranda
>gi|1037036635|ref|XP_017140280.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037023363|ref|XP_017139481.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037023379|ref|XP_017139483.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037036695|ref|XP_017140285.1| PREDICTED: larval cuticle protein A3A [Drosophila miranda]
>gi|1037037731|ref|XP_017140344.1| PREDICTED: uncharacterized protein LOC108154552 [Drosophila
>gi|1037038864|ref|XP_017140413.1| PREDICTED: cuticle protein 19.8 [Drosophila miranda]
>gi|1037037065|ref|XP_017140305.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037034464|ref|XP_017140161.1| PREDICTED: cuticle protein 7-like [Drosophila miranda]
>gi|1037101693|ref|XP_017144899.1| PREDICTED: homeotic protein proboscipedia [Drosophila mirand
>gi|1037101703|ref|XP_017144900.1| PREDICTED: homeotic protein proboscipedia [Drosophila mirand
>gi|1037030845|ref|XP_017139943.1| PREDICTED: protein zerknuellt 2 [Drosophila miranda]
>gi|1037030871|ref|XP_017139945.1| PREDICTED: tetratricopeptide repeat protein 12 [Drosophila m
>gi|1037030973|ref|XP_017139950.1| PREDICTED: protein zerknuellt [Drosophila miranda]
>gi|1037058367|ref|XP_017141727.1| PREDICTED: homeotic protein bicoid isoform X1 [Drosophila mi
>gi|1037058383|ref|XP_017141728.1| PREDICTED: homeotic protein bicoid isoform X2 [Drosophila mi
>gi|1037058431|ref|XP_017141731.1| PREDICTED: protein amalgam [Drosophila miranda]
>gi|1037058447|ref|XP_017141733.1| PREDICTED: uncharacterized protein LOC108155432 [Drosophila
>gi|1037058354|ref|XP_017141726.1| PREDICTED: homeotic protein deformed [Drosophila miranda]
>gi|1037058399|ref|XP_017141729.1| PREDICTED: homeotic protein Sex combs reduced [Drosophila mi
>gi|1037058415|ref|XP_017141730.1| PREDICTED: homeotic protein Sex combs reduced [Drosophila mi
>gi|1037101222|ref|XP_017144860.1| PREDICTED: segmentation protein fushi tarazu [Drosophila mir
>gi|1037032569|ref|XP_017140045.1| PREDICTED: uncharacterized protein LOC108154302 [Drosophila
>gi|1037066419|ref|XP_017142270.1| PREDICTED: homeotic protein antennapedia isoform X4 [Drosoph
>gi|1037066406|ref|XP_017142269.1| PREDICTED: homeotic protein antennapedia isoform X3 [Drosoph
>gi|1037066392|ref|XP_017142268.1| PREDICTED: homeotic protein antennapedia isoform X2 [Drosoph
>gi|1037066362|ref|XP_017142266.1| PREDICTED: homeotic protein antennapedia isoform X1 [Drosoph
>gi|1037066379|ref|XP_017142267.1| PREDICTED: homeotic protein antennapedia isoform X1 [Drosoph
>gi|1037049041|ref|XP_017141091.1| PREDICTED: sorbitol dehydrogenase [Drosophila miranda]
>gi|1037023395|ref|XP_017139484.1| PREDICTED: putative leucine-rich repeat-containing protein L
>gi|1037049012|ref|XP_017141089.1| PREDICTED: axoneme-associated protein mst101(2) [Drosophila
>gi|1037049028|ref|XP_017141090.1| PREDICTED: arrestin domain-containing protein 2 [Drosophila
>gi|1037048891|ref|XP_017141080.1| PREDICTED: PAX3- and PAX7-binding protein 1 isoform X1 [Dros
>gi|1037048923|ref|XP_017141082.1| PREDICTED: uncharacterized protein LOC108155030 isoform X3
>gi|1037048907|ref|XP_017141081.1| PREDICTED: PAX3- and PAX7-binding protein 1 isoform X2 [Dros
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>gi|1037048941|ref|XP_017141083.1| PREDICTED: E3 UFM1-protein ligase 1 homolog [Drosophila mira
>gi|1037048980|ref|XP_017141087.1| PREDICTED: TBC1 domain family member 22B [Drosophila miranda
>gi|1037048956|ref|XP_017141084.1| PREDICTED: exocyst complex component 6 [Drosophila miranda]
>gi|1037048996|ref|XP_017141088.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1037064384|ref|XP_017142142.1| PREDICTED: ras-related protein Rab-11A [Drosophila miranda]
>gi|1037064350|ref|XP_017142140.1| PREDICTED: protein Peter pan [Drosophila miranda]
>gi|1037064366|ref|XP_017142141.1| PREDICTED: peptidyl-prolyl cis-trans isomerase D [Drosophila
>gi|1037064334|ref|XP_017142138.1| PREDICTED: F-box/WD repeat-containing protein 11 [Drosophila
>gi|1037085749|ref|XP_017143636.1| PREDICTED: acylpyruvase FAHD1, mitochondrial [Drosophila mir
>gi|1037085763|ref|XP_017143637.1| PREDICTED: uncharacterized protein LOC108156606 isoform X1
>gi|1037085777|ref|XP_017143638.1| PREDICTED: uncharacterized protein LOC108156606 isoform X2
>gi|1037085721|ref|XP_017143634.1| PREDICTED: 14-3-3 protein epsilon isoform X1 [Drosophila mir
>gi|1037085735|ref|XP_017143635.1| PREDICTED: 14-3-3 protein epsilon isoform X2 [Drosophila mir
>gi|1037061552|ref|XP_017141947.1| PREDICTED: uncharacterized protein LOC108155565 [Drosophila
>gi|1037061522|ref|XP_017141945.1| PREDICTED: protein AAR2 homolog [Drosophila miranda]
>gi|1037061478|ref|XP_017141942.1| PREDICTED: DNA-directed RNA polymerase II 16 kDa polypeptide
>gi|1037061462|ref|XP_017141941.1| PREDICTED: transcriptional adapter 2A isoform X1 [Drosophila
>gi|1037061542|ref|XP_017141946.1| PREDICTED: uncharacterized protein LOC108155564 [Drosophila
>gi|1037061492|ref|XP_017141943.1| PREDICTED: trimethylguanosine synthase [Drosophila miranda]
>gi|1037061506|ref|XP_017141944.1| PREDICTED: homeobox protein MOX-2 [Drosophila miranda]
>gi|1037061446|ref|XP_017141940.1| PREDICTED: vam6/Vps39-like protein [Drosophila miranda]
>gi|1037090890|ref|XP_017144008.1| PREDICTED: signal peptide peptidase-like 3 isoform X1 [Dros
>gi|1037090903|ref|XP_017144009.1| PREDICTED: signal peptide peptidase-like 3 isoform X2 [Dros
>gi|1037091459|ref|XP_017144051.1| PREDICTED: SH2B adapter protein 1 isoform X1 [Drosophila mi
>gi|1037091473|ref|XP_017144052.1| PREDICTED: SH2B adapter protein 1 isoform X1 [Drosophila mi
>gi|1037091487|ref|XP_017144053.1| PREDICTED: SH2B adapter protein 1 isoform X2 [Drosophila mi
>gi|1037091501|ref|XP_017144054.1| PREDICTED: SH2B adapter protein 1 isoform X3 [Drosophila mi
>gi|1037100115|ref|XP_017144753.1| PREDICTED: protein FAM98B [Drosophila miranda]
>gi|1037023411|ref|XP_017139485.1| PREDICTED: superoxide dismutase [Cu-Zn], chloroplastic [Dros
>gi|1037037083|ref|XP_017140306.1| PREDICTED: laccase [Drosophila miranda]
>gi|1037106636|ref|XP_017145524.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1037032936|ref|XP_017140068.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1037033784|ref|XP_017140117.1| PREDICTED: lipase 3-like [Drosophila miranda]
>gi|1037100802|ref|XP_017144819.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100772|ref|XP_017144816.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100822|ref|XP_017144821.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100812|ref|XP_017144820.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100792|ref|XP_017144818.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100782|ref|XP_017144817.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037104707|ref|XP_017145251.1| PREDICTED: uncharacterized protein LOC108157625 [Drosophila
>gi|1037104713|ref|XP_017145252.1| PREDICTED: uncharacterized protein LOC108157626 isoform X1
>gi|1037104721|ref|XP_017145253.1| PREDICTED: uncharacterized protein LOC108157626 isoform X2
>gi|1037032585|ref|XP_017140046.1| PREDICTED: cardioacceleratory peptide receptor [Drosophila m
>gi|1037105484|ref|XP_017145365.1| PREDICTED: lateral signaling target protein 2 homolog [Dros
>gi|1037103023|ref|XP_017145037.1| PREDICTED: LOW QUALITY PROTEIN: protein bride of sevenless
>gi|1037105332|ref|XP_017145342.1| PREDICTED: uncharacterized protein LOC108157686 [Drosophila
>gi|1037063070|ref|XP_017142052.1| PREDICTED: neurofibromin isoform X3 [Drosophila miranda]
>gi|1037063036|ref|XP_017142049.1| PREDICTED: neurofibromin isoform X1 [Drosophila miranda]
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>gi|1037063052|ref|XP_017142051.1| PREDICTED: neurofibromin isoform X2 [Drosophila miranda]
>gi|1037104380|ref|XP_017145206.1| PREDICTED: enhancer of split mdelta protein [Drosophila mira
>gi|1037104584|ref|XP_017145234.1| PREDICTED: enhancer of split mgamma protein [Drosophila mira
>gi|1037103882|ref|XP_017145140.1| PREDICTED: enhancer of split mbeta protein [Drosophila miran
>gi|1037031455|ref|XP_017139981.1| PREDICTED: enhancer of split malpha protein [Drosophila mira
>gi|1037032348|ref|XP_017140030.1| PREDICTED: enhancer of split M1 protein [Drosophila miranda]
>gi|1037031528|ref|XP_017139986.1| PREDICTED: enhancer of split M2 protein [Drosophila miranda]
>gi|1037104850|ref|XP_017145272.1| PREDICTED: enhancer of split m3 protein [Drosophila miranda]
>gi|1037031418|ref|XP_017139979.1| PREDICTED: enhancer of split m4 protein [Drosophila miranda]
>gi|1037030924|ref|XP_017139948.1| PREDICTED: enhancer of split m5 protein [Drosophila miranda]
>gi|1037031475|ref|XP_017139982.1| PREDICTED: enhancer of split m6 protein [Drosophila miranda]
>gi|1037103953|ref|XP_017145149.1| PREDICTED: enhancer of split m7 protein [Drosophila miranda]
>gi|1037103403|ref|XP_017145078.1| PREDICTED: enhancer of split m8 protein [Drosophila miranda]
>gi|1037103385|ref|XP_017145077.1| PREDICTED: actin-binding protein IPP-like [Drosophila miran
>gi|1037049648|ref|XP_017141132.1| PREDICTED: protein groucho isoform X1 [Drosophila miranda]
>gi|1037049664|ref|XP_017141133.1| PREDICTED: protein groucho isoform X1 [Drosophila miranda]
>gi|1037049680|ref|XP_017141134.1| PREDICTED: protein groucho isoform X1 [Drosophila miranda]
>gi|1037049696|ref|XP_017141135.1| PREDICTED: protein groucho isoform X2 [Drosophila miranda]
>gi|1037049713|ref|XP_017141136.1| PREDICTED: exocyst complex component 8 [Drosophila miranda]
>gi|1037049747|ref|XP_017141138.1| PREDICTED: charged multivesicular body protein 2a [Drosophil
>gi|1037049779|ref|XP_017141141.1| PREDICTED: UMP-CMP kinase [Drosophila miranda]
>gi|1037049794|ref|XP_017141142.1| PREDICTED: DNA replication complex GINS protein SLD5 [Drosop
>gi|1037049810|ref|XP_017141143.1| PREDICTED: pre-rRNA-processing protein TSR2 homolog [Drosoph
>gi|1037049860|ref|XP_017141146.1| PREDICTED: phosphatidylinositol N-acetylglucosaminyltransfe
>gi|1037049844|ref|XP_017141145.1| PREDICTED: uncharacterized protein LOC108155069 [Drosophila
>gi|1037049828|ref|XP_017141144.1| PREDICTED: 60S ribosomal protein L34 [Drosophila miranda]
>gi|1037049729|ref|XP_017141137.1| PREDICTED: tRNA (adenine(58)-N(1))-methyltransferase cataly
>gi|1037049763|ref|XP_017141139.1| PREDICTED: uncharacterized protein LOC108155064 [Drosophila
>gi|1037104120|ref|XP_017145170.1| PREDICTED: neuronal PAS domain-containing protein 4 [Drosoph
>gi|1037102894|ref|XP_017145022.1| PREDICTED: uncharacterized protein LOC108157460 isoform X2
>gi|1037102885|ref|XP_017145021.1| PREDICTED: uncharacterized protein LOC108157460 isoform X1
>gi|1037102876|ref|XP_017145020.1| PREDICTED: uncharacterized protein LOC108157459 [Drosophila
>gi|1037106644|ref|XP_017145525.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1037106540|ref|XP_017145511.1| PREDICTED: GPI ethanolamine phosphate transferase 1-like [D
>gi|1037106657|ref|XP_017145527.1| PREDICTED: suppressor protein SRP40-like isoform X2 [Drosoph
>gi|1037106650|ref|XP_017145526.1| PREDICTED: clumping factor A-like isoform X1 [Drosophila mi
>gi|1037085656|ref|XP_017143631.1| PREDICTED: uncharacterized protein LOC108156599 [Drosophila
>gi|1037085600|ref|XP_017143627.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085614|ref|XP_017143628.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085628|ref|XP_017143629.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085642|ref|XP_017143630.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085670|ref|XP_017143632.1| PREDICTED: natterin-3 [Drosophila miranda]
>gi|1037085573|ref|XP_017143626.1| PREDICTED: protein HGH1 homolog [Drosophila miranda]
>gi|1037038365|ref|XP_017140382.1| PREDICTED: LOW QUALITY PROTEIN: proteasome subunit beta typ
>gi|1037033508|ref|XP_017140101.1| PREDICTED: neurotrypsin-like [Drosophila miranda]
>gi|1037101313|ref|XP_017144867.1| PREDICTED: uncharacterized protein LOC108157356 [Drosophila
>gi|1037037714|ref|XP_017140343.1| PREDICTED: uncharacterized protein LOC108154551 [Drosophila
>gi|1037023427|ref|XP_017139486.1| PREDICTED: uncharacterized protein LOC108153899 [Drosophila
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>gi|1037083391|ref|XP_017143460.1| PREDICTED: proteasome subunit beta type-7-like [Drosophila r
>gi|1037083375|ref|XP_017143459.1| PREDICTED: fructose-bisphosphate aldolase isoform X3 [Droso
>gi|1037083333|ref|XP_017143456.1| PREDICTED: fructose-bisphosphate aldolase isoform X1 [Droso
>gi|1037083363|ref|XP_017143458.1| PREDICTED: fructose-bisphosphate aldolase isoform X1 [Droso
>gi|1037083349|ref|XP_017143457.1| PREDICTED: fructose-bisphosphate aldolase isoform X2 [Droso
>gi|1037083319|ref|XP_017143455.1| PREDICTED: dipeptidase 1 [Drosophila miranda]
>gi|1037098293|ref|XP_017144601.1| PREDICTED: ABC transporter G family member 20 [Drosophila m:
>gi|1037023441|ref|XP_017139487.1| PREDICTED: uncharacterized protein LOC108153900 [Drosophila
>gi|1037106346|ref|XP_017145483.1| PREDICTED: fructose-bisphosphate aldolase-like [Drosophila r
>gi|1037036816|ref|XP_017140289.1| PREDICTED: uncharacterized protein LOC108154500 [Drosophila
>gi|1037038333|ref|XP_017140380.1| PREDICTED: uncharacterized protein LOC108154587 [Drosophila
>gi|1037103069|ref|XP_017145042.1| PREDICTED: helix-loop-helix protein delilah [Drosophila mira
>gi|1037033142|ref|XP_017140081.1| PREDICTED: hexokinase type 1 [Drosophila miranda]
>gi|1037106025|ref|XP_017145440.1| PREDICTED: hexokinase type 2 [Drosophila miranda]
>gi|1037043816|ref|XP_017140744.1| PREDICTED: probable uridine-cytidine kinase isoform X1 [Dros
>gi|1037043830|ref|XP_017140745.1| PREDICTED: probable uridine-cytidine kinase isoform X2 [Dros
>gi|1037043784|ref|XP_017140742.1| PREDICTED: zinc metalloproteinase nas-1 isoform X2 [Drosophi
>gi|1037043768|ref|XP_017140741.1| PREDICTED: uncharacterized protein LOC108154836 isoform X1
>gi|1037043800|ref|XP_017140743.1| PREDICTED: uncharacterized protein LOC108154838 [Drosophila
>gi|1037043846|ref|XP_017140746.1| PREDICTED: uncharacterized protein LOC108154840 [Drosophila
>gi|1037043737|ref|XP_017140739.1| PREDICTED: protein crumbs isoform X1 [Drosophila miranda]
>gi|1037043755|ref|XP_017140740.1| PREDICTED: protein crumbs isoform X2 [Drosophila miranda]
>gi|1037044023|ref|XP_017140753.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1037043992|ref|XP_017140751.1| PREDICTED: zinc finger CCCH domain-containing protein 14 [D:
>gi|1037044008|ref|XP_017140752.1| PREDICTED: uncharacterized protein LOC108154847 [Drosophila
>gi|1037043950|ref|XP_017140748.1| PREDICTED: bromodomain and WD repeat-containing protein 3 is
>gi|1037043962|ref|XP_017140749.1| PREDICTED: bromodomain and WD repeat-containing protein 3 is
>gi|1037043976|ref|XP_017140750.1| PREDICTED: protein RRP5 homolog [Drosophila miranda]
>gi|1037089702|ref|XP_017143913.1| PREDICTED: exosome complex exonuclease RRP44 [Drosophila mi:
>gi|1037105468|ref|XP_017145363.1| PREDICTED: acyl-CoA synthetase short-chain family member 3,
>gi|1037106681|ref|XP_017145530.1| PREDICTED: dromyosuppressin [Drosophila miranda]
>gi|1037064318|ref|XP_017142137.1| PREDICTED: RNA-binding protein FUS isoform X4 [Drosophila m:
>gi|1037064288|ref|XP_017142135.1| PREDICTED: RNA-binding protein FUS isoform X2 [Drosophila m:
>gi|1037064302|ref|XP_017142136.1| PREDICTED: RNA-binding protein FUS isoform X3 [Drosophila m:
>gi|1037064272|ref|XP_017142134.1| PREDICTED: RNA-binding protein FUS isoform X1 [Drosophila m:
>gi|1037064210|ref|XP_017142130.1| PREDICTED: C2 domain-containing protein 5 isoform X1 [Droso
>gi|1037064226|ref|XP_017142131.1| PREDICTED: C2 domain-containing protein 5 isoform X2 [Droso
>gi|1037064242|ref|XP_017142132.1| PREDICTED: golgin-84 [Drosophila miranda]
>gi|1037064258|ref|XP_017142133.1| PREDICTED: golgin-84 [Drosophila miranda]
>gi|1037091762|ref|XP_017144074.1| PREDICTED: tRNA modification GTPase GTPBP3, mitochondrial [I
>gi|1037091898|ref|XP_017144083.1| PREDICTED: uncharacterized protein LOC108156866 [Drosophila
>gi|1037091794|ref|XP_017144076.1| PREDICTED: uncharacterized protein LOC108156860 isoform X1
>gi|1037091806|ref|XP_017144077.1| PREDICTED: uncharacterized protein LOC108156860 isoform X2
>gi|1037091826|ref|XP_017144078.1| PREDICTED: uncharacterized protein LOC108156861 [Drosophila
>gi|1037091840|ref|XP_017144079.1| PREDICTED: uncharacterized protein LOC108156862 [Drosophila
>gi|1037091856|ref|XP_017144080.1| PREDICTED: uncharacterized protein LOC108156863 [Drosophila
>gi|1037091884|ref|XP_017144082.1| PREDICTED: uncharacterized protein LOC108156865 [Drosophila
>gi|1037091870|ref|XP_017144081.1| PREDICTED: uncharacterized protein LOC108156864 [Drosophila
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>gi|1037091780|ref|XP_017144075.1| PREDICTED: uncharacterized protein LOC108156859 [Drosophila
>gi|1037035853|ref|XP_017140233.1| PREDICTED: uncharacterized protein LOC108154472 [Drosophila
>gi|1037023461|ref|XP_017139488.1| PREDICTED: uncharacterized protein LOC108153901 [Drosophila
>gi|1037035770|ref|XP_017140228.1| PREDICTED: uncharacterized protein LOC108154468 [Drosophila
>gi|1037023479|ref|XP_017139489.1| PREDICTED: uncharacterized protein LOC108153902 [Drosophila
>gi|1037034559|ref|XP_017140167.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037023499|ref|XP_017139490.1| PREDICTED: uncharacterized protein LOC108153904 [Drosophila
>gi|1037023517|ref|XP_017139491.1| PREDICTED: uncharacterized protein LOC108153905 [Drosophila
>gi|1037035700|ref|XP_017140223.1| PREDICTED: uncharacterized protein LOC108154464 [Drosophila
>gi|1037074554|ref|XP_017142830.1| PREDICTED: acetylcholine receptor subunit alpha-like 1 [Dros
>gi|1037074616|ref|XP_017142834.1| PREDICTED: uncharacterized protein LOC108156062 [Drosophila
>gi|1037074632|ref|XP_017142835.1| PREDICTED: uncharacterized protein LOC108156063 [Drosophila
>gi|1037074568|ref|XP_017142831.1| PREDICTED: acetylcholine receptor subunit alpha-like 2 [Dros
>gi|1037074584|ref|XP_017142832.1| PREDICTED: acetylcholine receptor subunit beta-like 2 [Droso
>gi|1037074507|ref|XP_017142826.1| PREDICTED: zinc finger protein DZIP1L isoform X1 [Drosophila
>gi|1037074522|ref|XP_017142827.1| PREDICTED: zinc finger protein DZIP1L isoform X2 [Drosophila
>gi|1037074538|ref|XP_017142828.1| PREDICTED: zinc finger protein DZIP1L isoform X3 [Drosophila
>gi|1037074600|ref|XP_017142833.1| PREDICTED: serine/threonine-protein phosphatase alpha-1 iso
>gi|1037035667|ref|XP_017140222.1| PREDICTED: uncharacterized protein LOC108154461 [Drosophila
>gi|1037105834|ref|XP_017145413.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037097586|ref|XP_017144529.1| PREDICTED: protein phosphatase 1 regulatory subunit 21 [Dros
>gi|1037097597|ref|XP_017144530.1| PREDICTED: iron-sulfur cluster assembly 2 homolog, mitochond
>gi|1037056760|ref|XP_017141613.1| PREDICTED: uncharacterized protein LOC108155362 [Drosophila
>gi|1037056776|ref|XP_017141614.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037056792|ref|XP_017141615.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037065176|ref|XP_017142186.1| PREDICTED: proton-associated sugar transporter A isoform X2
>gi|1037065161|ref|XP_017142185.1| PREDICTED: proton-associated sugar transporter A isoform X1
>gi|1037065191|ref|XP_017142188.1| PREDICTED: proton-associated sugar transporter A isoform X3
>gi|1037065250|ref|XP_017142192.1| PREDICTED: uncharacterized protein LOC108155709 isoform X2
>gi|1037065219|ref|XP_017142190.1| PREDICTED: uncharacterized protein LOC108155709 isoform X1
>gi|1037065234|ref|XP_017142191.1| PREDICTED: uncharacterized protein LOC108155709 isoform X1
>gi|1037065206|ref|XP_017142189.1| PREDICTED: survival of motor neuron-related-splicing factor
>gi|1037091596|ref|XP_017144062.1| PREDICTED: synaptosomal-associated protein 25 isoform X2 [Dr
>gi|1037091583|ref|XP_017144061.1| PREDICTED: synaptosomal-associated protein 25 isoform X1 [Dr
>gi|1037091610|ref|XP_017144063.1| PREDICTED: synaptosomal-associated protein 25 isoform X3 [Dr
>gi|1037091637|ref|XP_017144064.1| PREDICTED: synaptosomal-associated protein 25 isoform X4 [Dr
>gi|1037092359|ref|XP_017144113.1| PREDICTED: 28S ribosomal protein S5, mitochondrial [Drosoph
>gi|1037045773|ref|XP_017140873.1| PREDICTED: casein kinase II subunit alpha [Drosophila miran
>gi|1037045791|ref|XP_017140874.1| PREDICTED: casein kinase II subunit alpha [Drosophila miran
>gi|1037045744|ref|XP_017140870.1| PREDICTED: unconventional myosin heavy chain 6 isoform X1 [
>gi|1037045759|ref|XP_017140872.1| PREDICTED: myosin-I heavy chain isoform X2 [Drosophila mira
>gi|1037094869|ref|XP_017144293.1| PREDICTED: DEAD-box helicase Dbp80 [Drosophila miranda]
>gi|1037023535|ref|XP_017139493.1| PREDICTED: LOW QUALITY PROTEIN: probable E3 ubiquitin-prote
>gi|1037100223|ref|XP_017144763.1| PREDICTED: LOW QUALITY PROTEIN: 60S ribosomal protein L15 [
>gi|1037096957|ref|XP_017144470.1| PREDICTED: uncharacterized protein LOC108157108 [Drosophila
>gi|1037096968|ref|XP_017144472.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomple
>gi|1037098733|ref|XP_017144642.1| PREDICTED: uncharacterized protein LOC108157205 [Drosophila
>gi|1037023553|ref|XP_017139494.1| PREDICTED: cholesterol 7-desaturase, partial [Drosophila mi
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>gi|1037089209|ref|XP_017143874.1| PREDICTED: uncharacterized protein LOC108156730 isoform X1
>gi|1037089223|ref|XP_017143875.1| PREDICTED: uncharacterized protein LOC108156730 isoform X2
>gi|1037089195|ref|XP_017143873.1| PREDICTED: vacuolar protein-sorting-associated protein 11 ho
>gi|1037074377|ref|XP_017142818.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074395|ref|XP_017142819.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074411|ref|XP_017142820.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074443|ref|XP_017142822.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074427|ref|XP_017142821.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074459|ref|XP_017142823.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074476|ref|XP_017142824.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037074492|ref|XP_017142825.1| PREDICTED: acetylcholine receptor subunit alpha-like isoform
>gi|1037028462|ref|XP_017139799.1| PREDICTED: synaptic vesicle glycoprotein 2C-like isoform X2
>gi|1037028480|ref|XP_017139801.1| PREDICTED: synaptic vesicle glycoprotein 2C-like isoform X2
>gi|1037028498|ref|XP_017139802.1| PREDICTED: synaptic vesicle glycoprotein 2C-like isoform X2
>gi|1037028444|ref|XP_017139798.1| PREDICTED: synaptic vesicle glycoprotein 2B-like isoform X1
>gi|1037028516|ref|XP_017139803.1| PREDICTED: synaptic vesicle glycoprotein 2B-like isoform X3
>gi|1037028534|ref|XP_017139804.1| PREDICTED: synaptic vesicle glycoprotein 2B-like isoform X4
>gi|1037028676|ref|XP_017139813.1| PREDICTED: protein NATD1 [Drosophila miranda]
>gi|1037028549|ref|XP_017139805.1| PREDICTED: uncharacterized protein LOC108154156 [Drosophila
>gi|1037028567|ref|XP_017139806.1| PREDICTED: uncharacterized protein LOC108154156 [Drosophila
>gi|1037028354|ref|XP_017139793.1| PREDICTED: dnaJ homolog subfamily C member 16 [Drosophila mi
>gi|1037028374|ref|XP_017139794.1| PREDICTED: dnaJ homolog subfamily C member 16 [Drosophila mi
>gi|1037028392|ref|XP_017139795.1| PREDICTED: dnaJ homolog subfamily C member 16 [Drosophila mi
>gi|1037028656|ref|XP_017139811.1| PREDICTED: homeobox protein abdominal-A homolog [Drosophila
>gi|1037028638|ref|XP_017139810.1| PREDICTED: 60S ribosomal protein L10 [Drosophila miranda]
>gi|1037028723|ref|XP_017139815.1| PREDICTED: cytochrome c oxidase subunit NDUFA4 [Drosophila m
>gi|1037028741|ref|XP_017139816.1| PREDICTED: cytochrome c oxidase subunit NDUFA4 [Drosophila m
>gi|1037028170|ref|XP_017139779.1| PREDICTED: uncharacterized protein LOC108154148 isoform X2
>gi|1037028150|ref|XP_017139778.1| PREDICTED: proline-, glutamic acid- and leucine-rich protei
>gi|1037028186|ref|XP_017139780.1| PREDICTED: proline-, glutamic acid- and leucine-rich protei
>gi|1037028044|ref|XP_017139772.1| PREDICTED: uncharacterized protein LOC108154146 isoform X3
>gi|1037028028|ref|XP_017139771.1| PREDICTED: uncharacterized protein LOC108154146 isoform X2
>gi|1037028011|ref|XP_017139769.1| PREDICTED: uncharacterized protein LOC108154146 isoform X1
>gi|1037028256|ref|XP_017139784.1| PREDICTED: forkhead box protein P1-like isoform X7 [Drosophi
>gi|1037028062|ref|XP_017139773.1| PREDICTED: aminopeptidase N isoform X1 [Drosophila miranda]
>gi|1037028080|ref|XP_017139774.1| PREDICTED: aminopeptidase N isoform X1 [Drosophila miranda]
>gi|1037028132|ref|XP_017139777.1| PREDICTED: endoplasmic reticulum aminopeptidase 1 isoform X3
>gi|1037028274|ref|XP_017139788.1| PREDICTED: forkhead box protein P1-like isoform X8 [Drosophi
>gi|1037028096|ref|XP_017139775.1| PREDICTED: aminopeptidase N isoform X2 [Drosophila miranda]
>gi|1037028114|ref|XP_017139776.1| PREDICTED: aminopeptidase N isoform X2 [Drosophila miranda]
>gi|1037028220|ref|XP_017139782.1| PREDICTED: uncharacterized protein LOC108154148 isoform X5
>gi|1037028204|ref|XP_017139781.1| PREDICTED: forkhead box protein P1-like isoform X4 [Drosophi
>gi|1037028290|ref|XP_017139789.1| PREDICTED: forkhead box protein P1-like isoform X9 [Drosophi
>gi|1037028238|ref|XP_017139783.1| PREDICTED: forkhead box protein P1-like isoform X6 [Drosophi
>gi|1037028308|ref|XP_017139790.1| PREDICTED: uncharacterized protein LOC108154148 isoform X10
>gi|1037028410|ref|XP_017139796.1| PREDICTED: xaa-Pro aminopeptidase 1 [Drosophila miranda]
>gi|1037028692|ref|XP_017139814.1| PREDICTED: uncharacterized protein LOC108154164 [Drosophila
>gi|1037028426|ref|XP_017139797.1| PREDICTED: calreticulin [Drosophila miranda]
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>gi|1037028585|ref|XP_017139807.1| PREDICTED: spermidine synthase [Drosophila miranda]
>gi|1037028620|ref|XP_017139809.1| PREDICTED: pinin isoform X2 [Drosophila miranda]
>gi|1037028602|ref|XP_017139808.1| PREDICTED: pinin isoform X1 [Drosophila miranda]
>gi|1037028324|ref|XP_017139791.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037028336|ref|XP_017139792.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037053440|ref|XP_017141393.1| PREDICTED: eIF-2-alpha kinase activator GCN1 [Drosophila mi
>gi|1037026348|ref|XP_017139672.1| PREDICTED: eukaryotic translation initiation factor 4B [Dros
>gi|1037026402|ref|XP_017139675.1| PREDICTED: uncharacterized protein LOC108154079 isoform X1
>gi|1037026420|ref|XP_017139676.1| PREDICTED: uncharacterized protein LOC108154079 isoform X2
>gi|1037026438|ref|XP_017139677.1| PREDICTED: uncharacterized protein LOC108154079 isoform X3
>gi|1037026458|ref|XP_017139679.1| PREDICTED: thioredoxin domain-containing protein isoform X4
>gi|1037026168|ref|XP_017139660.1| PREDICTED: YTH domain-containing family protein 3 isoform X2
>gi|1037026150|ref|XP_017139659.1| PREDICTED: uncharacterized protein LOC108154074 isoform X1
>gi|1037026184|ref|XP_017139661.1| PREDICTED: uncharacterized protein LOC108154074 isoform X3
>gi|1037026476|ref|XP_017139680.1| PREDICTED: purine nucleoside phosphorylase isoform X1 [Droso
>gi|1037026494|ref|XP_017139681.1| PREDICTED: purine nucleoside phosphorylase isoform X2 [Droso
>gi|1037026606|ref|XP_017139688.1| PREDICTED: transmembrane emp24 domain-containing protein ba
>gi|1037026258|ref|XP_017139667.1| PREDICTED: type I inositol 1,4,5-trisphosphate 5-phosphatase
>gi|1037026330|ref|XP_017139671.1| PREDICTED: uncharacterized protein LOC108154075 isoform X8
>gi|1037026312|ref|XP_017139670.1| PREDICTED: type I inositol 1,4,5-trisphosphate 5-phosphatase
>gi|1037026276|ref|XP_017139668.1| PREDICTED: type I inositol 1,4,5-trisphosphate 5-phosphatase
>gi|1037026294|ref|XP_017139669.1| PREDICTED: type I inositol 1,4,5-trisphosphate 5-phosphatase
>gi|1037026222|ref|XP_017139665.1| PREDICTED: uncharacterized protein LOC108154075 isoform X2
>gi|1037026240|ref|XP_017139666.1| PREDICTED: uncharacterized protein LOC108154075 isoform X3
>gi|1037026202|ref|XP_017139664.1| PREDICTED: type I inositol 1,4,5-trisphosphate 5-phosphatase
>gi|1037026059|ref|XP_017139654.1| PREDICTED: E3 ubiquitin-protein ligase hyd isoform X3 [Droso
>gi|1037026044|ref|XP_017139653.1| PREDICTED: E3 ubiquitin-protein ligase hyd isoform X2 [Droso
>gi|1037026026|ref|XP_017139652.1| PREDICTED: E3 ubiquitin-protein ligase hyd isoform X1 [Droso
>gi|1037026512|ref|XP_017139682.1| PREDICTED: transforming growth factor beta regulator 1 [Dros
>gi|1037023573|ref|XP_017139495.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037026584|ref|XP_017139687.1| PREDICTED: zinc finger protein 830 [Drosophila miranda]
>gi|1037026624|ref|XP_017139689.1| PREDICTED: uncharacterized protein LOC108154086 [Drosophila
>gi|1037026530|ref|XP_017139683.1| PREDICTED: uncharacterized protein LOC108154082 [Drosophila
>gi|1037026548|ref|XP_017139684.1| PREDICTED: protein lin-7 homolog C isoform X1 [Drosophila mi
>gi|1037026564|ref|XP_017139685.1| PREDICTED: protein lin-7 homolog C isoform X2 [Drosophila mi
>gi|1037026366|ref|XP_017139673.1| PREDICTED: trypsin-1 isoform X1 [Drosophila miranda]
>gi|1037026384|ref|XP_017139674.1| PREDICTED: trypsin-1 isoform X2 [Drosophila miranda]
>gi|1037026640|ref|XP_017139690.1| PREDICTED: uncharacterized protein LOC108154087 [Drosophila
>gi|1037026077|ref|XP_017139655.1| PREDICTED: protein unc-13 homolog D isoform X1 [Drosophila r
>gi|1037026095|ref|XP_017139656.1| PREDICTED: protein unc-13 homolog D isoform X2 [Drosophila r
>gi|1037026130|ref|XP_017139658.1| PREDICTED: BAI1-associated protein 3 isoform X4 [Drosophila
>gi|1037026113|ref|XP_017139657.1| PREDICTED: BAI1-associated protein 3 isoform X3 [Drosophila
>gi|1037088894|ref|XP_017143849.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1037079071|ref|XP_017143145.1| PREDICTED: glucoside xylosyltransferase 2 isoform X1 [Drosop
>gi|1037079087|ref|XP_017143146.1| PREDICTED: glucoside xylosyltransferase 2 isoform X2 [Drosop
>gi|1037079101|ref|XP_017143147.1| PREDICTED: U3 small nucleolar ribonucleoprotein protein IMP4
>gi|1037058685|ref|XP_017141748.1| PREDICTED: fork head domain-containing protein FD4 [Drosophi
>gi|1037058717|ref|XP_017141750.1| PREDICTED: uncharacterized protein LOC108155445 [Drosophila
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>gi|1037023591|ref|XP_017139496.1| PREDICTED: uncharacterized protein LOC108153911 [Drosophila
>gi|1037058701|ref|XP_017141749.1| PREDICTED: uncharacterized protein LOC108155444 [Drosophila
>gi|1037083045|ref|XP_017143431.1| PREDICTED: protein ABHD17B [Drosophila miranda]
>gi|1037083029|ref|XP_017143429.1| PREDICTED: uncharacterized protein LOC108156473 [Drosophila
>gi|1037048466|ref|XP_017141050.1| PREDICTED: uncharacterized protein LOC108155010 [Drosophila
>gi|1037048480|ref|XP_017141051.1| PREDICTED: uncharacterized protein LOC108155011 [Drosophila
>gi|1037048436|ref|XP_017141048.1| PREDICTED: uncharacterized protein LOC108155008 [Drosophila
>gi|1037048452|ref|XP_017141049.1| PREDICTED: uncharacterized protein LOC108155009 [Drosophila
>gi|1037048418|ref|XP_017141047.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1037048407|ref|XP_017141045.1| PREDICTED: tubulin polyglutamylase TTLL5 isoform X4 [Drosop
>gi|1037048379|ref|XP_017141043.1| PREDICTED: tubulin polyglutamylase TTLL5 isoform X2 [Drosop
>gi|1037048395|ref|XP_017141044.1| PREDICTED: tubulin polyglutamylase TTLL5 isoform X3 [Drosop
>gi|1037048363|ref|XP_017141042.1| PREDICTED: uncharacterized protein LOC108155005 isoform X1
>gi|1037087870|ref|XP_017143769.1| PREDICTED: plasminogen activator inhibitor 1 RNA-binding pro
>gi|1037087882|ref|XP_017143770.1| PREDICTED: plasminogen activator inhibitor 1 RNA-binding pro
>gi|1037087896|ref|XP_017143771.1| PREDICTED: uncharacterized protein LOC108156664 isoform X2
>gi|1037097645|ref|XP_017144535.1| PREDICTED: molybdopterin synthase catalytic subunit isoform
>gi|1037097633|ref|XP_017144534.1| PREDICTED: molybdopterin synthase catalytic subunit isoform
>gi|1037097657|ref|XP_017144536.1| PREDICTED: molybdopterin synthase sulfur carrier subunit [D
>gi|1037088430|ref|XP_017143812.1| PREDICTED: nuclear export mediator factor NEMF homolog [Dros
>gi|1037094785|ref|XP_017144286.1| PREDICTED: FERM domain-containing protein 8 isoform X1 [Dros
>gi|1037094799|ref|XP_017144287.1| PREDICTED: probable basic-leucine zipper transcription facto
>gi|1037103454|ref|XP_017145084.1| PREDICTED: protein lunapark-B-like [Drosophila miranda]
>gi|1037034141|ref|XP_017140139.1| PREDICTED: protein distal antenna-related [Drosophila mirand
>gi|1037075815|ref|XP_017142917.1| PREDICTED: coiled-coil domain-containing protein lobo isofor
>gi|1037075847|ref|XP_017142920.1| PREDICTED: coiled-coil domain-containing protein lobo isofor
>gi|1037075863|ref|XP_017142921.1| PREDICTED: coiled-coil domain-containing protein lobo isofor
>gi|1037075831|ref|XP_017142919.1| PREDICTED: coiled-coil domain-containing protein lobo isofor
>gi|1037075895|ref|XP_017142923.1| PREDICTED: protein distal antenna [Drosophila miranda]
>gi|1037075879|ref|XP_017142922.1| PREDICTED: ras-like protein rasD [Drosophila miranda]
>gi|1037031546|ref|XP_017139987.1| PREDICTED: uncharacterized protein LOC108154262 [Drosophila
>gi|1037078805|ref|XP_017143124.1| PREDICTED: tyrosine kinase receptor Cad96Ca [Drosophila mira
>gi|1037078791|ref|XP_017143122.1| PREDICTED: uncharacterized protein LOC108156266 isoform X2
>gi|1037078763|ref|XP_017143120.1| PREDICTED: uncharacterized protein LOC108156266 isoform X1
>gi|1037078777|ref|XP_017143121.1| PREDICTED: uncharacterized protein LOC108156266 isoform X1
>gi|1037078819|ref|XP_017143125.1| PREDICTED: uncharacterized protein LOC108156268 [Drosophila
>gi|1037087982|ref|XP_017143778.1| PREDICTED: synaptic vesicle glycoprotein 2B-like isoform X1
>gi|1037087996|ref|XP_017143779.1| PREDICTED: synaptic vesicle glycoprotein 2A-like isoform X2
>gi|1037087924|ref|XP_017143773.1| PREDICTED: protein anoxia up-regulated-like isoform X2 [Dros
>gi|1037087910|ref|XP_017143772.1| PREDICTED: protein anoxia up-regulated-like isoform X1 [Dros
>gi|1037087938|ref|XP_017143774.1| PREDICTED: protein anoxia up-regulated-like isoform X3 [Dros
>gi|1037087968|ref|XP_017143777.1| PREDICTED: protein anoxia up-regulated-like isoform X5 [Dros
>gi|1037087952|ref|XP_017143775.1| PREDICTED: uncharacterized protein LOC108156666 isoform X4
>gi|1037102373|ref|XP_017144970.1| PREDICTED: myb-like protein X isoform X2 [Drosophila miranda
>gi|1037102363|ref|XP_017144969.1| PREDICTED: myb-like protein X isoform X1 [Drosophila miranda
>gi|1037102383|ref|XP_017144971.1| PREDICTED: uncharacterized protein LOC108157431 isoform X1
>gi|1037102393|ref|XP_017144972.1| PREDICTED: uncharacterized protein LOC108157431 isoform X1
>gi|1037102404|ref|XP_017144973.1| PREDICTED: uncharacterized protein LOC108157431 isoform X2
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>gi|1037032794|ref|XP_017140060.1| PREDICTED: uncharacterized protein LOC108154313 [Drosophila
>gi|1037105969|ref|XP_017145432.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037097494|ref|XP_017144520.1| PREDICTED: 39S ribosomal protein L37, mitochondrial [Drosop
>gi|1037023607|ref|XP_017139497.1| PREDICTED: protein arginine N-methyltransferase 1-B-like [D
>gi|1037069894|ref|XP_017142509.1| PREDICTED: protein arginine N-methyltransferase 1-B-like [D
>gi|1037069878|ref|XP_017142508.1| PREDICTED: DNA replication licensing factor Mcm5 [Drosophila
>gi|1037069910|ref|XP_017142510.1| PREDICTED: abnormal spindle-like microcephaly-associated pr
>gi|1037023625|ref|XP_017139498.1| PREDICTED: uncharacterized protein LOC108153914 [Drosophila
>gi|1037105087|ref|XP_017145306.1| PREDICTED: tubulin polyglutamylase TTLL6 isoform X1 [Drosop
>gi|1037105095|ref|XP_017145307.1| PREDICTED: tubulin polyglutamylase TTLL6 isoform X2 [Drosop
>gi|1037095750|ref|XP_017144363.1| PREDICTED: cadherin-86C isoform X1 [Drosophila miranda]
>gi|1037095764|ref|XP_017144364.1| PREDICTED: cadherin-86C isoform X2 [Drosophila miranda]
>gi|1037095776|ref|XP_017144365.1| PREDICTED: cadherin-86C isoform X3 [Drosophila miranda]
>gi|1037095788|ref|XP_017144366.1| PREDICTED: cadherin-86C isoform X4 [Drosophila miranda]
>gi|1037090723|ref|XP_017143996.1| PREDICTED: protein midgut expression 1 [Drosophila miranda]
>gi|1037023641|ref|XP_017139499.1| PREDICTED: histone-lysine N-methyltransferase SETMAR [Droso
>gi|1037054097|ref|XP_017141437.1| PREDICTED: lysophospholipase-like protein 1 [Drosophila mir
>gi|1037054017|ref|XP_017141431.1| PREDICTED: kelch-like ECH-associated protein 1 isoform X1 [
>gi|1037054029|ref|XP_017141432.1| PREDICTED: kelch-like ECH-associated protein 1 isoform X2 [
>gi|1037054045|ref|XP_017141433.1| PREDICTED: kelch-like ECH-associated protein 1 isoform X2 [
>gi|1037054109|ref|XP_017141438.1| PREDICTED: 28S ribosomal protein S11, mitochondrial [Drosop
>gi|1037054079|ref|XP_017141436.1| PREDICTED: guanine nucleotide-binding protein-like 3 homolog
>gi|1037054063|ref|XP_017141435.1| PREDICTED: mothers against decapentaplegic homolog 6 [Droso
>gi|1037081557|ref|XP_017143322.1| PREDICTED: protein PAT1 homolog 1 isoform X2 [Drosophila mi
>gi|1037081543|ref|XP_017143321.1| PREDICTED: protein PAT1 homolog 1 isoform X1 [Drosophila mi
>gi|1037093227|ref|XP_017144172.1| PREDICTED: uncharacterized protein LOC108156928 [Drosophila
>gi|1037093241|ref|XP_017144173.1| PREDICTED: putative tRNA (cytidine(32)/guanosine(34)-2'-O)-m
>gi|1037094134|ref|XP_017144237.1| PREDICTED: uncharacterized protein LOC108156966 isoform X1
>gi|1037094147|ref|XP_017144238.1| PREDICTED: uncharacterized protein LOC108156966 isoform X2
>gi|1037094175|ref|XP_017144240.1| PREDICTED: uncharacterized protein LOC108156966 isoform X4
>gi|1037094189|ref|XP_017144241.1| PREDICTED: uncharacterized protein LOC108156966 isoform X5
>gi|1037094163|ref|XP_017144239.1| PREDICTED: uncharacterized protein LOC108156966 isoform X3
>gi|1037094215|ref|XP_017144243.1| PREDICTED: uncharacterized protein LOC108156966 isoform X7
>gi|1037094203|ref|XP_017144242.1| PREDICTED: uncharacterized protein LOC108156966 isoform X6
>gi|1037103929|ref|XP_017145146.1| PREDICTED: peroxidase [Drosophila miranda]
>gi|1037023659|ref|XP_017139500.1| PREDICTED: uncharacterized protein LOC108153916 [Drosophila
>gi|1037080373|ref|XP_017143231.1| PREDICTED: peroxidase [Drosophila miranda]
>gi|1037080387|ref|XP_017143232.1| PREDICTED: adenylosuccinate lyase [Drosophila miranda]
>gi|1037080401|ref|XP_017143233.1| PREDICTED: adenylosuccinate lyase [Drosophila miranda]
>gi|1037080345|ref|XP_017143229.1| PREDICTED: epidermal growth factor receptor kinase substrate
>gi|1037080359|ref|XP_017143230.1| PREDICTED: chorion peroxidase [Drosophila miranda]
>gi|1037082241|ref|XP_017143377.1| PREDICTED: splicing factor 3A subunit 1 [Drosophila miranda]
>gi|1037082255|ref|XP_017143379.1| PREDICTED: xylulose kinase isoform X1 [Drosophila miranda]
>gi|1037082269|ref|XP_017143380.1| PREDICTED: xylulose kinase isoform X2 [Drosophila miranda]
>gi|1037083626|ref|XP_017143478.1| PREDICTED: ER membrane protein complex subunit 2 [Drosophila
>gi|1037083613|ref|XP_017143477.1| PREDICTED: uncharacterized protein LOC108156495 isoform X2
>gi|1037083600|ref|XP_017143476.1| PREDICTED: uncharacterized protein LOC108156495 isoform X1
>gi|1037083639|ref|XP_017143480.1| PREDICTED: uncharacterized protein LOC108156497 [Drosophila
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>gi|1037083653|ref|XP_017143481.1| PREDICTED: uncharacterized protein LOC108156497 [Drosophila
>gi|1037083667|ref|XP_017143482.1| PREDICTED: uncharacterized protein LOC108156497 [Drosophila
>gi|1037106827|ref|XP_017145552.1| PREDICTED: EH domain-binding protein 1 [Drosophila miranda]
>gi|1037106835|ref|XP_017145553.1| PREDICTED: V-type proton ATPase subunit e [Drosophila mirand
>gi|1037084508|ref|XP_017143548.1| PREDICTED: homeobox protein abdominal-B [Drosophila miranda]
>gi|1037053742|ref|XP_017141415.1| PREDICTED: uncharacterized protein LOC108155234 [Drosophila
>gi|1037053756|ref|XP_017141416.1| PREDICTED: uncharacterized protein LOC108155234 [Drosophila
>gi|1037053770|ref|XP_017141417.1| PREDICTED: uncharacterized protein LOC108155234 [Drosophila
>gi|1037053697|ref|XP_017141411.1| PREDICTED: homeobox protein abdominal-A isoform X1 [Drosophi
>gi|1037053713|ref|XP_017141413.1| PREDICTED: homeobox protein abdominal-A isoform X1 [Drosophi
>gi|1037053728|ref|XP_017141414.1| PREDICTED: homeobox protein abdominal-A isoform X2 [Drosophi
>gi|1037105324|ref|XP_017145341.1| PREDICTED: glucose transporter type 3 [Drosophila miranda]
>gi|1037085234|ref|XP_017143603.1| PREDICTED: dynein light chain roadblock-type 1 [Drosophila m
>gi|1037082843|ref|XP_017143418.1| PREDICTED: homeotic protein ultrabithorax isoform X1 [Drosop
>gi|1037082857|ref|XP_017143419.1| PREDICTED: homeotic protein ultrabithorax isoform X2 [Drosop
>gi|1037082869|ref|XP_017143420.1| PREDICTED: homeotic protein ultrabithorax isoform X3 [Drosop
>gi|1037079772|ref|XP_017143184.1| PREDICTED: uncharacterized protein LOC108156306 [Drosophila
>gi|1037079719|ref|XP_017143181.1| PREDICTED: suppressor of tumorigenicity 14 protein homolog
>gi|1037079733|ref|XP_017143182.1| PREDICTED: coiled-coil domain-containing protein 63 [Drosop
>gi|1037079760|ref|XP_017143183.1| PREDICTED: coiled-coil domain-containing protein 63 [Drosop
>gi|1037051843|ref|XP_017141285.1| PREDICTED: methyltransferase-like protein 4 [Drosophila mir
>gi|1037051761|ref|XP_017141279.1| PREDICTED: T-complex protein 1 subunit gamma [Drosophila mi
>gi|1037051777|ref|XP_017141280.1| PREDICTED: G patch domain and KOW motifs-containing protein
>gi|1037051871|ref|XP_017141287.1| PREDICTED: ARL-6-interacting protein 1 homolog [Drosophila m
>gi|1037051859|ref|XP_017141286.1| PREDICTED: uncharacterized protein LOC108155143 [Drosophila
>gi|1037051811|ref|XP_017141282.1| PREDICTED: UPF0609 protein CG1218 [Drosophila miranda]
>gi|1037051795|ref|XP_017141281.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 12
>gi|1037051827|ref|XP_017141283.1| PREDICTED: histone acetyltransferase type B catalytic subuni
>gi|1037051746|ref|XP_017141278.1| PREDICTED: katanin p60 ATPase-containing subunit A1 isoform
>gi|1037051730|ref|XP_017141277.1| PREDICTED: katanin p60 ATPase-containing subunit A1 isoform
>gi|1037051714|ref|XP_017141276.1| PREDICTED: katanin p60 ATPase-containing subunit A1 isoform
>gi|1037051699|ref|XP_017141275.1| PREDICTED: katanin p60 ATPase-containing subunit A1 isoform
>gi|1037065129|ref|XP_017142183.1| PREDICTED: protein jagunal isoform X1 [Drosophila miranda]
>gi|1037065145|ref|XP_017142184.1| PREDICTED: protein jagunal isoform X2 [Drosophila miranda]
>gi|1037064951|ref|XP_017142171.1| PREDICTED: exocyst complex component 4 isoform X1 [Drosophil
>gi|1037064967|ref|XP_017142173.1| PREDICTED: exocyst complex component 4 isoform X1 [Drosophil
>gi|1037064983|ref|XP_017142174.1| PREDICTED: exocyst complex component 4 isoform X2 [Drosophil
>gi|1037064999|ref|XP_017142175.1| PREDICTED: exocyst complex component 4 isoform X3 [Drosophil
>gi|1037065017|ref|XP_017142176.1| PREDICTED: exocyst complex component 4 isoform X4 [Drosophil
>gi|1037065051|ref|XP_017142178.1| PREDICTED: exocyst complex component 4 isoform X6 [Drosophil
>gi|1037065098|ref|XP_017142181.1| PREDICTED: uncharacterized protein ZK1073.1 isoform X9 [Dros
>gi|1037065114|ref|XP_017142182.1| PREDICTED: uncharacterized protein ZK1073.1 isoform X10 [Dro
>gi|1037065032|ref|XP_017142177.1| PREDICTED: exocyst complex component 4 isoform X5 [Drosophil
>gi|1037065066|ref|XP_017142179.1| PREDICTED: uncharacterized protein ZK1073.1 isoform X7 [Dros
>gi|1037065082|ref|XP_017142180.1| PREDICTED: uncharacterized protein ZK1073.1 isoform X8 [Dros
>gi|1037082677|ref|XP_017143407.1| PREDICTED: m7GpppX diphosphatase [Drosophila miranda]
>gi|1037082664|ref|XP_017143406.1| PREDICTED: nuclear receptor-binding protein homolog [Drosoph
>gi|1037069453|ref|XP_017142478.1| PREDICTED: symplekin [Drosophila miranda]
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>gi|1037069480|ref|XP_017142480.1| PREDICTED: glyoxylate reductase/hydroxypyruvate reductase [
>gi|1037069469|ref|XP_017142479.1| PREDICTED: CCA tRNA nucleotidyltransferase 1, mitochondrial
>gi|1037069496|ref|XP_017142481.1| PREDICTED: probable RNA methyltransferase CG1239 isoform X1
>gi|1037069512|ref|XP_017142482.1| PREDICTED: probable RNA methyltransferase CG1239 isoform X1
>gi|1037069528|ref|XP_017142483.1| PREDICTED: probable RNA methyltransferase CG1239 isoform X2
>gi|1037068800|ref|XP_017142431.1| PREDICTED: transcription factor castor [Drosophila miranda]
>gi|1037068739|ref|XP_017142427.1| PREDICTED: TBC1 domain family member 4 isoform X2 [Drosophil
>gi|1037068677|ref|XP_017142423.1| PREDICTED: TBC1 domain family member 4 isoform X1 [Drosophil
>gi|1037068693|ref|XP_017142424.1| PREDICTED: TBC1 domain family member 4 isoform X1 [Drosophil
>gi|1037068709|ref|XP_017142425.1| PREDICTED: TBC1 domain family member 4 isoform X1 [Drosophil
>gi|1037068725|ref|XP_017142426.1| PREDICTED: TBC1 domain family member 4 isoform X1 [Drosophil
>gi|1037068784|ref|XP_017142430.1| PREDICTED: TBC1 domain family member 4 isoform X5 [Drosophil
>gi|1037068755|ref|XP_017142428.1| PREDICTED: TBC1 domain family member 4 isoform X3 [Drosophil
>gi|1037068769|ref|XP_017142429.1| PREDICTED: TBC1 domain family member 4 isoform X4 [Drosophil
>gi|1037068814|ref|XP_017142432.1| PREDICTED: serine/threonine-protein phosphatase 2A activator
>gi|1037085379|ref|XP_017143612.1| PREDICTED: ras-related protein Rab-23 [Drosophila miranda]
>gi|1037085393|ref|XP_017143613.1| PREDICTED: ras-related protein Rab-23 [Drosophila miranda]
>gi|1037085407|ref|XP_017143614.1| PREDICTED: ras-related protein Rab-23 [Drosophila miranda]
>gi|1037085421|ref|XP_017143615.1| PREDICTED: ras-related protein Rab-23 [Drosophila miranda]
>gi|1037085435|ref|XP_017143616.1| PREDICTED: ras-related protein Rab-23 [Drosophila miranda]
>gi|1037075913|ref|XP_017142924.1| PREDICTED: RB1-inducible coiled-coil protein 1 [Drosophila m
>gi|1037075929|ref|XP_017142925.1| PREDICTED: RB1-inducible coiled-coil protein 1 [Drosophila m
>gi|1037075944|ref|XP_017142926.1| PREDICTED: RB1-inducible coiled-coil protein 1 [Drosophila m
>gi|1037075960|ref|XP_017142928.1| PREDICTED: RB1-inducible coiled-coil protein 1 [Drosophila m
>gi|1037041542|ref|XP_017140597.1| PREDICTED: A-kinase anchor protein 17A isoform X1 [Drosophil
>gi|1037041566|ref|XP_017140598.1| PREDICTED: A-kinase anchor protein 17A isoform X2 [Drosophil
>gi|1037041526|ref|XP_017140595.1| PREDICTED: pre-mRNA 3' end processing protein WDR33 [Drosoph
>gi|1037041697|ref|XP_017140606.1| PREDICTED: glucose 1-dehydrogenase-like [Drosophila miranda]
>gi|1037041681|ref|XP_017140605.1| PREDICTED: cyclopentanol dehydrogenase-like [Drosophila mira
>gi|1037041667|ref|XP_017140604.1| PREDICTED: uncharacterized oxidoreductase TM_0325-like [Dros
>gi|1037041713|ref|XP_017140607.1| PREDICTED: 60S ribosomal protein L13a [Drosophila miranda]
>gi|1037041729|ref|XP_017140608.1| PREDICTED: DPH4 homolog isoform X1 [Drosophila miranda]
>gi|1037041745|ref|XP_017140609.1| PREDICTED: DPH4 homolog isoform X2 [Drosophila miranda]
>gi|1037041757|ref|XP_017140610.1| PREDICTED: CDC42 small effector protein homolog [Drosophila
>gi|1037041574|ref|XP_017140599.1| PREDICTED: another transcription unit protein [Drosophila mi
>gi|1037041587|ref|XP_017140601.1| PREDICTED: regulator of gene activity isoform X1 [Drosophila
>gi|1037041636|ref|XP_017140602.1| PREDICTED: regulator of gene activity isoform X2 [Drosophila
>gi|1037041652|ref|XP_017140603.1| PREDICTED: regulator of gene activity isoform X3 [Drosophila
>gi|1037080511|ref|XP_017143242.1| PREDICTED: CAP-Gly domain-containing linker protein 1 [Droso
>gi|1037080525|ref|XP_017143243.1| PREDICTED: CAP-Gly domain-containing linker protein 1 [Droso
>gi|1037080539|ref|XP_017143244.1| PREDICTED: O-phosphoseryl-tRNA(Sec) selenium transferase [Dr
>gi|1037085823|ref|XP_017143642.1| PREDICTED: uncharacterized protein LOC108156609 isoform X1
>gi|1037085837|ref|XP_017143643.1| PREDICTED: uncharacterized protein LOC108156609 isoform X2
>gi|1037085851|ref|XP_017143644.1| PREDICTED: uncharacterized protein LOC108156609 isoform X3
>gi|1037035252|ref|XP_017140204.1| PREDICTED: uncharacterized protein LOC108154443 [Drosophila
>gi|1037023677|ref|XP_017139502.1| PREDICTED: putative odorant receptor 85e [Drosophila miranda
>gi|1037106550|ref|XP_017145512.1| PREDICTED: probable cytochrome P450 313b1 [Drosophila mirand
>gi|1037069377|ref|XP_017142472.1| PREDICTED: dual specificity mitogen-activated protein kinase
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>gi|1037069423|ref|XP_017142476.1| PREDICTED: SRA stem-loop-interacting RNA-binding protein, m
>gi|1037069393|ref|XP_017142473.1| PREDICTED: dihydroorotate dehydrogenase (quinone), mitochond
>gi|1037069407|ref|XP_017142474.1| PREDICTED: protein SYS1 homolog [Drosophila miranda]
>gi|1037069439|ref|XP_017142477.1| PREDICTED: uncharacterized protein LOC108155869 [Drosophila
>gi|1037070410|ref|XP_017142544.1| PREDICTED: ATP-dependent RNA helicase bel isoform X2 [Drosop
>gi|1037070392|ref|XP_017142543.1| PREDICTED: ATP-dependent RNA helicase bel isoform X1 [Drosop
>gi|1037070376|ref|XP_017142541.1| PREDICTED: Hermansky-Pudlak syndrome 5 protein homolog [Dros
>gi|1037023695|ref|XP_017139503.1| PREDICTED: RNA-directed DNA polymerase from mobile element j
>gi|1037090295|ref|XP_017143962.1| PREDICTED: histone H2A-like [Drosophila miranda]
>gi|1037090309|ref|XP_017143963.1| PREDICTED: histone H4 [Drosophila miranda]
>gi|1037090281|ref|XP_017143960.1| PREDICTED: histone H3 [Drosophila miranda]
>gi|1037088785|ref|XP_017143841.1| PREDICTED: histone H1-like [Drosophila miranda]
>gi|1037088812|ref|XP_017143843.1| PREDICTED: histone H2B [Drosophila miranda]
>gi|1037088800|ref|XP_017143842.1| PREDICTED: histone H2A [Drosophila miranda]
>gi|1037075326|ref|XP_017142883.1| PREDICTED: RNA-directed DNA polymerase from mobile element j
>gi|1037065492|ref|XP_017142206.1| PREDICTED: putative adenosylhomocysteinase 2 [Drosophila mi
>gi|1037065511|ref|XP_017142207.1| PREDICTED: putative adenosylhomocysteinase 2 [Drosophila mi
>gi|1037065439|ref|XP_017142202.1| PREDICTED: peroxisomal N(1)-acetyl-spermine/spermidine oxida
>gi|1037065461|ref|XP_017142204.1| PREDICTED: peroxisomal N(1)-acetyl-spermine/spermidine oxida
>gi|1037065476|ref|XP_017142205.1| PREDICTED: spermine oxidase isoform X2 [Drosophila miranda]
>gi|1037065423|ref|XP_017142201.1| PREDICTED: transketolase-like protein 2 [Drosophila miranda]
>gi|1037100350|ref|XP_017144775.1| PREDICTED: transcription termination factor 2 isoform X1 [Dr
>gi|1037100360|ref|XP_017144776.1| PREDICTED: transcription termination factor 2 isoform X2 [Dr
>gi|1037100372|ref|XP_017144777.1| PREDICTED: transcription termination factor 2 isoform X2 [Dr
>gi|1037075198|ref|XP_017142874.1| PREDICTED: transcription termination factor 2 [Drosophila m
>gi|1037075216|ref|XP_017142875.1| PREDICTED: protein doublesex isoform X1 [Drosophila miranda]
>gi|1037075233|ref|XP_017142876.1| PREDICTED: protein doublesex isoform X2 [Drosophila miranda]
>gi|1037092574|ref|XP_017144130.1| PREDICTED: neutral and basic amino acid transport protein rh
>gi|1037104947|ref|XP_017145286.1| PREDICTED: antennal-specific protein OS-C isoform X1 [Drosop
>gi|1037104955|ref|XP_017145287.1| PREDICTED: antennal-specific protein OS-C isoform X2 [Drosop
>gi|1037104939|ref|XP_017145285.1| PREDICTED: uncharacterized protein LOC108157647 [Drosophila
>gi|1037102953|ref|XP_017145029.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037102934|ref|XP_017145027.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037102943|ref|XP_017145028.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037080002|ref|XP_017143203.1| PREDICTED: integrator complex subunit 5 [Drosophila miranda]
>gi|1037080016|ref|XP_017143204.1| PREDICTED: integrator complex subunit 5 [Drosophila miranda]
>gi|1037080030|ref|XP_017143205.1| PREDICTED: serine/threonine-protein kinase stk11 [Drosophila
>gi|1037080057|ref|XP_017143206.1| PREDICTED: serine/threonine-protein kinase stk11 [Drosophila
>gi|1037097881|ref|XP_017144558.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 9 [
>gi|1037097860|ref|XP_017144556.1| PREDICTED: two pore potassium channel protein sup-9 isoform
>gi|1037097872|ref|XP_017144557.1| PREDICTED: two pore potassium channel protein sup-9 isoform
>gi|1037051616|ref|XP_017141269.1| PREDICTED: serine-arginine protein 55 isoform X2 [Drosophila
>gi|1037051632|ref|XP_017141270.1| PREDICTED: serine-arginine protein 55 isoform X3 [Drosophila
>gi|1037051598|ref|XP_017141268.1| PREDICTED: serine-arginine protein 55 isoform X1 [Drosophila
>gi|1037051649|ref|XP_017141271.1| PREDICTED: serine-arginine protein 55 isoform X4 [Drosophila
>gi|1037051521|ref|XP_017141263.1| PREDICTED: heterogeneous nuclear ribonucleoprotein 87F isof
>gi|1037051537|ref|XP_017141264.1| PREDICTED: heterogeneous nuclear ribonucleoprotein 87F isof
>gi|1037051550|ref|XP_017141265.1| PREDICTED: heterogeneous nuclear ribonucleoprotein 87F isof
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>gi|1037051566|ref|XP_017141266.1| PREDICTED: ribonucleoprotein RB97D [Drosophila miranda]
>gi|1037051582|ref|XP_017141267.1| PREDICTED: ribonucleoprotein RB97D [Drosophila miranda]
>gi|1037051665|ref|XP_017141273.1| PREDICTED: ras GTPase-activating protein-binding protein 1
>gi|1037051683|ref|XP_017141274.1| PREDICTED: ras GTPase-activating protein-binding protein 1
>gi|1037059376|ref|XP_017141796.1| PREDICTED: RNA-binding protein squid isoform X1 [Drosophila
>gi|1037059405|ref|XP_017141798.1| PREDICTED: RNA-binding protein squid isoform X3 [Drosophila
>gi|1037059391|ref|XP_017141797.1| PREDICTED: RNA-binding protein squid isoform X2 [Drosophila
>gi|1037059362|ref|XP_017141795.1| PREDICTED: uncharacterized protein LOC108155482 [Drosophila
>gi|1037059421|ref|XP_017141799.1| PREDICTED: uncharacterized protein LOC108155484 [Drosophila
>gi|1037059437|ref|XP_017141800.1| PREDICTED: prisilkin-39 [Drosophila miranda]
>gi|1037059453|ref|XP_017141801.1| PREDICTED: keratin, type II cytoskeletal 2 epidermal [Drosop
>gi|1037059469|ref|XP_017141802.1| PREDICTED: cold shock domain-containing protein 4 [Drosophil
>gi|1037023717|ref|XP_017139504.1| PREDICTED: uncharacterized protein LOC108153919 [Drosophila
>gi|1037059485|ref|XP_017141803.1| PREDICTED: uncharacterized protein LOC108155489 [Drosophila
>gi|1037023735|ref|XP_017139505.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037092615|ref|XP_017144133.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037092629|ref|XP_017144134.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037092643|ref|XP_017144135.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037092659|ref|XP_017144136.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037092673|ref|XP_017144137.1| PREDICTED: hemicentin-1 [Drosophila miranda]
>gi|1037034322|ref|XP_017140152.1| PREDICTED: uncharacterized protein LOC108154397 [Drosophila
>gi|1037073479|ref|XP_017142756.1| PREDICTED: uncharacterized protein LOC108156023 isoform X1
>gi|1037073545|ref|XP_017142760.1| PREDICTED: band 4.1-like protein 5 isoform X5 [Drosophila m
>gi|1037073511|ref|XP_017142758.1| PREDICTED: uncharacterized protein LOC108156023 isoform X3
>gi|1037073495|ref|XP_017142757.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 2
>gi|1037073561|ref|XP_017142761.1| PREDICTED: band 4.1-like protein 5 isoform X6 [Drosophila m
>gi|1037073529|ref|XP_017142759.1| PREDICTED: band 4.1-like protein 5 isoform X4 [Drosophila m
>gi|1037073579|ref|XP_017142762.1| PREDICTED: actin-87E [Drosophila miranda]
>gi|1037023753|ref|XP_017139506.1| PREDICTED: uncharacterized protein LOC108153921 [Drosophila
>gi|1037105156|ref|XP_017145316.1| PREDICTED: L-dopachrome tautomerase yellow-f2 [Drosophila m
>gi|1037099458|ref|XP_017144704.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1037099470|ref|XP_017144705.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1037093763|ref|XP_017144208.1| PREDICTED: microprocessor complex subunit DGCR8 [Drosophila
>gi|1037104842|ref|XP_017145271.1| PREDICTED: uncharacterized protein LOC108157638 [Drosophila
>gi|1037045059|ref|XP_017140826.1| PREDICTED: protein kinase C-binding protein 1 [Drosophila m
>gi|1037045177|ref|XP_017140833.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophi
>gi|1037045207|ref|XP_017140836.1| PREDICTED: uncharacterized protein LOC108154898 [Drosophila
>gi|1037023771|ref|XP_017139507.1| PREDICTED: uncharacterized protein LOC108153922 [Drosophila
>gi|1037045074|ref|XP_017140828.1| PREDICTED: protein tramtrack, alpha isoform isoform X1 [Dros
>gi|1037045088|ref|XP_017140829.1| PREDICTED: protein tramtrack, alpha isoform isoform X1 [Dros
>gi|1037045104|ref|XP_017140830.1| PREDICTED: protein tramtrack, alpha isoform isoform X1 [Dros
>gi|1037045121|ref|XP_017140831.1| PREDICTED: protein tramtrack, alpha isoform isoform X1 [Dros
>gi|1037045133|ref|XP_017140832.1| PREDICTED: protein tramtrack, beta isoform isoform X2 [Droso
>gi|1037045193|ref|XP_017140834.1| PREDICTED: cleavage stimulation factor subunit 1 [Drosophila
>gi|1037039590|ref|XP_017140460.1| PREDICTED: nucleolar protein 16 [Drosophila miranda]
>gi|1037039538|ref|XP_017140455.1| PREDICTED: elongation factor 1-alpha 2 [Drosophila miranda]
>gi|1037039620|ref|XP_017140462.1| PREDICTED: tubulin-specific chaperone A [Drosophila miranda]
>gi|1037039574|ref|XP_017140459.1| PREDICTED: uncharacterized protein LOC108154641 [Drosophila
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>gi|1037039606|ref|XP_017140461.1| PREDICTED: nucleoside diphosphate kinase [Drosophila miranda
>gi|1037039560|ref|XP_017140457.1| PREDICTED: deoxyhypusine hydroxylase [Drosophila miranda]
>gi|1037039476|ref|XP_017140451.1| PREDICTED: cell surface glycoprotein 1 isoform X1 [Drosophil
>gi|1037039490|ref|XP_017140452.1| PREDICTED: cell surface glycoprotein 1 isoform X2 [Drosophil
>gi|1037039550|ref|XP_017140456.1| PREDICTED: ectopic P granules protein 5 homolog [Drosophila
>gi|1037039506|ref|XP_017140453.1| PREDICTED: cell surface glycoprotein 1 isoform X3 [Drosophil
>gi|1037039522|ref|XP_017140454.1| PREDICTED: cell surface glycoprotein 1 isoform X4 [Drosophil
>gi|1037039428|ref|XP_017140447.1| PREDICTED: ectopic P granules protein 5 homolog [Drosophila
>gi|1037039460|ref|XP_017140450.1| PREDICTED: serine/threonine-protein kinase greatwall isoform
>gi|1037039444|ref|XP_017140448.1| PREDICTED: serine/threonine-protein kinase greatwall isoform
>gi|1037039635|ref|XP_017140463.1| PREDICTED: RE1-silencing transcription factor [Drosophila mi
>gi|1037039650|ref|XP_017140464.1| PREDICTED: RE1-silencing transcription factor [Drosophila mi
>gi|1037077701|ref|XP_017143045.1| PREDICTED: saccharopine dehydrogenase-like oxidoreductase [D
>gi|1037077729|ref|XP_017143047.1| PREDICTED: cysteine-rich DPF motif domain-containing protei
>gi|1037077715|ref|XP_017143046.1| PREDICTED: putative transferase CAF17 homolog, mitochondrial
>gi|1037077663|ref|XP_017143042.1| PREDICTED: autophagy-related protein 16-1 isoform X1 [Droso
>gi|1037077676|ref|XP_017143043.1| PREDICTED: autophagy-related protein 16-1 isoform X2 [Droso
>gi|1037077689|ref|XP_017143044.1| PREDICTED: autophagy-related protein 16 isoform X3 [Drosophi
>gi|1037094530|ref|XP_017144268.1| PREDICTED: RNA-binding protein 34 [Drosophila miranda]
>gi|1037094544|ref|XP_017144269.1| PREDICTED: uncharacterized protein LOC108156986 [Drosophila
>gi|1037074896|ref|XP_017142852.1| PREDICTED: zinc finger protein weckle [Drosophila miranda]
>gi|1037074858|ref|XP_017142850.1| PREDICTED: zinc finger protein weckle [Drosophila miranda]
>gi|1037074912|ref|XP_017142853.1| PREDICTED: N-alpha-acetyltransferase 20-like [Drosophila mi
>gi|1037074880|ref|XP_017142851.1| PREDICTED: spermatogenesis-defective protein 39 homolog [Dr
>gi|1037058744|ref|XP_017141752.1| PREDICTED: uncharacterized protein LOC108155447 [Drosophila
>gi|1037058758|ref|XP_017141753.1| PREDICTED: protein claret segregational [Drosophila miranda]
>gi|1037058774|ref|XP_017141754.1| PREDICTED: 3'(2'),5'-bisphosphate nucleotidase 1 [Drosophila
>gi|1037086919|ref|XP_017143727.1| PREDICTED: electron transfer flavoprotein subunit beta [Dros
>gi|1037086933|ref|XP_017143728.1| PREDICTED: electron transfer flavoprotein subunit beta [Dros
>gi|1037086947|ref|XP_017143729.1| PREDICTED: electron transfer flavoprotein subunit beta [Dros
>gi|1037086905|ref|XP_017143726.1| PREDICTED: caspase [Drosophila miranda]
>gi|1037078196|ref|XP_017143084.1| PREDICTED: cytochrome c-type heme lyase [Drosophila miranda]
>gi|1037078182|ref|XP_017143083.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1037078168|ref|XP_017143082.1| PREDICTED: transmembrane protein 181 [Drosophila miranda]
>gi|1037078154|ref|XP_017143081.1| PREDICTED: pre-mRNA-processing factor 17 [Drosophila miranda
>gi|1037078210|ref|XP_017143085.1| PREDICTED: uncharacterized protein LOC108156247 [Drosophila
>gi|1037101741|ref|XP_017144904.1| PREDICTED: fas-associated death domain protein [Drosophila m
>gi|1037092259|ref|XP_017144110.1| PREDICTED: probable ATP-dependent RNA helicase pitchoune [Dr
>gi|1037058593|ref|XP_017141742.1| PREDICTED: protein held out wings isoform X1 [Drosophila mir
>gi|1037058609|ref|XP_017141743.1| PREDICTED: protein held out wings isoform X1 [Drosophila mir
>gi|1037058623|ref|XP_017141744.1| PREDICTED: protein held out wings isoform X2 [Drosophila mir
>gi|1037058639|ref|XP_017141745.1| PREDICTED: protein held out wings isoform X3 [Drosophila mir
>gi|1037058655|ref|XP_017141746.1| PREDICTED: protein held out wings isoform X4 [Drosophila mir
>gi|1037058671|ref|XP_017141747.1| PREDICTED: protein held out wings isoform X5 [Drosophila mir
>gi|1037035047|ref|XP_017140194.1| PREDICTED: uncharacterized protein LOC108154435 isoform X1
>gi|1037035063|ref|XP_017140195.1| PREDICTED: uncharacterized protein LOC108154435 isoform X2
>gi|1037099662|ref|XP_017144722.1| PREDICTED: serine protease HTRA2, mitochondrial [Drosophila
>gi|1037053425|ref|XP_017141392.1| PREDICTED: uncharacterized protein LOC108155219 [Drosophila
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>gi|1037053372|ref|XP_017141387.1| PREDICTED: platelet binding protein GspB-like, partial [Dros
>gi|1037053385|ref|XP_017141389.1| PREDICTED: uncharacterized protein LOC108155216 [Drosophila
>gi|1037053401|ref|XP_017141390.1| PREDICTED: stAR-related lipid transfer protein 13-like [Dros
>gi|1037053413|ref|XP_017141391.1| PREDICTED: GTPase-activating protein CdGAPr-like, partial [D
>gi|1037030991|ref|XP_017139952.1| PREDICTED: uncharacterized protein LOC108154236 [Drosophila
>gi|1037033216|ref|XP_017140085.1| PREDICTED: male-specific protein scotti [Drosophila miranda]
>gi|1037049095|ref|XP_017141094.1| PREDICTED: dynein assembly factor 5, axonemal [Drosophila mi
>gi|1037049111|ref|XP_017141095.1| PREDICTED: dynein assembly factor 5, axonemal [Drosophila mi
>gi|1037049127|ref|XP_017141096.1| PREDICTED: dynein assembly factor 5, axonemal [Drosophila mi
>gi|1037049143|ref|XP_017141097.1| PREDICTED: dynein assembly factor 5, axonemal [Drosophila mi
>gi|1037049161|ref|XP_017141098.1| PREDICTED: dynein assembly factor 5, axonemal [Drosophila mi
>gi|1037049175|ref|XP_017141100.1| PREDICTED: dynein assembly factor 5, axonemal [Drosophila mi
>gi|1037049191|ref|XP_017141101.1| PREDICTED: homeobox protein abdominal-B [Drosophila miranda]
>gi|1037049209|ref|XP_017141102.1| PREDICTED: homeobox protein abdominal-B [Drosophila miranda]
>gi|1037049227|ref|XP_017141103.1| PREDICTED: homeobox protein abdominal-B [Drosophila miranda]
>gi|1037049255|ref|XP_017141105.1| PREDICTED: DNA repair protein XRCC3 [Drosophila miranda]
>gi|1037049268|ref|XP_017141106.1| PREDICTED: ATP synthase subunit e, mitochondrial [Drosophila
>gi|1037049059|ref|XP_017141092.1| PREDICTED: uncharacterized protein LOC108155038 isoform X1
>gi|1037049079|ref|XP_017141093.1| PREDICTED: actin cytoskeleton-regulatory complex protein PAN
>gi|1037049239|ref|XP_017141104.1| PREDICTED: histone-lysine N-methyltransferase pr-set7 [Droso
>gi|1037056330|ref|XP_017141586.1| PREDICTED: bromodomain adjacent to zinc finger domain prote
>gi|1037056364|ref|XP_017141589.1| PREDICTED: DDB1- and CUL4-associated factor 5 isoform X2 [D
>gi|1037056346|ref|XP_017141588.1| PREDICTED: DDB1- and CUL4-associated factor 5 isoform X1 [D
>gi|1037056396|ref|XP_017141591.1| PREDICTED: uncharacterized protein LOC108155344 [Drosophila
>gi|1037056380|ref|XP_017141590.1| PREDICTED: DDB1- and CUL4-associated factor 5 isoform X3 [D
>gi|1037056446|ref|XP_017141594.1| PREDICTED: uncharacterized protein LOC108155347 [Drosophila
>gi|1037056412|ref|XP_017141592.1| PREDICTED: solute carrier family 52, riboflavin transporter
>gi|1037056430|ref|XP_017141593.1| PREDICTED: ferrochelatase, mitochondrial [Drosophila mirand
>gi|1037035411|ref|XP_017140212.1| PREDICTED: E3 ubiquitin-protein ligase msl-2-like isoform X
>gi|1037035395|ref|XP_017140211.1| PREDICTED: E3 ubiquitin-protein ligase msl-2-like isoform X
>gi|1037099293|ref|XP_017144686.1| PREDICTED: rho GTPase-activating protein 100F isoform X1 [D
>gi|1037099302|ref|XP_017144687.1| PREDICTED: rho GTPase-activating protein 100F isoform X2 [D
>gi|1037099338|ref|XP_017144692.1| PREDICTED: rho GTPase-activating protein 100F isoform X5 [D
>gi|1037099350|ref|XP_017144693.1| PREDICTED: rho GTPase-activating protein 100F isoform X6 [D
>gi|1037099314|ref|XP_017144688.1| PREDICTED: rho GTPase-activating protein 100F isoform X3 [D
>gi|1037099326|ref|XP_017144689.1| PREDICTED: rho GTPase-activating protein 100F isoform X4 [D
>gi|1037099362|ref|XP_017144694.1| PREDICTED: rho GTPase-activating protein 100F isoform X7 [D
>gi|1037034514|ref|XP_017140164.1| PREDICTED: uncharacterized protein LOC108154405 [Drosophila
>gi|1037102691|ref|XP_017145001.1| PREDICTED: uncharacterized protein LOC108157451 [Drosophila
>gi|1037102700|ref|XP_017145002.1| PREDICTED: uncharacterized protein LOC108157452 isoform X1
>gi|1037102711|ref|XP_017145003.1| PREDICTED: uncharacterized protein LOC108157452 isoform X2
>gi|1037102722|ref|XP_017145004.1| PREDICTED: uncharacterized protein LOC108157452 isoform X3
>gi|1037102731|ref|XP_017145005.1| PREDICTED: uncharacterized protein LOC108157452 isoform X4
>gi|1037102682|ref|XP_017145000.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1037102672|ref|XP_017144999.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1037103675|ref|XP_017145113.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037103685|ref|XP_017145115.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037106354|ref|XP_017145484.1| PREDICTED: farnesol dehydrogenase-like [Drosophila miranda]
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>gi|1037032688|ref|XP_017140053.1| PREDICTED: farnesol dehydrogenase-like [Drosophila miranda]
>gi|1037065741|ref|XP_017142223.1| PREDICTED: farnesol dehydrogenase-like [Drosophila miranda]
>gi|1037065625|ref|XP_017142216.1| PREDICTED: protein dispatched isoform X1 [Drosophila miranda
>gi|1037065641|ref|XP_017142217.1| PREDICTED: protein dispatched isoform X2 [Drosophila miranda
>gi|1037065725|ref|XP_017142222.1| PREDICTED: evolutionarily conserved signaling intermediate
>gi|1037065659|ref|XP_017142218.1| PREDICTED: GTP-binding protein 2 [Drosophila miranda]
>gi|1037065675|ref|XP_017142219.1| PREDICTED: GTP-binding protein 2 [Drosophila miranda]
>gi|1037065693|ref|XP_017142220.1| PREDICTED: GTP-binding protein 2 [Drosophila miranda]
>gi|1037065709|ref|XP_017142221.1| PREDICTED: GTP-binding protein 2 [Drosophila miranda]
>gi|1037097145|ref|XP_017144488.1| PREDICTED: uncharacterized protein LOC108157117 [Drosophila
>gi|1037032205|ref|XP_017140022.1| PREDICTED: general odorant-binding protein 83a [Drosophila m
>gi|1037032227|ref|XP_017140023.1| PREDICTED: general odorant-binding protein 83a [Drosophila m
>gi|1037032603|ref|XP_017140047.1| PREDICTED: pheromone-binding protein-related protein 6 [Dros
>gi|1037034696|ref|XP_017140176.1| PREDICTED: akirin-like [Drosophila miranda]
>gi|1037075991|ref|XP_017142930.1| PREDICTED: ATP-dependent RNA helicase p62 isoform X2 [Drosop
>gi|1037075976|ref|XP_017142929.1| PREDICTED: ATP-dependent RNA helicase p62 isoform X1 [Drosop
>gi|1037076034|ref|XP_017142932.1| PREDICTED: probable small nuclear ribonucleoprotein G [Drosc
>gi|1037076022|ref|XP_017142931.1| PREDICTED: low choriolytic enzyme [Drosophila miranda]
>gi|1037106197|ref|XP_017145464.1| PREDICTED: uncharacterized protein LOC108157785 [Drosophila
>gi|1037023789|ref|XP_017139508.1| PREDICTED: putative odorant receptor 83c [Drosophila miranda
>gi|1037101567|ref|XP_017144885.1| PREDICTED: uncharacterized protein LOC108157370 isoform X2
>gi|1037101558|ref|XP_017144884.1| PREDICTED: uncharacterized protein LOC108157370 isoform X1
>gi|1037101605|ref|XP_017144890.1| PREDICTED: uncharacterized protein LOC108157373 [Drosophila
>gi|1037101595|ref|XP_017144889.1| PREDICTED: uncharacterized protein LOC108157371 isoform X3
>gi|1037101577|ref|XP_017144887.1| PREDICTED: uncharacterized protein LOC108157371 isoform X1
>gi|1037101587|ref|XP_017144888.1| PREDICTED: general odorant-binding protein 99b isoform X2 [[
>gi|1037036759|ref|XP_017140286.1| PREDICTED: V-type proton ATPase subunit F [Drosophila miranc
>gi|1037038993|ref|XP_017140421.1| PREDICTED: glutaredoxin domain-containing cysteine-rich prot
>gi|1037033833|ref|XP_017140121.1| PREDICTED: uncharacterized protein LOC108154367 [Drosophila
>gi|1037039696|ref|XP_017140468.1| PREDICTED: uncharacterized protein LOC108154648 [Drosophila
>gi|1037101901|ref|XP_017144921.1| PREDICTED: uncharacterized protein LOC108157391 [Drosophila
>gi|1037101861|ref|XP_017144917.1| PREDICTED: neuropeptide F receptor isoform X1 [Drosophila mi
>gi|1037101869|ref|XP_017144918.1| PREDICTED: neuropeptide F receptor isoform X1 [Drosophila mi
>gi|1037101889|ref|XP_017144920.1| PREDICTED: neuropeptide F receptor isoform X3 [Drosophila mi
>gi|1037101879|ref|XP_017144919.1| PREDICTED: neuropeptide F receptor isoform X2 [Drosophila mi
>gi|1037101851|ref|XP_017144915.1| PREDICTED: uncharacterized protein LOC108157389 [Drosophila
>gi|1037037842|ref|XP_017140348.1| PREDICTED: uncharacterized protein LOC108154559 [Drosophila
>gi|1037023807|ref|XP_017139509.1| PREDICTED: uncharacterized protein LOC108153925 [Drosophila
>gi|1037037031|ref|XP_017140302.1| PREDICTED: uncharacterized protein LOC108154513 [Drosophila
>gi|1037039990|ref|XP_017140489.1| PREDICTED: uncharacterized protein LOC108154663 isoform X1
>gi|1037040006|ref|XP_017140490.1| PREDICTED: uncharacterized protein LOC108154663 isoform X2
>gi|1037038143|ref|XP_017140367.1| PREDICTED: keratin, type II cytoskeletal 2 oral [Drosophila
>gi|1037038219|ref|XP_017140372.1| PREDICTED: uncharacterized protein LOC108154581 [Drosophila
>gi|1037040022|ref|XP_017140491.1| PREDICTED: uncharacterized protein LOC108154664 [Drosophila
>gi|1037106821|ref|XP_017145551.1| PREDICTED: uncharacterized protein LOC108157856 [Drosophila
>gi|1037023824|ref|XP_017139510.1| PREDICTED: uncharacterized protein LOC108153926, partial [Di
>gi|1037023852|ref|XP_017139512.1| PREDICTED: uncharacterized protein LOC108153927 [Drosophila
>gi|1037023870|ref|XP_017139513.1| PREDICTED: uncharacterized protein LOC108153928 [Drosophila
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>gi|1037036922|ref|XP_017140296.1| PREDICTED: acanthoscurrin-1 [Drosophila miranda]
>gi|1037023886|ref|XP_017139514.1| PREDICTED: uncharacterized protein LOC108153929 [Drosophila
>gi|1037040843|ref|XP_017140547.1| PREDICTED: uncharacterized protein LOC108154708 [Drosophila
>gi|1037036777|ref|XP_017140287.1| PREDICTED: uncharacterized protein LOC108154498 [Drosophila
>gi|1037039712|ref|XP_017140469.1| PREDICTED: uncharacterized protein LOC108154649 [Drosophila
>gi|1037023904|ref|XP_017139515.1| PREDICTED: uncharacterized protein LOC108153930 [Drosophila
>gi|1037023922|ref|XP_017139516.1| PREDICTED: uncharacterized protein LOC108153931 [Drosophila
>gi|1037023938|ref|XP_017139517.1| PREDICTED: uncharacterized protein LOC108153932 [Drosophila
>gi|1037105602|ref|XP_017145381.1| PREDICTED: uncharacterized protein LOC108157723 isoform X1
>gi|1037105610|ref|XP_017145382.1| PREDICTED: uncharacterized protein LOC108157723 isoform X2
>gi|1037039666|ref|XP_017140465.1| PREDICTED: uncharacterized protein LOC108154646 [Drosophila
>gi|1037023956|ref|XP_017139519.1| PREDICTED: uncharacterized protein LOC108153933 [Drosophila
>gi|1037099393|ref|XP_017144698.1| PREDICTED: uncharacterized protein LOC108157239 [Drosophila
>gi|1037099374|ref|XP_017144696.1| PREDICTED: uncharacterized protein LOC108157238 [Drosophila
>gi|1037099385|ref|XP_017144697.1| PREDICTED: uncharacterized protein LOC108157238 [Drosophila
>gi|1037099403|ref|XP_017144699.1| PREDICTED: protein D2 [Drosophila miranda]
>gi|1037099415|ref|XP_017144700.1| PREDICTED: probable small nuclear ribonucleoprotein Sm D2 [
>gi|1037098088|ref|XP_017144577.1| PREDICTED: tRNA:m(4)X modification enzyme TRM13 homolog [Dro
>gi|1037089373|ref|XP_017143886.1| PREDICTED: vesicle-fusing ATPase 2 [Drosophila miranda]
>gi|1037099708|ref|XP_017144727.1| PREDICTED: peptide tarsal-less AA [Drosophila miranda]
>gi|1037077288|ref|XP_017143013.1| PREDICTED: putative aminopeptidase W07G4.4 [Drosophila mira
>gi|1037077303|ref|XP_017143014.1| PREDICTED: putative aminopeptidase W07G4.4 [Drosophila mira
>gi|1037077318|ref|XP_017143016.1| PREDICTED: putative aminopeptidase W07G4.4 [Drosophila mira
>gi|1037077332|ref|XP_017143017.1| PREDICTED: protein BCCIP homolog [Drosophila miranda]
>gi|1037077360|ref|XP_017143019.1| PREDICTED: LOW QUALITY PROTEIN: mediator of RNA polymerase
>gi|1037077374|ref|XP_017143020.1| PREDICTED: COX assembly mitochondrial protein 2 homolog [Dr
>gi|1037077346|ref|XP_017143018.1| PREDICTED: FAM206 family protein CG9288 [Drosophila miranda]
>gi|1037091912|ref|XP_017144084.1| PREDICTED: LOW QUALITY PROTEIN: protein fork head [Drosophil
>gi|1037096719|ref|XP_017144449.1| PREDICTED: zinc finger protein 330 homolog [Drosophila mira
>gi|1037105100|ref|XP_017145308.1| PREDICTED: uncharacterized protein LOC108157661 [Drosophila
>gi|1037031563|ref|XP_017139988.1| PREDICTED: uncharacterized protein LOC108154263 [Drosophila
>gi|1037070486|ref|XP_017142549.1| PREDICTED: signal transducer and transcription activator is
>gi|1037070502|ref|XP_017142550.1| PREDICTED: signal transducer and transcription activator is
>gi|1037070534|ref|XP_017142552.1| PREDICTED: signal transducer and transcription activator is
>gi|1037070518|ref|XP_017142551.1| PREDICTED: signal transducer and transcription activator is
>gi|1037070550|ref|XP_017142553.1| PREDICTED: signal transducer and transcription activator is
>gi|1037070565|ref|XP_017142554.1| PREDICTED: signal transducer and transcription activator is
>gi|1037070581|ref|XP_017142556.1| PREDICTED: metallothionein-4 [Drosophila miranda]
>gi|1037070597|ref|XP_017142557.1| PREDICTED: metallothionein-2 [Drosophila miranda]
>gi|1037104776|ref|XP_017145261.1| PREDICTED: vesicular glutamate transporter 1 [Drosophila mi
>gi|1037038157|ref|XP_017140368.1| PREDICTED: metallothionein-2 [Drosophila miranda]
>gi|1037106338|ref|XP_017145482.1| PREDICTED: uncharacterized protein LOC108157801 [Drosophila
>gi|1037106332|ref|XP_017145481.1| PREDICTED: mitochondrial dicarboxylate carrier [Drosophila
>gi|1037096053|ref|XP_017144389.1| PREDICTED: uncharacterized protein LOC108157065 [Drosophila
>gi|1037096062|ref|XP_017144390.1| PREDICTED: serine/threonine-protein phosphatase 2A regulato
>gi|1037096084|ref|XP_017144393.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037096072|ref|XP_017144391.1| PREDICTED: trimethyllysine dioxygenase, mitochondrial [Dros
>gi|1037036834|ref|XP_017140290.1| PREDICTED: metallothionein-3-like [Drosophila miranda]
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>gi|1037023970|ref|XP_017139520.1| PREDICTED: uncharacterized protein LOC108153934 [Drosophila
>gi|1037057274|ref|XP_017141649.1| PREDICTED: NAD-dependent protein deacetylase Sirt2 [Drosophi
>gi|1037057243|ref|XP_017141647.1| PREDICTED: zinc finger protein 836 isoform X1 [Drosophila mi
>gi|1037057259|ref|XP_017141648.1| PREDICTED: zinc finger protein 501 isoform X2 [Drosophila mi
>gi|1037057306|ref|XP_017141651.1| PREDICTED: oxysterol-binding protein-related protein 8 isofo
>gi|1037057290|ref|XP_017141650.1| PREDICTED: oxysterol-binding protein-related protein 8 isofo
>gi|1037057322|ref|XP_017141653.1| PREDICTED: oxysterol-binding protein-related protein 8 isofo
>gi|1037057338|ref|XP_017141654.1| PREDICTED: oxysterol-binding protein-related protein 8 isofo
>gi|1037057354|ref|XP_017141655.1| PREDICTED: oxysterol-binding protein-related protein 8 isofo
>gi|1037057370|ref|XP_017141656.1| PREDICTED: uncharacterized protein LOC108155387 [Drosophila
>gi|1037057386|ref|XP_017141657.1| PREDICTED: thioredoxin domain-containing protein 9 [Drosophi
>gi|1037078977|ref|XP_017143137.1| PREDICTED: 205 kDa microtubule-associated protein [Drosophil
>gi|1037094049|ref|XP_017144231.1| PREDICTED: DNA-binding protein modulo [Drosophila miranda]
>gi|1037088773|ref|XP_017143839.1| PREDICTED: beta-arrestin-1 [Drosophila miranda]
>gi|1037023988|ref|XP_017139521.1| PREDICTED: uncharacterized protein LOC108153935 [Drosophila
>gi|1037100832|ref|XP_017144823.1| PREDICTED: uncharacterized protein LOC108157319 [Drosophila
>gi|1037100920|ref|XP_017144833.1| PREDICTED: uncharacterized protein LOC108157327 [Drosophila
>gi|1037100910|ref|XP_017144832.1| PREDICTED: cell differentiation protein RCD1 homolog [Drosop
>gi|1037101271|ref|XP_017144863.1| PREDICTED: protein strawberry notch-like isoform X2 [Drosoph
>gi|1037101261|ref|XP_017144862.1| PREDICTED: protein strawberry notch-like isoform X1 [Drosoph
>gi|1037032263|ref|XP_017140025.1| PREDICTED: uncharacterized protein CG13380-like [Drosophila
>gi|1037035923|ref|XP_017140238.1| PREDICTED: uncharacterized protein LOC108154475 [Drosophila
>gi|1037048847|ref|XP_017141077.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1037048859|ref|XP_017141078.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1037048875|ref|XP_017141079.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1037048780|ref|XP_017141072.1| PREDICTED: polypyrimidine tract-binding protein 1 isoform X2
>gi|1037048800|ref|XP_017141073.1| PREDICTED: polypyrimidine tract-binding protein 1 isoform X2
>gi|1037048764|ref|XP_017141071.1| PREDICTED: polypyrimidine tract-binding protein 1 isoform X1
>gi|1037048816|ref|XP_017141074.1| PREDICTED: polypyrimidine tract-binding protein 1 isoform X3
>gi|1037048832|ref|XP_017141075.1| PREDICTED: polypyrimidine tract-binding protein 1 isoform X4
>gi|1037067691|ref|XP_017142353.1| PREDICTED: uncharacterized protein LOC108155793 isoform X2
>gi|1037067677|ref|XP_017142352.1| PREDICTED: uncharacterized protein LOC108155793 isoform X1
>gi|1037067661|ref|XP_017142351.1| PREDICTED: telomerase-binding protein EST1A [Drosophila mira
>gi|1037061625|ref|XP_017141952.1| PREDICTED: WD repeat-containing protein 91 [Drosophila miran
>gi|1037061593|ref|XP_017141949.1| PREDICTED: uncharacterized protein LOC108155567 isoform X2
>gi|1037061609|ref|XP_017141950.1| PREDICTED: uncharacterized protein LOC108155567 isoform X3
>gi|1037061568|ref|XP_017141948.1| PREDICTED: uncharacterized protein LOC108155567 isoform X1
>gi|1037081655|ref|XP_017143330.1| PREDICTED: U1 small nuclear ribonucleoprotein C [Drosophila
>gi|1037081641|ref|XP_017143329.1| PREDICTED: deoxynucleoside kinase [Drosophila miranda]
>gi|1037081627|ref|XP_017143328.1| PREDICTED: WD40 repeat-containing protein SMU1 [Drosophila m
>gi|1037081669|ref|XP_017143332.1| PREDICTED: uncharacterized protein LOC108156404 isoform X1
>gi|1037081685|ref|XP_017143333.1| PREDICTED: uncharacterized protein LOC108156404 isoform X2
>gi|1037097980|ref|XP_017144568.1| PREDICTED: tolloid-like protein 2 isoform X2 [Drosophila mi
>gi|1037097958|ref|XP_017144566.1| PREDICTED: tolloid-like protein 2 isoform X1 [Drosophila mi
>gi|1037097969|ref|XP_017144567.1| PREDICTED: tolloid-like protein 2 isoform X1 [Drosophila mi
>gi|1037098051|ref|XP_017144573.1| PREDICTED: tolloid-like protein 2 isoform X6 [Drosophila mi
>gi|1037097991|ref|XP_017144569.1| PREDICTED: tolloid-like protein 2 isoform X3 [Drosophila mi
>gi|1037098000|ref|XP_017144570.1| PREDICTED: tolloid-like protein 2 isoform X4 [Drosophila mi
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>gi|1037098031|ref|XP_017144571.1| PREDICTED: tolloid-like protein 2 isoform X5 [Drosophila mi
>gi|1037098061|ref|XP_017144574.1| PREDICTED: uncharacterized protein LOC108157167 [Drosophila
>gi|1037098069|ref|XP_017144575.1| PREDICTED: uncharacterized protein LOC108157167 [Drosophila
>gi|1037086044|ref|XP_017143659.1| PREDICTED: mitochondrial pyruvate carrier 1 [Drosophila mira
>gi|1037086030|ref|XP_017143658.1| PREDICTED: uncharacterized protein LOC108156615 [Drosophila
>gi|1037102993|ref|XP_017145033.1| PREDICTED: N-sulphoglucosamine sulphohydrolase [Drosophila m
>gi|1037094265|ref|XP_017144247.1| PREDICTED: endophilin-A [Drosophila miranda]
>gi|1037094279|ref|XP_017144248.1| PREDICTED: endophilin-A [Drosophila miranda]
>gi|1037094293|ref|XP_017144249.1| PREDICTED: endophilin-A [Drosophila miranda]
>gi|1037098208|ref|XP_017144592.1| PREDICTED: uncharacterized protein LOC108157180 [Drosophila
>gi|1037098191|ref|XP_017144590.1| PREDICTED: uncharacterized protein LOC108157178 [Drosophila
>gi|1037098199|ref|XP_017144591.1| PREDICTED: uncharacterized protein LOC108157179 [Drosophila
>gi|1037098174|ref|XP_017144587.1| PREDICTED: opsin Rh2 [Drosophila miranda]
>gi|1037098181|ref|XP_017144588.1| PREDICTED: low molecular weight phosphotyrosine protein phos
>gi|1037036602|ref|XP_017140278.1| PREDICTED: kunitz-type serine protease inhibitor Hg1 [Drosop
>gi|1037105672|ref|XP_017145390.1| PREDICTED: alpha-2Db adrenergic receptor isoform X1 [Drosoph
>gi|1037105680|ref|XP_017145392.1| PREDICTED: alpha-2Db adrenergic receptor isoform X2 [Drosoph
>gi|1037105688|ref|XP_017145393.1| PREDICTED: alpha-2Db adrenergic receptor isoform X3 [Drosoph
>gi|1037100173|ref|XP_017144759.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1037100185|ref|XP_017144760.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1037100197|ref|XP_017144761.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1037103693|ref|XP_017145116.1| PREDICTED: sodium-coupled monocarboxylate transporter 2 [Dro
>gi|1037097723|ref|XP_017144542.1| PREDICTED: sodium-coupled monocarboxylate transporter 2-like
>gi|1037097734|ref|XP_017144543.1| PREDICTED: sodium-coupled monocarboxylate transporter 2-like
>gi|1037097744|ref|XP_017144544.1| PREDICTED: sodium-coupled monocarboxylate transporter 2-like
>gi|1037097756|ref|XP_017144545.1| PREDICTED: sodium-coupled monocarboxylate transporter 2 [Dro
>gi|1037097774|ref|XP_017144547.1| PREDICTED: putative glycine-rich cell wall structural prote
>gi|1037097764|ref|XP_017144546.1| PREDICTED: glycine-rich cell wall structural protein 2 isof
>gi|1037034238|ref|XP_017140146.1| PREDICTED: cytochrome c oxidase subunit 7A-related protein,
>gi|1037095925|ref|XP_017144376.1| PREDICTED: uncharacterized protein F21D5.5 [Drosophila mira
>gi|1037095937|ref|XP_017144377.1| PREDICTED: ADP-ribosylation factor-like protein 2 [Drosophi
>gi|1037101956|ref|XP_017144926.1| PREDICTED: cytochrome c oxidase subunit 7A, mitochondrial [
>gi|1037067964|ref|XP_017142373.1| PREDICTED: paired box pox-meso protein [Drosophila miranda]
>gi|1037068036|ref|XP_017142378.1| PREDICTED: actin cytoskeleton-regulatory complex protein pa
>gi|1037068022|ref|XP_017142377.1| PREDICTED: high affinity copper uptake protein 1 [Drosophila
>gi|1037067949|ref|XP_017142372.1| PREDICTED: 4-hydroxybenzoate polyprenyltransferase, mitochon
>gi|1037067935|ref|XP_017142371.1| PREDICTED: eukaryotic initiation factor 4A-III [Drosophila m
>gi|1037068008|ref|XP_017142376.1| PREDICTED: uncharacterized protein LOC108155807 [Drosophila
>gi|1037067994|ref|XP_017142375.1| PREDICTED: uncharacterized protein LOC108155805 [Drosophila
>gi|1037068052|ref|XP_017142380.1| PREDICTED: basic-leucine zipper transcription factor A [Dros
>gi|1037067980|ref|XP_017142374.1| PREDICTED: THO complex subunit 3 [Drosophila miranda]
>gi|1037024004|ref|XP_017139522.1| PREDICTED: protein SGT1 homolog [Drosophila miranda]
>gi|1037084346|ref|XP_017143535.1| PREDICTED: protein SGT1 homolog [Drosophila miranda]
>gi|1037084333|ref|XP_017143534.1| PREDICTED: GDP-fucose transporter 1 [Drosophila miranda]
>gi|1037084319|ref|XP_017143533.1| PREDICTED: 39S ribosomal protein L1, mitochondrial [Drosophi
>gi|1037096005|ref|XP_017144384.1| PREDICTED: uncharacterized protein LOC108157063 isoform X1
>gi|1037096015|ref|XP_017144385.1| PREDICTED: uncharacterized protein LOC108157063 isoform X1
>gi|1037096035|ref|XP_017144387.1| PREDICTED: uncharacterized protein LOC108157063 isoform X3
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>gi|1037096025|ref|XP_017144386.1| PREDICTED: uncharacterized protein LOC108157063 isoform X2
>gi|1037096043|ref|XP_017144388.1| PREDICTED: uncharacterized protein LOC108157064 [Drosophila
>gi|1037034482|ref|XP_017140162.1| PREDICTED: uncharacterized protein LOC108154403 [Drosophila
>gi|1037033378|ref|XP_017140093.1| PREDICTED: uncharacterized protein LOC108154342 [Drosophila
>gi|1037032396|ref|XP_017140033.1| PREDICTED: protein C19orf12 homolog [Drosophila miranda]
>gi|1037031510|ref|XP_017139984.1| PREDICTED: protein atonal [Drosophila miranda]
>gi|1037047436|ref|XP_017140985.1| PREDICTED: phenylalanine--tRNA ligase beta subunit [Drosoph
>gi|1037047468|ref|XP_017140988.1| PREDICTED: ankyrin repeat and KH domain-containing protein r
>gi|1037047452|ref|XP_017140986.1| PREDICTED: ankyrin repeat and KH domain-containing protein r
>gi|1037090183|ref|XP_017143953.1| PREDICTED: signal peptidase complex subunit 3 [Drosophila mi
>gi|1037090169|ref|XP_017143951.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1037040716|ref|XP_017140540.1| PREDICTED: transcription factor Sp4 [Drosophila miranda]
>gi|1037040785|ref|XP_017140545.1| PREDICTED: telomere-binding protein cav [Drosophila miranda]
>gi|1037040748|ref|XP_017140542.1| PREDICTED: CCR4-NOT transcription complex subunit 6 [Drosoph
>gi|1037040762|ref|XP_017140543.1| PREDICTED: CCR4-NOT transcription complex subunit 6 [Drosoph
>gi|1037040732|ref|XP_017140541.1| PREDICTED: uncharacterized protein LOC108154702 [Drosophila
>gi|1037024024|ref|XP_017139523.1| PREDICTED: LOW QUALITY PROTEIN: NFX1-type zinc finger-contai
>gi|1037040696|ref|XP_017140539.1| PREDICTED: LOW QUALITY PROTEIN: structural maintenance of ch
>gi|1037040801|ref|XP_017140546.1| PREDICTED: RWD domain-containing protein 1 [Drosophila miran
>gi|1037040770|ref|XP_017140544.1| PREDICTED: cysteine and histidine-rich domain-containing pro
>gi|1037103840|ref|XP_017145134.1| PREDICTED: uncharacterized protein LOC108157542 isoform X2
>gi|1037103832|ref|XP_017145133.1| PREDICTED: uncharacterized protein LOC108157542 isoform X1
>gi|1037070628|ref|XP_017142559.1| PREDICTED: vesicular integral-membrane protein VIP36 [Droso
>gi|1037070613|ref|XP_017142558.1| PREDICTED: adenomatous polyposis coli protein [Drosophila mi
>gi|1037070660|ref|XP_017142561.1| PREDICTED: 28S ribosomal protein S24, mitochondrial [Drosoph
>gi|1037070644|ref|XP_017142560.1| PREDICTED: TBC1 domain family member 7 [Drosophila miranda]
>gi|1037080758|ref|XP_017143261.1| PREDICTED: unconventional myosin-Ib isoform X1 [Drosophila m
>gi|1037080774|ref|XP_017143262.1| PREDICTED: unconventional myosin-Ia isoform X2 [Drosophila m
>gi|1037080788|ref|XP_017143263.1| PREDICTED: uncharacterized protein LOC108156368 isoform X3
>gi|1037080802|ref|XP_017143264.1| PREDICTED: 4-coumarate--CoA ligase 1 [Drosophila miranda]
>gi|1037080816|ref|XP_017143265.1| PREDICTED: 4-coumarate--CoA ligase 1 [Drosophila miranda]
>gi|1037080830|ref|XP_017143267.1| PREDICTED: 4-coumarate--CoA ligase 1 [Drosophila miranda]
>gi|1037099590|ref|XP_017144716.1| PREDICTED: mitogen-activated protein kinase 14A [Drosophila
>gi|1037099578|ref|XP_017144715.1| PREDICTED: mitogen-activated protein kinase 14A-like [Drosop
>gi|1037096402|ref|XP_017144422.1| PREDICTED: serine--tRNA synthetase-like protein Slimp [Dros
>gi|1037096390|ref|XP_017144421.1| PREDICTED: anosmin-1 [Drosophila miranda]
>gi|1037088975|ref|XP_017143856.1| PREDICTED: epididymal secretory protein E1 isoform X2 [Dros
>gi|1037088963|ref|XP_017143855.1| PREDICTED: epididymal secretory protein E1 isoform X1 [Dros
>gi|1037089003|ref|XP_017143857.1| PREDICTED: epididymal secretory protein E1 isoform X3 [Dros
>gi|1037088949|ref|XP_017143854.1| PREDICTED: exocyst complex component 5 [Drosophila miranda]
>gi|1037024042|ref|XP_017139524.1| PREDICTED: hamartin, partial [Drosophila miranda]
>gi|1037086849|ref|XP_017143721.1| PREDICTED: glutamyl-tRNA(Gln) amidotransferase subunit B, m
>gi|1037086807|ref|XP_017143718.1| PREDICTED: rootletin [Drosophila miranda]
>gi|1037086821|ref|XP_017143719.1| PREDICTED: rootletin [Drosophila miranda]
>gi|1037086835|ref|XP_017143720.1| PREDICTED: uncharacterized protein LOC108156633 [Drosophila
>gi|1037093587|ref|XP_017144196.1| PREDICTED: uncharacterized protein LOC108156943 [Drosophila
>gi|1037093599|ref|XP_017144198.1| PREDICTED: ADP-ribosylation factor-like protein 6-interactin
>gi|1037045574|ref|XP_017140861.1| PREDICTED: eukaryotic translation initiation factor 4E type
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>gi|1037045522|ref|XP_017140857.1| PREDICTED: eukaryotic translation initiation factor 4E type
>gi|1037045538|ref|XP_017140858.1| PREDICTED: eukaryotic translation initiation factor 4E type
>gi|1037045588|ref|XP_017140862.1| PREDICTED: eukaryotic translation initiation factor 4E type
>gi|1037045490|ref|XP_017140855.1| PREDICTED: UV excision repair protein RAD23 homolog A [Droso
>gi|1037045558|ref|XP_017140859.1| PREDICTED: eukaryotic translation initiation factor 4E type
>gi|1037045506|ref|XP_017140856.1| PREDICTED: syntaxin-1A [Drosophila miranda]
>gi|1037045442|ref|XP_017140852.1| PREDICTED: structural maintenance of chromosomes protein 1A
>gi|1037045474|ref|XP_017140854.1| PREDICTED: heat shock protein 68 isoform X2 [Drosophila mira
>gi|1037045458|ref|XP_017140853.1| PREDICTED: heat shock protein 68 isoform X1 [Drosophila mira
>gi|1037044949|ref|XP_017140819.1| PREDICTED: heat shock protein 67B2 [Drosophila miranda]
>gi|1037044910|ref|XP_017140816.1| PREDICTED: phosphatidylserine decarboxylase proenzyme, mito
>gi|1037044924|ref|XP_017140817.1| PREDICTED: beclin-1-like protein [Drosophila miranda]
>gi|1037044934|ref|XP_017140818.1| PREDICTED: protein SDE2 homolog [Drosophila miranda]
>gi|1037044874|ref|XP_017140813.1| PREDICTED: nucleolysin TIAR isoform X1 [Drosophila miranda]
>gi|1037044890|ref|XP_017140814.1| PREDICTED: nucleolysin TIAR isoform X2 [Drosophila miranda]
>gi|1037044779|ref|XP_017140807.1| PREDICTED: spastin isoform X2 [Drosophila miranda]
>gi|1037044763|ref|XP_017140806.1| PREDICTED: spastin isoform X1 [Drosophila miranda]
>gi|1037044795|ref|XP_017140808.1| PREDICTED: spastin isoform X3 [Drosophila miranda]
>gi|1037044811|ref|XP_017140809.1| PREDICTED: mitochondrial Rho GTPase isoform X1 [Drosophila m
>gi|1037044827|ref|XP_017140810.1| PREDICTED: mitochondrial Rho GTPase isoform X2 [Drosophila m
>gi|1037044843|ref|XP_017140811.1| PREDICTED: mitochondrial Rho GTPase isoform X3 [Drosophila m
>gi|1037044859|ref|XP_017140812.1| PREDICTED: probable N6-adenosine-methyltransferase MT-A70-li
>gi|1037063468|ref|XP_017142080.1| PREDICTED: phosphofurin acidic cluster sorting protein 2 iso
>gi|1037063482|ref|XP_017142081.1| PREDICTED: phosphofurin acidic cluster sorting protein 2 iso
>gi|1037063498|ref|XP_017142082.1| PREDICTED: UPF0692 protein CG33108 [Drosophila miranda]
>gi|1037063545|ref|XP_017142085.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm3 [Drosoph
>gi|1037063513|ref|XP_017142083.1| PREDICTED: ras-related protein Rab-7a [Drosophila miranda]
>gi|1037063529|ref|XP_017142084.1| PREDICTED: ras-related protein Rab-7a [Drosophila miranda]
>gi|1037104499|ref|XP_017145222.1| PREDICTED: uncharacterized protein LOC108157605 [Drosophila
>gi|1037075639|ref|XP_017142905.1| PREDICTED: protein UBASH3A homolog isoform X1 [Drosophila mi
>gi|1037075655|ref|XP_017142906.1| PREDICTED: protein UBASH3A homolog isoform X2 [Drosophila mi
>gi|1037075671|ref|XP_017142907.1| PREDICTED: uncharacterized protein CG5902 [Drosophila mirand
>gi|1037075687|ref|XP_017142908.1| PREDICTED: uncharacterized protein CG5902 [Drosophila mirand
>gi|1037075701|ref|XP_017142910.1| PREDICTED: uncharacterized protein CG5902 [Drosophila mirand
>gi|1037075717|ref|XP_017142911.1| PREDICTED: uncharacterized protein CG5902 [Drosophila mirand
>gi|1037075733|ref|XP_017142912.1| PREDICTED: transmembrane protein 179 [Drosophila miranda]
>gi|1037090877|ref|XP_017144007.1| PREDICTED: diacylglycerol kinase theta isoform X2 [Drosophil
>gi|1037090863|ref|XP_017144006.1| PREDICTED: diacylglycerol kinase theta isoform X1 [Drosophil
>gi|1037093787|ref|XP_017144210.1| PREDICTED: AP-1 complex subunit sigma-2 isoform X1 [Drosophi
>gi|1037093815|ref|XP_017144212.1| PREDICTED: AP-1 complex subunit sigma-2 isoform X3 [Drosophi
>gi|1037093829|ref|XP_017144214.1| PREDICTED: AP-1 complex subunit sigma-2 isoform X3 [Drosophi
>gi|1037093843|ref|XP_017144215.1| PREDICTED: AP-1 complex subunit sigma-2 isoform X4 [Drosophi
>gi|1037093801|ref|XP_017144211.1| PREDICTED: AP-1 complex subunit sigma-2 isoform X2 [Drosophi
>gi|1037093856|ref|XP_017144216.1| PREDICTED: AP-1 complex subunit sigma-2 isoform X5 [Drosophi
>gi|1037076095|ref|XP_017142935.1| PREDICTED: bifunctional glutamate/proline--tRNA ligase [Dros
>gi|1037078239|ref|XP_017143088.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 isoform
>gi|1037078251|ref|XP_017143089.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 isoform
>gi|1037078277|ref|XP_017143091.1| PREDICTED: ras-related protein Ral-a-like [Drosophila mirand
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>gi|1037078225|ref|XP_017143086.1| PREDICTED: LOW QUALITY PROTEIN: glutamate dehydrogenase, mit
>gi|1037078265|ref|XP_017143090.1| PREDICTED: uncharacterized protein LOC108156251 [Drosophila
>gi|1037100648|ref|XP_017144802.1| PREDICTED: LOW QUALITY PROTEIN: 40S ribosomal protein S19b
>gi|1037100638|ref|XP_017144801.1| PREDICTED: uncharacterized protein LOC108157308 [Drosophila
>gi|1037072539|ref|XP_017142688.1| PREDICTED: 26S protease regulatory subunit 4 [Drosophila mi
>gi|1037072649|ref|XP_017142696.1| PREDICTED: uncharacterized protein YAE1 [Drosophila miranda]
>gi|1037072601|ref|XP_017142693.1| PREDICTED: probable DNA-directed RNA polymerase III subunit
>gi|1037072617|ref|XP_017142694.1| PREDICTED: probable DNA-directed RNA polymerase III subunit
>gi|1037072555|ref|XP_017142689.1| PREDICTED: selenide, water dikinase 2 [Drosophila miranda]
>gi|1037072585|ref|XP_017142692.1| PREDICTED: zinc finger protein-like 1 homolog [Drosophila mi
>gi|1037072569|ref|XP_017142690.1| PREDICTED: transmembrane protein 199 [Drosophila miranda]
>gi|1037072633|ref|XP_017142695.1| PREDICTED: biogenesis of lysosome-related organelles complex
>gi|1037085793|ref|XP_017143640.1| PREDICTED: bifunctional arginine demethylase and lysyl-hydro
>gi|1037085809|ref|XP_017143641.1| PREDICTED: GTP-binding protein SAR1b [Drosophila miranda]
>gi|1037079677|ref|XP_017143177.1| PREDICTED: uncharacterized protein LOC108156302 isoform X3
>gi|1037079663|ref|XP_017143176.1| PREDICTED: uncharacterized protein LOC108156302 isoform X2
>gi|1037079649|ref|XP_017143175.1| PREDICTED: uncharacterized protein LOC108156302 isoform X1
>gi|1037079705|ref|XP_017143179.1| PREDICTED: phosphoinositide 3-kinase adapter protein 1 isof
>gi|1037079691|ref|XP_017143178.1| PREDICTED: phosphoinositide 3-kinase adapter protein 1 isof
>gi|1037038076|ref|XP_017140361.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1037076583|ref|XP_017142965.1| PREDICTED: uncharacterized protein LOC108156152 [Drosophila
>gi|1037076615|ref|XP_017142967.1| PREDICTED: cystatin-like protein [Drosophila miranda]
>gi|1037076631|ref|XP_017142969.1| PREDICTED: cystatin-like protein [Drosophila miranda]
>gi|1037076599|ref|XP_017142966.1| PREDICTED: uncharacterized protein LOC108156153 [Drosophila
>gi|1037024060|ref|XP_017139525.1| PREDICTED: LOW QUALITY PROTEIN: bypass of stop codon protein
>gi|1037037955|ref|XP_017140355.1| PREDICTED: bypass of stop codon protein 1-like [Drosophila m
>gi|1037035941|ref|XP_017140239.1| PREDICTED: glyceraldehyde-3-phosphate dehydrogenase 2-like
>gi|1037034725|ref|XP_017140177.1| PREDICTED: uncharacterized protein LOC108154417 [Drosophila
>gi|1037095076|ref|XP_017144309.1| PREDICTED: LOW QUALITY PROTEIN: cGMP-specific 3',5'-cyclic p
>gi|1037095089|ref|XP_017144310.1| PREDICTED: putative uncharacterized protein DDB_G0271974 [Dr
>gi|1037037645|ref|XP_017140339.1| PREDICTED: protein new-glue 2 [Drosophila miranda]
>gi|1037098777|ref|XP_017144647.1| PREDICTED: protein PDF [Drosophila miranda]
>gi|1037098790|ref|XP_017144648.1| PREDICTED: ELMO domain-containing protein C-like [Drosophila
>gi|1037098766|ref|XP_017144646.1| PREDICTED: golgin subfamily A member 7 [Drosophila miranda]
>gi|1037098144|ref|XP_017144584.1| PREDICTED: elastase-1 isoform X2 [Drosophila miranda]
>gi|1037098134|ref|XP_017144583.1| PREDICTED: serine protease 44 isoform X1 [Drosophila miranda
>gi|1037098116|ref|XP_017144581.1| PREDICTED: organic cation transporter protein isoform X1 [Dr
>gi|1037098126|ref|XP_017144582.1| PREDICTED: organic cation transporter protein isoform X2 [Dr
>gi|1037089068|ref|XP_017143863.1| PREDICTED: uncharacterized protein LOC108156723 [Drosophila
>gi|1037089056|ref|XP_017143861.1| PREDICTED: organic cation transporter protein [Drosophila mi
>gi|1037089044|ref|XP_017143860.1| PREDICTED: organic cation transporter-like protein [Drosophi
>gi|1037079129|ref|XP_017143149.1| PREDICTED: myosin heavy chain 95F isoform X1 [Drosophila mir
>gi|1037079155|ref|XP_017143151.1| PREDICTED: myosin heavy chain 95F isoform X1 [Drosophila mir
>gi|1037079169|ref|XP_017143152.1| PREDICTED: myosin heavy chain 95F isoform X1 [Drosophila mir
>gi|1037079143|ref|XP_017143150.1| PREDICTED: myosin heavy chain 95F isoform X2 [Drosophila mir
>gi|1037070877|ref|XP_017142576.1| PREDICTED: probable ATP-dependent RNA helicase DDX55 homolog
>gi|1037070861|ref|XP_017142575.1| PREDICTED: DNA replication licensing factor Mcm2 [Drosophila
>gi|1037070925|ref|XP_017142579.1| PREDICTED: replication protein A 70 kDa DNA-binding subunit
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>gi|1037070907|ref|XP_017142578.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1037070891|ref|XP_017142577.1| PREDICTED: uncharacterized protein LOC108155932 isoform X1
>gi|1037097157|ref|XP_017144489.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isoform
>gi|1037097169|ref|XP_017144490.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isoform
>gi|1037097181|ref|XP_017144491.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Drosop
>gi|1037097193|ref|XP_017144492.1| PREDICTED: tether containing UBX domain for GLUT4 [Drosophil
>gi|1037099224|ref|XP_017144679.1| PREDICTED: methylcrotonoyl-CoA carboxylase subunit alpha, m
>gi|1037070845|ref|XP_017142574.1| PREDICTED: probable ubiquitin carboxyl-terminal hydrolase F
>gi|1037070829|ref|XP_017142573.1| PREDICTED: probable ubiquitin carboxyl-terminal hydrolase F
>gi|1037070813|ref|XP_017142572.1| PREDICTED: probable ubiquitin carboxyl-terminal hydrolase F
>gi|1037070781|ref|XP_017142570.1| PREDICTED: probable ubiquitin carboxyl-terminal hydrolase F
>gi|1037070797|ref|XP_017142571.1| PREDICTED: probable ubiquitin carboxyl-terminal hydrolase F
>gi|1037087815|ref|XP_017143765.1| PREDICTED: JNK1/MAPK8-associated membrane protein [Drosophil
>gi|1037087787|ref|XP_017143763.1| PREDICTED: E3 ubiquitin-protein ligase HRD1 [Drosophila mir
>gi|1037087801|ref|XP_017143764.1| PREDICTED: E3 ubiquitin-protein ligase HRD1 [Drosophila mir
>gi|1037082541|ref|XP_017143396.1| PREDICTED: dipeptidyl peptidase 3 isoform X1 [Drosophila mi
>gi|1037082555|ref|XP_017143397.1| PREDICTED: dipeptidyl peptidase 3 isoform X2 [Drosophila mi
>gi|1037082569|ref|XP_017143398.1| PREDICTED: trichohyalin-like [Drosophila miranda]
>gi|1037093774|ref|XP_017144209.1| PREDICTED: autophagy-related protein 13 homolog [Drosophila
>gi|1037024076|ref|XP_017139526.1| PREDICTED: uncharacterized protein LOC108153941 [Drosophila
>gi|1037076646|ref|XP_017142970.1| PREDICTED: LOW QUALITY PROTEIN: adenomatous polyposis coli
>gi|1037024094|ref|XP_017139527.1| PREDICTED: solute carrier family 41 member 2-like [Drosophil
>gi|1037091210|ref|XP_017144032.1| PREDICTED: T-complex protein 1 subunit alpha [Drosophila mi
>gi|1037091222|ref|XP_017144033.1| PREDICTED: zinc finger protein 853 [Drosophila miranda]
>gi|1037091236|ref|XP_017144034.1| PREDICTED: probable prefoldin subunit 5 [Drosophila miranda
>gi|1037057619|ref|XP_017141674.1| PREDICTED: neurexin-1 isoform X3 [Drosophila miranda]
>gi|1037057635|ref|XP_017141675.1| PREDICTED: neurexin-1 isoform X4 [Drosophila miranda]
>gi|1037057603|ref|XP_017141673.1| PREDICTED: neurexin-1 isoform X2 [Drosophila miranda]
>gi|1037057588|ref|XP_017141672.1| PREDICTED: neurexin-1 isoform X1 [Drosophila miranda]
>gi|1037057655|ref|XP_017141676.1| PREDICTED: neurexin-1 isoform X5 [Drosophila miranda]
>gi|1037057668|ref|XP_017141677.1| PREDICTED: neurexin-1 isoform X6 [Drosophila miranda]
>gi|1037057700|ref|XP_017141679.1| PREDICTED: neurexin-3b isoform X8 [Drosophila miranda]
>gi|1037057684|ref|XP_017141678.1| PREDICTED: neurexin-3b isoform X7 [Drosophila miranda]
>gi|1037057828|ref|XP_017141688.1| PREDICTED: valacyclovir hydrolase [Drosophila miranda]
>gi|1037057860|ref|XP_017141690.1| PREDICTED: protein D3 [Drosophila miranda]
>gi|1037057844|ref|XP_017141689.1| PREDICTED: protein D3 [Drosophila miranda]
>gi|1037057796|ref|XP_017141686.1| PREDICTED: AP-2 complex subunit mu [Drosophila miranda]
>gi|1037057812|ref|XP_017141687.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 6 [
>gi|1037057731|ref|XP_017141681.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1037057715|ref|XP_017141680.1| PREDICTED: broad-complex core protein isoform X1 [Drosophila
>gi|1037057763|ref|XP_017141683.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1037057747|ref|XP_017141682.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1037057778|ref|XP_017141685.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1037032297|ref|XP_017140028.1| PREDICTED: uncharacterized protein LOC108154288 [Drosophila
>gi|1037073779|ref|XP_017142776.1| PREDICTED: uncharacterized protein LOC108156035 [Drosophila
>gi|1037106119|ref|XP_017145453.1| PREDICTED: transcription factor SPT20 homolog isoform X1 [Di
>gi|1037106127|ref|XP_017145454.1| PREDICTED: transcription factor SPT20 homolog isoform X2 [Di
>gi|1037094402|ref|XP_017144258.1| PREDICTED: uncharacterized protein LOC108156977 [Drosophila
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>gi|1037094416|ref|XP_017144259.1| PREDICTED: MOB kinase activator-like 1 [Drosophila miranda]
>gi|1037089634|ref|XP_017143905.1| PREDICTED: LOW QUALITY PROTEIN: integrator complex subunit
>gi|1037065817|ref|XP_017142228.1| PREDICTED: nucleolar protein 56 [Drosophila miranda]
>gi|1037065771|ref|XP_017142225.1| PREDICTED: clathrin interactor 1 isoform X2 [Drosophila mira
>gi|1037065757|ref|XP_017142224.1| PREDICTED: telomere length regulation protein TEL2 homolog
>gi|1037065801|ref|XP_017142227.1| PREDICTED: protein TBRG4 [Drosophila miranda]
>gi|1037065785|ref|XP_017142226.1| PREDICTED: IQ and ubiquitin-like domain-containing protein
>gi|1037032500|ref|XP_017140041.1| PREDICTED: uncharacterized protein LOC108154299 isoform X1
>gi|1037032518|ref|XP_017140042.1| PREDICTED: uncharacterized protein LOC108154299 isoform X2
>gi|1037053035|ref|XP_017141365.1| PREDICTED: uncharacterized protein LOC108155202 [Drosophila
>gi|1037053078|ref|XP_017141368.1| PREDICTED: phosphoglycerate mutase 2 isoform X3 [Drosophila
>gi|1037053065|ref|XP_017141367.1| PREDICTED: phosphoglycerate mutase 2 isoform X2 [Drosophila
>gi|1037053050|ref|XP_017141366.1| PREDICTED: phosphoglycerate mutase 2 isoform X1 [Drosophila
>gi|1037053087|ref|XP_017141369.1| PREDICTED: phosphoglycerate mutase 2 isoform X4 [Drosophila
>gi|1037053102|ref|XP_017141370.1| PREDICTED: phosphoglycerate mutase 2 isoform X5 [Drosophila
>gi|1037053118|ref|XP_017141371.1| PREDICTED: phosphoglycerate mutase 2 isoform X5 [Drosophila
>gi|1037053152|ref|XP_017141373.1| PREDICTED: carbonic anhydrase 15 [Drosophila miranda]
>gi|1037052927|ref|XP_017141357.1| PREDICTED: titin [Drosophila miranda]
>gi|1037053134|ref|XP_017141372.1| PREDICTED: protein suppressor of variegation 3-7 [Drosophila
>gi|1037053022|ref|XP_017141364.1| PREDICTED: protein suppressor of variegation 3-7 [Drosophila
>gi|1037052942|ref|XP_017141358.1| PREDICTED: uncharacterized protein LOC108155197 [Drosophila
>gi|1037052958|ref|XP_017141359.1| PREDICTED: pyruvate kinase-like [Drosophila miranda]
>gi|1037052974|ref|XP_017141360.1| PREDICTED: pyruvate kinase isoform X1 [Drosophila miranda]
>gi|1037053006|ref|XP_017141362.1| PREDICTED: pyruvate kinase isoform X3 [Drosophila miranda]
>gi|1037052990|ref|XP_017141361.1| PREDICTED: pyruvate kinase isoform X2 [Drosophila miranda]
>gi|1037089596|ref|XP_017143902.1| PREDICTED: putative OPA3-like protein CG13603 [Drosophila mi
>gi|1037089588|ref|XP_017143901.1| PREDICTED: putative OPA3-like protein CG43998 [Drosophila mi
>gi|1037089580|ref|XP_017143900.1| PREDICTED: putative OPA3-like protein CG13603 [Drosophila mi
>gi|1037089572|ref|XP_017143899.1| PREDICTED: nucleoporin Ndc1 [Drosophila miranda]
>gi|1037081417|ref|XP_017143311.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037081431|ref|XP_017143312.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037081473|ref|XP_017143316.1| PREDICTED: mucin-5AC isoform X4 [Drosophila miranda]
>gi|1037081459|ref|XP_017143315.1| PREDICTED: mucin-5AC isoform X3 [Drosophila miranda]
>gi|1037081487|ref|XP_017143317.1| PREDICTED: mucin-5AC isoform X5 [Drosophila miranda]
>gi|1037081445|ref|XP_017143314.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1037081501|ref|XP_017143318.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X6
>gi|1037081515|ref|XP_017143319.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X7
>gi|1037081529|ref|XP_017143320.1| PREDICTED: uncharacterized protein LOC108156396 [Drosophila
>gi|1037055029|ref|XP_017141504.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037054998|ref|XP_017141502.1| PREDICTED: serine/threonine-protein kinase fray2-like [Droso
>gi|1037024111|ref|XP_017139528.1| PREDICTED: uncharacterized protein LOC108153943 [Drosophila
>gi|1037055013|ref|XP_017141503.1| PREDICTED: uncharacterized protein LOC108155299 [Drosophila
>gi|1037053951|ref|XP_017141426.1| PREDICTED: progestin and adipoQ receptor family member 4 iso
>gi|1037053967|ref|XP_017141427.1| PREDICTED: progestin and adipoQ receptor family member 4 iso
>gi|1037053923|ref|XP_017141424.1| PREDICTED: neprilysin-21-like [Drosophila miranda]
>gi|1037053935|ref|XP_017141425.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1037053997|ref|XP_017141429.1| PREDICTED: cytochrome b-c1 complex subunit 2, mitochondrial-
>gi|1037053983|ref|XP_017141428.1| PREDICTED: progestin and adipoQ receptor family member 4 iso
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>gi|1037053912|ref|XP_017141423.1| PREDICTED: dedicator of cytokinesis protein 3 isoform X2 [D]
>gi|1037053870|ref|XP_017141422.1| PREDICTED: dedicator of cytokinesis protein 3 isoform X1 [D]
>gi|1037054007|ref|XP_017141430.1| PREDICTED: innexin inx3 [Drosophila miranda]
>gi|1037088527|ref|XP_017143820.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037088499|ref|XP_017143818.1| PREDICTED: LIM and senescent cell antigen-like-containing do
>gi|1037088486|ref|XP_017143817.1| PREDICTED: LIM and senescent cell antigen-like-containing do
>gi|1037088513|ref|XP_017143819.1| PREDICTED: LIM and senescent cell antigen-like-containing do
>gi|1037103890|ref|XP_017145141.1| PREDICTED: uncharacterized protein LOC108157547 [Drosophila
>gi|1037035447|ref|XP_017140214.1| PREDICTED: calcineurin subunit B isoform X1 [Drosophila mira
>gi|1037035465|ref|XP_017140215.1| PREDICTED: calcineurin subunit B isoform X2 [Drosophila mira
>gi|1037105975|ref|XP_017145433.1| PREDICTED: homeotic protein empty spiracles [Drosophila mira
>gi|1037101321|ref|XP_017144869.1| PREDICTED: homeotic protein empty spiracles [Drosophila mira
>gi|1037106524|ref|XP_017145508.1| PREDICTED: protein arginine N-methyltransferase 1 [Drosophil
>gi|1037106301|ref|XP_017145478.1| PREDICTED: protein arginine N-methyltransferase 1 [Drosophil
>gi|1037024125|ref|XP_017139530.1| PREDICTED: uncharacterized protein LOC108153944 [Drosophila
>gi|1037037484|ref|XP_017140330.1| PREDICTED: hepatoma-derived growth factor-related protein 2
>gi|1037046455|ref|XP_017140919.1| PREDICTED: inositol polyphosphate 1-phosphatase [Drosophila
>gi|1037046379|ref|XP_017140914.1| PREDICTED: uncharacterized protein LOC108154936 [Drosophila
>gi|1037046423|ref|XP_017140917.1| PREDICTED: origin recognition complex subunit 2 [Drosophila
>gi|1037046393|ref|XP_017140915.1| PREDICTED: protein roadkill isoform X1 [Drosophila miranda]
>gi|1037046407|ref|XP_017140916.1| PREDICTED: protein roadkill isoform X2 [Drosophila miranda]
>gi|1037046439|ref|XP_017140918.1| PREDICTED: probable cytochrome P450 6d5 [Drosophila miranda]
>gi|1037093156|ref|XP_017144166.1| PREDICTED: dnaJ homolog subfamily B member 12 isoform X1 [D]
>gi|1037093168|ref|XP_017144167.1| PREDICTED: dnaJ homolog subfamily B member 12 isoform X2 [D]
>gi|1037093186|ref|XP_017144169.1| PREDICTED: huntingtin-interacting protein K [Drosophila mira
>gi|1037084741|ref|XP_017143567.1| PREDICTED: forkhead box protein O isoform X1 [Drosophila mir
>gi|1037084757|ref|XP_017143568.1| PREDICTED: forkhead box protein O isoform X1 [Drosophila mir
>gi|1037084770|ref|XP_017143569.1| PREDICTED: forkhead box protein O isoform X1 [Drosophila mir
>gi|1037084784|ref|XP_017143570.1| PREDICTED: forkhead box protein O isoform X2 [Drosophila mir
>gi|1037033254|ref|XP_017140087.1| PREDICTED: attacin-A-like [Drosophila miranda]
>gi|1037104640|ref|XP_017145242.1| PREDICTED: ecdysteroid-regulated 16 kDa protein [Drosophila
>gi|1037104648|ref|XP_017145243.1| PREDICTED: ecdysteroid-regulated 16 kDa protein [Drosophila
>gi|1037104632|ref|XP_017145240.1| PREDICTED: neuropeptide CCHamide-1 [Drosophila miranda]
>gi|1037033867|ref|XP_017140123.1| PREDICTED: 10 kDa heat shock protein, mitochondrial [Drosop]
>gi|1037106189|ref|XP_017145463.1| PREDICTED: neuromedin-U receptor 2 [Drosophila miranda]
>gi|1037100973|ref|XP_017144837.1| PREDICTED: uncharacterized protein LOC108157332 isoform X1
>gi|1037100983|ref|XP_017144838.1| PREDICTED: uncharacterized protein LOC108157332 isoform X2
>gi|1037100993|ref|XP_017144839.1| PREDICTED: odorant receptor 88a [Drosophila miranda]
>gi|1037059140|ref|XP_017141780.1| PREDICTED: kinesin-like protein KIF19 [Drosophila miranda]
>gi|1037059201|ref|XP_017141784.1| PREDICTED: abl interactor 2 isoform X4 [Drosophila miranda]
>gi|1037059170|ref|XP_017141782.1| PREDICTED: abl interactor 2 isoform X2 [Drosophila miranda]
>gi|1037059185|ref|XP_017141783.1| PREDICTED: abl interactor 2 isoform X3 [Drosophila miranda]
>gi|1037059152|ref|XP_017141781.1| PREDICTED: abl interactor 2 isoform X1 [Drosophila miranda]
>gi|1037059217|ref|XP_017141785.1| PREDICTED: twinfilin [Drosophila miranda]
>gi|1037059233|ref|XP_017141786.1| PREDICTED: RPII140-upstream gene protein [Drosophila miranda
>gi|1037059127|ref|XP_017141779.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB2 [Dros
>gi|1037034496|ref|XP_017140163.1| PREDICTED: uncharacterized protein LOC108154404 [Drosophila
>gi|1037032724|ref|XP_017140055.1| PREDICTED: protein lifeguard 1 [Drosophila miranda]
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>gi|1037032740|ref|XP_017140056.1| PREDICTED: protein lifeguard 1 [Drosophila miranda]
>gi|1037038898|ref|XP_017140415.1| PREDICTED: transcription initiation factor IIA subunit 2-lik
>gi|1037103637|ref|XP_017145108.1| PREDICTED: uncharacterized protein LOC108157517 isoform X1
>gi|1037103645|ref|XP_017145109.1| PREDICTED: uncharacterized protein LOC108157517 isoform X2
>gi|1037105664|ref|XP_017145389.1| PREDICTED: uncharacterized protein LOC108157729 [Drosophila
>gi|1037034859|ref|XP_017140183.1| PREDICTED: dopamine receptor 1 [Drosophila miranda]
>gi|1037103708|ref|XP_017145117.1| PREDICTED: serine protease gd [Drosophila miranda]
>gi|1037103716|ref|XP_017145118.1| PREDICTED: enteropeptidase [Drosophila miranda]
>gi|1037103724|ref|XP_017145119.1| PREDICTED: uncharacterized protein LOC108157526 [Drosophila
>gi|1037104493|ref|XP_017145221.1| PREDICTED: LOW QUALITY PROTEIN: proclotting enzyme [Drosophi
>gi|1037104485|ref|XP_017145220.1| PREDICTED: serine protease gd [Drosophila miranda]
>gi|1037106703|ref|XP_017145533.1| PREDICTED: uncharacterized protein LOC108157841 [Drosophila
>gi|1037095258|ref|XP_017144324.1| PREDICTED: methionine--tRNA ligase, mitochondrial [Drosophil
>gi|1037034155|ref|XP_017140140.1| PREDICTED: uncharacterized protein LOC108154388 [Drosophila
>gi|1037059516|ref|XP_017141804.1| PREDICTED: uncharacterized protein LOC108155491 [Drosophila
>gi|1037059530|ref|XP_017141806.1| PREDICTED: uncharacterized protein LOC108155492 [Drosophila
>gi|1037059546|ref|XP_017141807.1| PREDICTED: histone-lysine N-methyltransferase trithorax [Dro
>gi|1037100678|ref|XP_017144806.1| PREDICTED: bromodomain-containing protein 4B-like [Drosophil
>gi|1037100688|ref|XP_017144807.1| PREDICTED: bromodomain-containing protein 4B-like [Drosophil
>gi|1037100668|ref|XP_017144805.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1037100698|ref|XP_017144808.1| PREDICTED: lachesin [Drosophila miranda]
>gi|1037102436|ref|XP_017144977.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037038880|ref|XP_017140414.1| PREDICTED: uncharacterized protein LOC108154608 [Drosophila
>gi|1037083797|ref|XP_017143492.1| PREDICTED: SCY1-like protein 2 [Drosophila miranda]
>gi|1037083811|ref|XP_017143493.1| PREDICTED: adipocyte plasma membrane-associated protein [Dro
>gi|1037083825|ref|XP_017143494.1| PREDICTED: adipocyte plasma membrane-associated protein [Dro
>gi|1037083838|ref|XP_017143495.1| PREDICTED: beta-1,4-N-acetylgalactosaminyltransferase bre-4
>gi|1037082815|ref|XP_017143416.1| PREDICTED: probable cleavage and polyadenylation specificit
>gi|1037082829|ref|XP_017143417.1| PREDICTED: pre-mRNA-splicing factor Slu7 [Drosophila mirand
>gi|1037088359|ref|XP_017143807.1| PREDICTED: protein kinase C isoform X1 [Drosophila miranda]
>gi|1037088375|ref|XP_017143808.1| PREDICTED: protein kinase C isoform X2 [Drosophila miranda]
>gi|1037088389|ref|XP_017143809.1| PREDICTED: protein kinase C isoform X3 [Drosophila miranda]
>gi|1037088403|ref|XP_017143810.1| PREDICTED: protein kinase C isoform X4 [Drosophila miranda]
>gi|1037088417|ref|XP_017143811.1| PREDICTED: protein kinase C isoform X5 [Drosophila miranda]
>gi|1037079891|ref|XP_017143195.1| PREDICTED: probable dimethyladenosine transferase [Drosophil
>gi|1037079865|ref|XP_017143193.1| PREDICTED: serine protease snake-like [Drosophila miranda]
>gi|1037079854|ref|XP_017143192.1| PREDICTED: serine protease snake-like [Drosophila miranda]
>gi|1037079877|ref|XP_017143194.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1037079842|ref|XP_017143191.1| PREDICTED: NAD-dependent protein deacetylase Sirt7 [Drosophi
>gi|1037079905|ref|XP_017143196.1| PREDICTED: attacin-A-like [Drosophila miranda]
>gi|1037091692|ref|XP_017144067.1| PREDICTED: cullin-5 [Drosophila miranda]
>gi|1037090323|ref|XP_017143964.1| PREDICTED: tyrosine-protein phosphatase vhp-1 [Drosophila m
>gi|1037090921|ref|XP_017144010.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila mira
>gi|1037090935|ref|XP_017144011.1| PREDICTED: pre-mRNA-splicing factor ISY1 homolog [Drosophila
>gi|1037094556|ref|XP_017144270.1| PREDICTED: ADP-ribosylation factor-like protein 8 [Drosophil
>gi|1037093213|ref|XP_017144171.1| PREDICTED: GATA-binding factor C isoform X2 [Drosophila mira
>gi|1037093200|ref|XP_017144170.1| PREDICTED: GATA-binding factor C isoform X1 [Drosophila mira
>gi|1037038171|ref|XP_017140369.1| PREDICTED: uncharacterized protein LOC108154578 [Drosophila
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>gi|1037104089|ref|XP_017145166.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1037024143|ref|XP_017139531.1| PREDICTED: uncharacterized protein LOC108153945 [Drosophila
>gi|1037077475|ref|XP_017143028.1| PREDICTED: uncharacterized protein LOC108156204 [Drosophila
>gi|1037077447|ref|XP_017143025.1| PREDICTED: DNA polymerase iota [Drosophila miranda]
>gi|1037077461|ref|XP_017143026.1| PREDICTED: DNA polymerase iota [Drosophila miranda]
>gi|1037077503|ref|XP_017143030.1| PREDICTED: transcriptional adapter 2B isoform X2 [Drosophila
>gi|1037077489|ref|XP_017143029.1| PREDICTED: transcriptional adapter 2B isoform X1 [Drosophila
>gi|1037077517|ref|XP_017143031.1| PREDICTED: transcriptional adapter 2B isoform X3 [Drosophila
>gi|1037093661|ref|XP_017144200.1| PREDICTED: CUE domain-containing protein 2 [Drosophila miran
>gi|1037093675|ref|XP_017144201.1| PREDICTED: CUE domain-containing protein 2 [Drosophila miran
>gi|1037093689|ref|XP_017144202.1| PREDICTED: CUE domain-containing protein 2 [Drosophila miran
>gi|1037074992|ref|XP_017142859.1| PREDICTED: pro-corazonin [Drosophila miranda]
>gi|1037074960|ref|XP_017142857.1| PREDICTED: uncharacterized protein LOC108156080 [Drosophila
>gi|1037075007|ref|XP_017142860.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB9 [Dros
>gi|1037074928|ref|XP_017142854.1| PREDICTED: protein suppressor of hairy wing [Drosophila mira
>gi|1037074942|ref|XP_017142855.1| PREDICTED: protein suppressor of hairy wing [Drosophila mira
>gi|1037074976|ref|XP_017142858.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1037092738|ref|XP_017144138.1| PREDICTED: uncharacterized protein LOC108156904 isoform X1
>gi|1037092752|ref|XP_017144139.1| PREDICTED: uncharacterized protein LOC108156904 isoform X2
>gi|1037092766|ref|XP_017144140.1| PREDICTED: uncharacterized protein LOC108156904 isoform X3
>gi|1037092778|ref|XP_017144141.1| PREDICTED: uncharacterized protein LOC108156904 isoform X3
>gi|1037092792|ref|XP_017144142.1| PREDICTED: uncharacterized protein LOC108156904 isoform X4
>gi|1037101821|ref|XP_017144912.1| PREDICTED: transcription initiation factor IIA subunit 2 [Dr
>gi|1037056692|ref|XP_017141611.1| PREDICTED: protein pellino [Drosophila miranda]
>gi|1037056712|ref|XP_017141612.1| PREDICTED: X-linked retinitis pigmentosa GTPase regulator-in
>gi|1037056644|ref|XP_017141608.1| PREDICTED: larval serum protein 1 beta chain [Drosophila mi
>gi|1037056676|ref|XP_017141610.1| PREDICTED: protein Skeletor, isoforms B/C isoform X2 [Drosop
>gi|1037056660|ref|XP_017141609.1| PREDICTED: protein Skeletor, isoforms B/C isoform X1 [Drosop
>gi|1037061641|ref|XP_017141953.1| PREDICTED: eukaryotic translation initiation factor 4G isof
>gi|1037061657|ref|XP_017141954.1| PREDICTED: uncharacterized protein LOC108155569 isoform X2
>gi|1037061672|ref|XP_017141955.1| PREDICTED: shootin-1 isoform X3 [Drosophila miranda]
>gi|1037061687|ref|XP_017141957.1| PREDICTED: shootin-1 isoform X3 [Drosophila miranda]
>gi|1037061718|ref|XP_017141958.1| PREDICTED: myosin heavy chain, striated muscle isoform X4 [
>gi|1037064506|ref|XP_017142151.1| PREDICTED: dedicator of cytokinesis protein 1 isoform X2 [D
>gi|1037064492|ref|XP_017142150.1| PREDICTED: dedicator of cytokinesis protein 1 isoform X1 [D
>gi|1037071742|ref|XP_017142635.1| PREDICTED: LOW QUALITY PROTEIN: nuclear pore complex protei
>gi|1037071759|ref|XP_017142636.1| PREDICTED: uncharacterized protein C11D3.03c [Drosophila mi
>gi|1037040068|ref|XP_017140493.1| PREDICTED: helicase SKI2W [Drosophila miranda]
>gi|1037040246|ref|XP_017140505.1| PREDICTED: dnaJ homolog subfamily C member 8 [Drosophila mi
>gi|1037040259|ref|XP_017140507.1| PREDICTED: probable oligoribonuclease [Drosophila miranda]
>gi|1037040215|ref|XP_017140503.1| PREDICTED: lipid storage droplets surface-binding protein 1
>gi|1037040229|ref|XP_017140504.1| PREDICTED: lipid storage droplets surface-binding protein 1
>gi|1037040168|ref|XP_017140500.1| PREDICTED: uncharacterized protein LOC108154670 [Drosophila
>gi|1037040288|ref|XP_017140509.1| PREDICTED: succinate dehydrogenase [ubiquinone] cytochrome
>gi|1037040275|ref|XP_017140508.1| PREDICTED: phosphatidylglycerophosphatase and protein-tyros
>gi|1037040105|ref|XP_017140496.1| PREDICTED: protein sel-1 homolog 1 [Drosophila miranda]
>gi|1037040119|ref|XP_017140497.1| PREDICTED: protein sel-1 homolog 1 [Drosophila miranda]
>gi|1037040134|ref|XP_017140498.1| PREDICTED: protein sel-1 homolog 1 [Drosophila miranda]
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>gi|1037040152|ref|XP_017140499.1| PREDICTED: protein sel-1 homolog 1 [Drosophila miranda]
>gi|1037040199|ref|XP_017140502.1| PREDICTED: 26S protease regulatory subunit 6A-B [Drosophila
>gi|1037040184|ref|XP_017140501.1| PREDICTED: ran-binding protein 3 [Drosophila miranda]
>gi|1037040090|ref|XP_017140495.1| PREDICTED: 3-hydroxy-3-methylglutaryl-coenzyme A reductase
>gi|1037064055|ref|XP_017142120.1| PREDICTED: adenylosuccinate synthetase [Drosophila miranda]
>gi|1037064087|ref|XP_017142122.1| PREDICTED: calmodulin-like protein 4 [Drosophila miranda]
>gi|1037064105|ref|XP_017142123.1| PREDICTED: 40S ribosomal protein S20 [Drosophila miranda]
>gi|1037064071|ref|XP_017142121.1| PREDICTED: zinc finger HIT domain-containing protein 2 [Dros
>gi|1037064037|ref|XP_017142119.1| PREDICTED: protein kinase C and casein kinase II substrate p
>gi|1037064019|ref|XP_017142118.1| PREDICTED: protein kinase C and casein kinase II substrate p
>gi|1037103732|ref|XP_017145120.1| PREDICTED: uncharacterized protein LOC108157527 [Drosophila
>gi|1037105169|ref|XP_017145319.1| PREDICTED: uncharacterized protein LOC108157668 isoform X2
>gi|1037105164|ref|XP_017145318.1| PREDICTED: uncharacterized protein LOC108157668 isoform X1
>gi|1037032887|ref|XP_017140065.1| PREDICTED: homeobox protein EMX2 [Drosophila miranda]
>gi|1037088921|ref|XP_017143852.1| PREDICTED: uncharacterized protein LOC108156714 [Drosophila
>gi|1037088935|ref|XP_017143853.1| PREDICTED: glutamate receptor ionotropic, delta-2 [Drosophil
>gi|1037088908|ref|XP_017143850.1| PREDICTED: glutamate receptor ionotropic, kainate 2 [Drosoph
>gi|1037038349|ref|XP_017140381.1| PREDICTED: protein Turandot B1-like [Drosophila miranda]
>gi|1037105106|ref|XP_017145310.1| PREDICTED: BTB/POZ domain-containing protein KCTD12 [Drosoph
>gi|1037024161|ref|XP_017139532.1| PREDICTED: protein Turandot X [Drosophila miranda]
>gi|1037034357|ref|XP_017140154.1| PREDICTED: protein Turandot E-like [Drosophila miranda]
>gi|1037106558|ref|XP_017145513.1| PREDICTED: glutamate receptor ionotropic, kainate 2 isoform
>gi|1037106565|ref|XP_017145514.1| PREDICTED: glutamate receptor ionotropic, kainate 2 isoform
>gi|1037024177|ref|XP_017139533.1| PREDICTED: protein bunched, class 2/F/G isoform [Drosophila
>gi|1037097039|ref|XP_017144478.1| PREDICTED: uncharacterized protein LOC108157113 isoform X1
>gi|1037097050|ref|XP_017144479.1| PREDICTED: uncharacterized protein LOC108157113 isoform X1
>gi|1037097061|ref|XP_017144481.1| PREDICTED: uncharacterized protein LOC108157113 isoform X1
>gi|1037097073|ref|XP_017144482.1| PREDICTED: uncharacterized protein LOC108157113 isoform X1
>gi|1037097085|ref|XP_017144483.1| PREDICTED: uncharacterized protein LOC108157113 isoform X2
>gi|1037063130|ref|XP_017142056.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
>gi|1037063098|ref|XP_017142054.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
>gi|1037063083|ref|XP_017142053.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
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>gi|1037063146|ref|XP_017142057.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
>gi|1037063164|ref|XP_017142058.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
>gi|1037088229|ref|XP_017143797.1| PREDICTED: sodium/calcium exchanger 3 isoform X2 [Drosophila
>gi|1037088243|ref|XP_017143798.1| PREDICTED: sodium/calcium exchanger 3 isoform X3 [Drosophila
>gi|1037088255|ref|XP_017143799.1| PREDICTED: sodium/calcium exchanger 3 isoform X4 [Drosophila
>gi|1037088175|ref|XP_017143793.1| PREDICTED: sodium/calcium exchanger 3 isoform X1 [Drosophila
>gi|1037088189|ref|XP_017143794.1| PREDICTED: sodium/calcium exchanger 3 isoform X1 [Drosophila
>gi|1037088205|ref|XP_017143795.1| PREDICTED: sodium/calcium exchanger 3 isoform X1 [Drosophila
>gi|1037088217|ref|XP_017143796.1| PREDICTED: sodium/calcium exchanger 3 isoform X1 [Drosophila
>gi|1037088269|ref|XP_017143800.1| PREDICTED: sodium/calcium exchanger 3 isoform X5 [Drosophila
>gi|1037088315|ref|XP_017143804.1| PREDICTED: sodium/calcium exchanger 3 isoform X8 [Drosophila
>gi|1037088297|ref|XP_017143803.1| PREDICTED: sodium/calcium exchanger 3 isoform X7 [Drosophila
>gi|1037088329|ref|XP_017143805.1| PREDICTED: sodium/calcium exchanger 3 isoform X9 [Drosophila
>gi|1037088283|ref|XP_017143801.1| PREDICTED: sodium/calcium exchanger 3 isoform X6 [Drosophila
>gi|1037088345|ref|XP_017143806.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila m
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>gi|1037103329|ref|XP_017145071.1| PREDICTED: uncharacterized protein LOC108157487 [Drosophila
>gi|1037103319|ref|XP_017145070.1| PREDICTED: chorion peroxidase [Drosophila miranda]
>gi|1037092516|ref|XP_017144125.1| PREDICTED: wolframin isoform X1 [Drosophila miranda]
>gi|1037092534|ref|XP_017144127.1| PREDICTED: wolframin isoform X2 [Drosophila miranda]
>gi|1037069722|ref|XP_017142497.1| PREDICTED: glutamate--cysteine ligase regulatory subunit [Di
>gi|1037069752|ref|XP_017142499.1| PREDICTED: uncharacterized protein LOC108155881 isoform X2
>gi|1037069738|ref|XP_017142498.1| PREDICTED: uncharacterized protein LOC108155881 isoform X1
>gi|1037024193|ref|XP_017139534.1| PREDICTED: putative uncharacterized protein DDB_G0277255 [Di
>gi|1037095649|ref|XP_017144354.1| PREDICTED: protein FAM50 homolog [Drosophila miranda]
>gi|1037066061|ref|XP_017142244.1| PREDICTED: N-acetyl-D-glucosamine kinase [Drosophila miranda
>gi|1037066046|ref|XP_017142243.1| PREDICTED: serine--tRNA ligase, mitochondrial [Drosophila mi
>gi|1037066030|ref|XP_017142242.1| PREDICTED: cytoplasmic FMR1-interacting protein [Drosophila
>gi|1037104097|ref|XP_017145167.1| PREDICTED: uncharacterized protein LOC108157562 isoform X1
>gi|1037104104|ref|XP_017145168.1| PREDICTED: uncharacterized protein LOC108157562 isoform X1
>gi|1037104112|ref|XP_017145169.1| PREDICTED: uncharacterized protein LOC108157562 isoform X2
>gi|1037079552|ref|XP_017143168.1| PREDICTED: 39 kDa FK506-binding nuclear protein [Drosophila
>gi|1037079565|ref|XP_017143169.1| PREDICTED: 39S ribosomal protein L9, mitochondrial [Drosophi
>gi|1037024211|ref|XP_017139535.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037079579|ref|XP_017143170.1| PREDICTED: uncharacterized protein LOC108156297 [Drosophila
>gi|1037079593|ref|XP_017143171.1| PREDICTED: uncharacterized protein LOC108156298 [Drosophila
>gi|1037080111|ref|XP_017143212.1| PREDICTED: serine/threonine-protein phosphatase 5 [Drosophi
>gi|1037080097|ref|XP_017143209.1| PREDICTED: TELO2-interacting protein 1 homolog [Drosophila i
>gi|1037105902|ref|XP_017145422.1| PREDICTED: 26S protease regulatory subunit 6B [Drosophila mi
>gi|1037056808|ref|XP_017141616.1| PREDICTED: probable Dol-P-Man:Man(7)GlcNAc(2)-PP-Dol alpha-i
>gi|1037056824|ref|XP_017141617.1| PREDICTED: probable Dol-P-Man:Man(7)GlcNAc(2)-PP-Dol alpha-i
>gi|1037056856|ref|XP_017141620.1| PREDICTED: mannose-6-phosphate isomerase [Drosophila miranda
>gi|1037056885|ref|XP_017141622.1| PREDICTED: synaptosomal-associated protein 25 [Drosophila mi
>gi|1037056840|ref|XP_017141619.1| PREDICTED: uncharacterized protein LOC108155366 [Drosophila
>gi|1037056871|ref|XP_017141621.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1037056917|ref|XP_017141624.1| PREDICTED: N-acetyltransferase 6 isoform X2 [Drosophila mira
>gi|1037056901|ref|XP_017141623.1| PREDICTED: N-acetyltransferase 6 isoform X1 [Drosophila mira
>gi|1037084687|ref|XP_017143562.1| PREDICTED: flavin reductase (NADPH) [Drosophila miranda]
>gi|1037084660|ref|XP_017143560.1| PREDICTED: formin-like protein 3 isoform X1 [Drosophila mira
>gi|1037084674|ref|XP_017143561.1| PREDICTED: formin-like protein 3 isoform X2 [Drosophila mira
>gi|1037084634|ref|XP_017143558.1| PREDICTED: testis-specific zinc finger protein topi isoform
>gi|1037084647|ref|XP_017143559.1| PREDICTED: testis-specific zinc finger protein topi isoform
>gi|1037084700|ref|XP_017143564.1| PREDICTED: 40S ribosomal protein S29 [Drosophila miranda]
>gi|1037084727|ref|XP_017143566.1| PREDICTED: uncharacterized protein LOC108156555 isoform X2
>gi|1037084713|ref|XP_017143565.1| PREDICTED: uncharacterized protein LOC108156555 isoform X1
>gi|1037094021|ref|XP_017144228.1| PREDICTED: metallothionein-1 [Drosophila miranda]
>gi|1037094007|ref|XP_017144227.1| PREDICTED: GTP-binding protein Di-Ras2 [Drosophila miranda]
>gi|1037048331|ref|XP_017141040.1| PREDICTED: uncharacterized protein LOC108155003 [Drosophila
>gi|1037048347|ref|XP_017141041.1| PREDICTED: alpha-2-macroglobulin receptor-associated protein
>gi|1037048279|ref|XP_017141036.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mir
>gi|1037048299|ref|XP_017141038.1| PREDICTED: BTB/POZ domain-containing protein KCTD3 [Drosophi
>gi|1037048315|ref|XP_017141039.1| PREDICTED: L-asparaginase 1 [Drosophila miranda]
>gi|1037048259|ref|XP_017141035.1| PREDICTED: F-box only protein 11 [Drosophila miranda]
>gi|1037035199|ref|XP_017140202.1| PREDICTED: uncharacterized protein LOC108154440 [Drosophila
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>gi|1037032670|ref|XP_017140052.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037033650|ref|XP_017140110.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037101499|ref|XP_017144879.1| PREDICTED: uncharacterized protein LOC108157364 [Drosophila
>gi|1037101373|ref|XP_017144872.1| PREDICTED: COP9 signalosome complex subunit 1b-like [Drosopl
>gi|1037101383|ref|XP_017144873.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1037101461|ref|XP_017144874.1| PREDICTED: pre-mRNA-splicing factor cwc24-like isoform X1 [
>gi|1037101471|ref|XP_017144876.1| PREDICTED: E3 ubiquitin-protein ligase RNF185-like isoform
>gi|1037101483|ref|XP_017144877.1| PREDICTED: uncharacterized protein LOC108157363 isoform X1
>gi|1037101490|ref|XP_017144878.1| PREDICTED: uncharacterized protein LOC108157363 isoform X2
>gi|1037035287|ref|XP_017140205.1| PREDICTED: uncharacterized protein LOC108154445 [Drosophila
>gi|1037105203|ref|XP_017145323.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1037105211|ref|XP_017145324.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1037105500|ref|XP_017145367.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1037105508|ref|XP_017145368.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1037076405|ref|XP_017142955.1| PREDICTED: fasciclin-1 isoform X8 [Drosophila miranda]
>gi|1037076296|ref|XP_017142948.1| PREDICTED: fasciclin-1 isoform X1 [Drosophila miranda]
>gi|1037076360|ref|XP_017142952.1| PREDICTED: fasciclin-1 isoform X5 [Drosophila miranda]
>gi|1037076328|ref|XP_017142950.1| PREDICTED: fasciclin-1 isoform X3 [Drosophila miranda]
>gi|1037076373|ref|XP_017142953.1| PREDICTED: fasciclin-1 isoform X6 [Drosophila miranda]
>gi|1037076312|ref|XP_017142949.1| PREDICTED: fasciclin-1 isoform X2 [Drosophila miranda]
>gi|1037076344|ref|XP_017142951.1| PREDICTED: fasciclin-1 isoform X4 [Drosophila miranda]
>gi|1037076391|ref|XP_017142954.1| PREDICTED: fasciclin-1 isoform X7 [Drosophila miranda]
>gi|1037088472|ref|XP_017143815.1| PREDICTED: zinc finger protein 800 [Drosophila miranda]
>gi|1037084116|ref|XP_017143517.1| PREDICTED: uncharacterized protein LOC108156524 [Drosophila
>gi|1037084130|ref|XP_017143519.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037084143|ref|XP_017143520.1| PREDICTED: LOW QUALITY PROTEIN: THO complex subunit 1 [Drosc
>gi|1037099055|ref|XP_017144669.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miranc
>gi|1037099043|ref|XP_017144668.1| PREDICTED: vesicle transport protein SEC20 [Drosophila miran
>gi|1037099203|ref|XP_017144677.1| PREDICTED: E3 ubiquitin-protein ligase complex slx8-rfp subu
>gi|1037099213|ref|XP_017144678.1| PREDICTED: E3 ubiquitin-protein ligase complex slx8-rfp subu
>gi|1037106145|ref|XP_017145456.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1037024227|ref|XP_017139536.1| PREDICTED: protein FAM170B [Drosophila miranda]
>gi|1037036795|ref|XP_017140288.1| PREDICTED: uncharacterized protein LOC108154499 [Drosophila
>gi|1037034054|ref|XP_017140134.1| PREDICTED: uncharacterized protein LOC108154381 [Drosophila
>gi|1037038832|ref|XP_017140411.1| PREDICTED: uncharacterized protein LOC108154605 [Drosophila
>gi|1037024245|ref|XP_017139538.1| PREDICTED: uncharacterized protein LOC108153951 [Drosophila
>gi|1037037558|ref|XP_017140335.1| PREDICTED: transcription factor SPT20 homolog [Drosophila m
>gi|1037105876|ref|XP_017145419.1| PREDICTED: uncharacterized protein LOC108157749 [Drosophila
>gi|1037042106|ref|XP_017140631.1| PREDICTED: carcinine transporter [Drosophila miranda]
>gi|1037042122|ref|XP_017140633.1| PREDICTED: uncharacterized protein LOC108154762 [Drosophila
>gi|1037042031|ref|XP_017140626.1| PREDICTED: uncharacterized protein LOC108154760 isoform X3
>gi|1037042015|ref|XP_017140625.1| PREDICTED: uncharacterized protein LOC108154760 isoform X2
>gi|1037042001|ref|XP_017140624.1| PREDICTED: uncharacterized protein LOC108154760 isoform X1
>gi|1037042045|ref|XP_017140627.1| PREDICTED: uncharacterized protein LOC108154760 isoform X4
>gi|1037042061|ref|XP_017140628.1| PREDICTED: sterile alpha motif domain-containing protein 5-
>gi|1037042076|ref|XP_017140629.1| PREDICTED: sterile alpha motif domain-containing protein 5-
>gi|1037042090|ref|XP_017140630.1| PREDICTED: sterile alpha motif domain-containing protein 5-
>gi|1037041986|ref|XP_017140623.1| PREDICTED: uncharacterized protein LOC108154759 [Drosophila
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>gi|1037024261|ref|XP_017139539.1| PREDICTED: uncharacterized protein LOC108153952 [Drosophila
>gi|1037024279|ref|XP_017139540.1| PREDICTED: uncharacterized protein LOC108153953 [Drosophila
>gi|1037024297|ref|XP_017139541.1| PREDICTED: uncharacterized protein LOC108153954 [Drosophila
>gi|1037024323|ref|XP_017139542.1| PREDICTED: uncharacterized protein LOC108153955 [Drosophila
>gi|1037024339|ref|XP_017139543.1| PREDICTED: uncharacterized protein LOC108153956 [Drosophila
>gi|1037097620|ref|XP_017144533.1| PREDICTED: serine/threonine-protein kinase D1 isoform X2 [Dr
>gi|1037097609|ref|XP_017144532.1| PREDICTED: serine/threonine-protein kinase D1 isoform X1 [Dr
>gi|1037036587|ref|XP_017140277.1| PREDICTED: uncharacterized protein LOC108154487 [Drosophila
>gi|1037105794|ref|XP_017145408.1| PREDICTED: retinol dehydrogenase 13 [Drosophila miranda]
>gi|1037102653|ref|XP_017144997.1| PREDICTED: patj homolog isoform X3 [Drosophila miranda]
>gi|1037102645|ref|XP_017144996.1| PREDICTED: patj homolog isoform X2 [Drosophila miranda]
>gi|1037102636|ref|XP_017144995.1| PREDICTED: patj homolog isoform X1 [Drosophila miranda]
>gi|1037097671|ref|XP_017144537.1| PREDICTED: protein hemingway [Drosophila miranda]
>gi|1037097685|ref|XP_017144538.1| PREDICTED: Werner Syndrome-like exonuclease [Drosophila mira
>gi|1037095397|ref|XP_017144334.1| PREDICTED: nucleoporin Nup43 [Drosophila miranda]
>gi|1037095411|ref|XP_017144335.1| PREDICTED: zinc finger protein 283-like [Drosophila miranda]
>gi|1037095383|ref|XP_017144333.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1037105382|ref|XP_017145350.1| PREDICTED: retinol dehydrogenase 14 isoform X1 [Drosophila m
>gi|1037105390|ref|XP_017145351.1| PREDICTED: retinol dehydrogenase 14 isoform X2 [Drosophila m
>gi|1037105476|ref|XP_017145364.1| PREDICTED: uncharacterized protein LOC108157705 [Drosophila
>gi|1037034088|ref|XP_017140136.1| PREDICTED: LOW QUALITY PROTEIN: V-type proton ATPase 116 kDa
>gi|1037084846|ref|XP_017143573.1| PREDICTED: V-type proton ATPase 116 kDa subunit a [Drosophil
>gi|1037061835|ref|XP_017141966.1| PREDICTED: LOW QUALITY PROTEIN: vesicle transport through in
>gi|1037061804|ref|XP_017141964.1| PREDICTED: LOW QUALITY PROTEIN: WD repeat-containing protein
>gi|1037061820|ref|XP_017141965.1| PREDICTED: cyclin-related protein FAM58A [Drosophila miranda
>gi|1037061943|ref|XP_017141974.1| PREDICTED: glucosidase 2 subunit beta isoform X1 [Drosophila
>gi|1037061958|ref|XP_017141975.1| PREDICTED: glucosidase 2 subunit beta isoform X2 [Drosophila
>gi|1037061881|ref|XP_017141969.1| PREDICTED: uncharacterized protein LOC108155577 isoform X2
>gi|1037061865|ref|XP_017141968.1| PREDICTED: UPF0565 protein C2orf69 homolog isoform X1 [Droso
>gi|1037061897|ref|XP_017141970.1| PREDICTED: TWiK family of potassium channels protein 18 [Dro
>gi|1037061915|ref|XP_017141971.1| PREDICTED: uncharacterized protein LOC108155580 [Drosophila
>gi|1037061930|ref|XP_017141972.1| PREDICTED: uncharacterized protein LOC108155580 [Drosophila
>gi|1037061974|ref|XP_017141976.1| PREDICTED: uncharacterized protein LOC108155582 [Drosophila
>gi|1037061994|ref|XP_017141977.1| PREDICTED: uncharacterized protein LOC108155582 [Drosophila
>gi|1037061849|ref|XP_017141967.1| PREDICTED: V-type proton ATPase 116 kDa subunit a-like [Dros
>gi|1037091390|ref|XP_017144046.1| PREDICTED: sex determination protein fruitless isoform X3 [D
>gi|1037091431|ref|XP_017144049.1| PREDICTED: GATA zinc finger domain-containing protein 10-lik
>gi|1037091417|ref|XP_017144048.1| PREDICTED: transcription factor mef2A-like isoform X1 [Droso
>gi|1037091376|ref|XP_017144045.1| PREDICTED: sex determination protein fruitless isoform X2 [D
>gi|1037091363|ref|XP_017144044.1| PREDICTED: sex determination protein fruitless isoform X1 [D
>gi|1037091404|ref|XP_017144047.1| PREDICTED: sex determination protein fruitless isoform X4 [D
>gi|1037032953|ref|XP_017140069.1| PREDICTED: uncharacterized protein LOC108154321 [Drosophila
>gi|1037102289|ref|XP_017144961.1| PREDICTED: zinc finger protein interacting with ribonucleopr
>gi|1037102307|ref|XP_017144963.1| PREDICTED: uncharacterized protein LOC108157424 [Drosophila
>gi|1037102299|ref|XP_017144962.1| PREDICTED: hemotin [Drosophila miranda]
>gi|1037024357|ref|XP_017139544.1| PREDICTED: microtubule-associated protein RP/EB family membe
>gi|1037091968|ref|XP_017144088.1| PREDICTED: serine/threonine-protein kinase OSR1 [Drosophila
>gi|1037091996|ref|XP_017144090.1| PREDICTED: E3 ubiquitin-protein ligase RNF181 homolog [Droso
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>gi|1037092010|ref|XP_017144091.1| PREDICTED: E3 ubiquitin-protein ligase RNF181 homolog [Dros
>gi|1037091982|ref|XP_017144089.1| PREDICTED: dentin sialophosphoprotein-like [Drosophila mira
>gi|1037069543|ref|XP_017142485.1| PREDICTED: uncharacterized protein LOC108155874 [Drosophila
>gi|1037069571|ref|XP_017142487.1| PREDICTED: trithorax group protein osa isoform X2 [Drosophil
>gi|1037069558|ref|XP_017142486.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X1
>gi|1037069583|ref|XP_017142488.1| PREDICTED: testis-specific serine/threonine-protein kinase
>gi|1037069598|ref|XP_017142489.1| PREDICTED: uncharacterized protein LOC108155877 [Drosophila
>gi|1037031492|ref|XP_017139983.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1037047835|ref|XP_017141013.1| PREDICTED: uncharacterized protein LOC108154985 [Drosophila
>gi|1037047849|ref|XP_017141015.1| PREDICTED: uncharacterized protein LOC108154985 [Drosophila
>gi|1037048017|ref|XP_017141021.1| PREDICTED: cleavage stimulation factor subunit 2 tau variant
>gi|1037047865|ref|XP_017141016.1| PREDICTED: cleavage and polyadenylation specificity factor
>gi|1037048035|ref|XP_017141023.1| PREDICTED: Golgi SNAP receptor complex member 1 [Drosophila
>gi|1037048065|ref|XP_017141025.1| PREDICTED: mitochondrial import receptor subunit TOM22 homol
>gi|1037048049|ref|XP_017141024.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037047970|ref|XP_017141018.1| PREDICTED: toll-like receptor 9 [Drosophila miranda]
>gi|1037047988|ref|XP_017141019.1| PREDICTED: toll-like receptor 9 [Drosophila miranda]
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>gi|1037047954|ref|XP_017141017.1| PREDICTED: uncharacterized protein LOC108154987 [Drosophila
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>gi|1037090253|ref|XP_017143958.1| PREDICTED: NADP-dependent malic enzyme isoform X1 [Drosophil
>gi|1037034912|ref|XP_017140186.1| PREDICTED: perlucin-like [Drosophila miranda]
>gi|1037035718|ref|XP_017140224.1| PREDICTED: lectin subunit alpha-like [Drosophila miranda]
>gi|1037033160|ref|XP_017140082.1| PREDICTED: uncharacterized protein LOC108154331 [Drosophila
>gi|1037106261|ref|XP_017145472.1| PREDICTED: calcyphosin-like protein isoform X1 [Drosophila m
>gi|1037106269|ref|XP_017145473.1| PREDICTED: calcyphosin-like protein isoform X2 [Drosophila m
>gi|1037106277|ref|XP_017145475.1| PREDICTED: calcyphosin-like protein isoform X2 [Drosophila m
>gi|1037106735|ref|XP_017145537.1| PREDICTED: uncharacterized protein LOC108157845 [Drosophila
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>gi|1037032450|ref|XP_017140037.1| PREDICTED: calcyphosin-like protein isoform X2 [Drosophila m
>gi|1037033683|ref|XP_017140111.1| PREDICTED: rRNA 2'-O-methyltransferase fibrillarin [Drosophi
>gi|1037033703|ref|XP_017140112.1| PREDICTED: uncharacterized protein LOC108154358 [Drosophila
>gi|1037033720|ref|XP_017140113.1| PREDICTED: uncharacterized protein LOC108154358 [Drosophila
>gi|1037038848|ref|XP_017140412.1| PREDICTED: LOW QUALITY PROTEIN: odorant receptor 67a [Drosop
>gi|1037096379|ref|XP_017144420.1| PREDICTED: probable aminopeptidase NPEPL1 [Drosophila miran
>gi|1037030827|ref|XP_017139942.1| PREDICTED: kinetochore protein Spc25 [Drosophila miranda]
>gi|1037104691|ref|XP_017145248.1| PREDICTED: probable cytochrome P450 304a1 [Drosophila miran
>gi|1037104699|ref|XP_017145250.1| PREDICTED: uncharacterized protein LOC108157624 [Drosophila
>gi|1037100424|ref|XP_017144782.1| PREDICTED: cell death abnormality protein 1 isoform X2 [Dros
>gi|1037100412|ref|XP_017144780.1| PREDICTED: cell death abnormality protein 1 isoform X1 [Dros
>gi|1037100436|ref|XP_017144783.1| PREDICTED: sialin isoform X1 [Drosophila miranda]
>gi|1037100448|ref|XP_017144784.1| PREDICTED: sialin isoform X2 [Drosophila miranda]
>gi|1037029110|ref|XP_017139838.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1037029130|ref|XP_017139839.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1037028794|ref|XP_017139819.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037029075|ref|XP_017139836.1| PREDICTED: uncharacterized protein LOC108154171 [Drosophila
>gi|1037029148|ref|XP_017139840.1| PREDICTED: uncharacterized protein LOC108154175 [Drosophila
>gi|1037029019|ref|XP_017139833.1| PREDICTED: L-dopachrome tautomerase yellow-f2-like isoform
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>gi|1037029001|ref|XP_017139832.1| PREDICTED: L-dopachrome tautomerase yellow-f2-like isoform
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>gi|1037029057|ref|XP_017139835.1| PREDICTED: L-dopachrome tautomerase yellow-f2-like isoform
>gi|1037028776|ref|XP_017139817.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037029166|ref|XP_017139842.1| PREDICTED: LOW QUALITY PROTEIN: protein MEMO1 [Drosophila mi
>gi|1037029093|ref|XP_017139837.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037028812|ref|XP_017139820.1| PREDICTED: C-terminal-binding protein isoform X1 [Drosophila
>gi|1037028826|ref|XP_017139821.1| PREDICTED: C-terminal-binding protein isoform X1 [Drosophila
>gi|1037028844|ref|XP_017139822.1| PREDICTED: C-terminal-binding protein isoform X1 [Drosophila
>gi|1037028882|ref|XP_017139824.1| PREDICTED: C-terminal-binding protein isoform X3 [Drosophila
>gi|1037028914|ref|XP_017139826.1| PREDICTED: C-terminal-binding protein isoform X5 [Drosophila
>gi|1037028864|ref|XP_017139823.1| PREDICTED: C-terminal-binding protein isoform X2 [Drosophila
>gi|1037028900|ref|XP_017139825.1| PREDICTED: C-terminal-binding protein isoform X4 [Drosophila
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>gi|1037028984|ref|XP_017139831.1| PREDICTED: C-terminal-binding protein isoform X9 [Drosophila
>gi|1037028966|ref|XP_017139830.1| PREDICTED: C-terminal-binding protein isoform X8 [Drosophila
>gi|1037029245|ref|XP_017139844.1| PREDICTED: traB domain-containing protein isoform X1 [Drosop
>gi|1037029262|ref|XP_017139845.1| PREDICTED: uncharacterized protein LOC108154181 isoform X2
>gi|1037029280|ref|XP_017139846.1| PREDICTED: cytochrome c oxidase protein 20 homolog [Drosophi
>gi|1037029227|ref|XP_017139843.1| PREDICTED: xanthine dehydrogenase [Drosophila miranda]
>gi|1037088134|ref|XP_017143790.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1037088092|ref|XP_017143786.1| PREDICTED: serine protease snake isoform X1 [Drosophila mira
>gi|1037088106|ref|XP_017143787.1| PREDICTED: serine protease snake isoform X2 [Drosophila mira
>gi|1037088120|ref|XP_017143789.1| PREDICTED: serine protease snake isoform X3 [Drosophila mira
>gi|1037088064|ref|XP_017143784.1| PREDICTED: serine protease snake isoform X1 [Drosophila mira
>gi|1037088078|ref|XP_017143785.1| PREDICTED: serine protease snake isoform X2 [Drosophila mira
>gi|1037088160|ref|XP_017143792.1| PREDICTED: small lysine-rich protein 1 [Drosophila miranda]
>gi|1037088050|ref|XP_017143783.1| PREDICTED: heat shock 70 kDa protein cognate 2 [Drosophila m
>gi|1037088147|ref|XP_017143791.1| PREDICTED: transmembrane protein 135-like [Drosophila mirand
>gi|1037104281|ref|XP_017145192.1| PREDICTED: uncharacterized protein LOC108157581 [Drosophila
>gi|1037104273|ref|XP_017145190.1| PREDICTED: selenium-binding protein 1 [Drosophila miranda]
>gi|1037059330|ref|XP_017141792.1| PREDICTED: 3-oxoacyl-[acyl-carrier-protein] synthase, mitoch
>gi|1037059278|ref|XP_017141789.1| PREDICTED: metastasis-associated protein MTA3 isoform X3 [Dr
>gi|1037059264|ref|XP_017141788.1| PREDICTED: metastasis-associated protein MTA3 isoform X2 [Dr
>gi|1037059249|ref|XP_017141787.1| PREDICTED: metastasis-associated protein MTA3 isoform X1 [Dr
>gi|1037059298|ref|XP_017141790.1| PREDICTED: protein transport protein Sec23A isoform X1 [Dros
>gi|1037059314|ref|XP_017141791.1| PREDICTED: protein transport protein Sec23A isoform X2 [Dros
>gi|1037059346|ref|XP_017141793.1| PREDICTED: calcineurin B homologous protein 1 [Drosophila mi
>gi|1037088023|ref|XP_017143781.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1037088010|ref|XP_017143780.1| PREDICTED: gametogenetin-binding protein 2-like [Drosophila
>gi|1037072460|ref|XP_017142682.1| PREDICTED: DNA damage-binding protein 1 [Drosophila miranda]
>gi|1037072523|ref|XP_017142687.1| PREDICTED: nuclear envelope phosphatase-regulatory subunit
>gi|1037072492|ref|XP_017142685.1| PREDICTED: protein single-minded isoform X2 [Drosophila mira
>gi|1037072476|ref|XP_017142684.1| PREDICTED: protein single-minded isoform X1 [Drosophila mira
>gi|1037072507|ref|XP_017142686.1| PREDICTED: protein single-minded isoform X3 [Drosophila mira
>gi|1037036939|ref|XP_017140297.1| PREDICTED: uncharacterized protein LOC108154508 [Drosophila
>gi|1037077147|ref|XP_017143003.1| PREDICTED: protein timeless homolog [Drosophila miranda]
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>gi|1037077162|ref|XP_017143004.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037106471|ref|XP_017145501.1| PREDICTED: uncharacterized protein C4orf22 homolog [Drosophi
>gi|1037034895|ref|XP_017140185.1| PREDICTED: uncharacterized protein LOC108154427 [Drosophila
>gi|1037105426|ref|XP_017145356.1| PREDICTED: peptidase inhibitor 15 [Drosophila miranda]
>gi|1037035029|ref|XP_017140193.1| PREDICTED: chaperone protein DnaJ [Drosophila miranda]
>gi|1037090391|ref|XP_017143970.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1037090403|ref|XP_017143971.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1037090417|ref|XP_017143972.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1037098316|ref|XP_017144603.1| PREDICTED: uncharacterized protein LOC108157186 [Drosophila
>gi|1037098304|ref|XP_017144602.1| PREDICTED: bifunctional lysine-specific demethylase and hist
>gi|1037062844|ref|XP_017142036.1| PREDICTED: protein suppressor of variegation 3-7 [Drosophila
>gi|1037062880|ref|XP_017142040.1| PREDICTED: TBC1 domain family member 5 [Drosophila miranda]
>gi|1037062856|ref|XP_017142037.1| PREDICTED: ankyrin repeat and sterile alpha motif domain-cor
>gi|1037062868|ref|XP_017142038.1| PREDICTED: ankyrin repeat and sterile alpha motif domain-cor
>gi|1037094035|ref|XP_017144229.1| PREDICTED: uncharacterized protein LOC108156962 [Drosophila
>gi|1037103165|ref|XP_017145054.1| PREDICTED: non-canonical poly(A) RNA polymerase PAPD5 [Drosc
>gi|1037055977|ref|XP_017141561.1| PREDICTED: oligopeptidase A [Drosophila miranda]
>gi|1037055962|ref|XP_017141560.1| PREDICTED: oxysterol-binding protein 1 [Drosophila miranda]
>gi|1037055869|ref|XP_017141554.1| PREDICTED: protein phosphatase Slingshot isoform X2 [Drosopk
>gi|1037055855|ref|XP_017141553.1| PREDICTED: protein phosphatase Slingshot isoform X1 [Drosopk
>gi|1037055885|ref|XP_017141555.1| PREDICTED: protein phosphatase Slingshot isoform X3 [Drosopk
>gi|1037055947|ref|XP_017141559.1| PREDICTED: protein phosphatase Slingshot isoform X7 [Drosopk
>gi|1037055919|ref|XP_017141557.1| PREDICTED: protein phosphatase Slingshot isoform X5 [Drosopk
>gi|1037055903|ref|XP_017141556.1| PREDICTED: protein phosphatase Slingshot isoform X4 [Drosopk
>gi|1037055931|ref|XP_017141558.1| PREDICTED: protein phosphatase Slingshot isoform X6 [Drosopk
>gi|1037055993|ref|XP_017141562.1| PREDICTED: nicotinamide/nicotinic acid mononucleotide adenyl
>gi|1037056011|ref|XP_017141564.1| PREDICTED: nicotinamide/nicotinic acid mononucleotide adenyl
>gi|1037035735|ref|XP_017140225.1| PREDICTED: sodium-coupled neutral amino acid transporter 1
>gi|1037071297|ref|XP_017142603.1| PREDICTED: mucin-2 [Drosophila miranda]
>gi|1037071326|ref|XP_017142606.1| PREDICTED: 26S protease regulatory subunit 8-like [Drosophil
>gi|1037071342|ref|XP_017142607.1| PREDICTED: beta-1,4-galactosyltransferase 7 [Drosophila mira
>gi|1037071358|ref|XP_017142608.1| PREDICTED: ER membrane protein complex subunit 6 [Drosophila
>gi|1037071312|ref|XP_017142605.1| PREDICTED: nuclear hormone receptor HR96 [Drosophila miranda
>gi|1037083235|ref|XP_017143446.1| PREDICTED: histone transcription regulator 3 homolog isoforr
>gi|1037083249|ref|XP_017143447.1| PREDICTED: uncharacterized protein LOC108156480 isoform X2
>gi|1037083221|ref|XP_017143445.1| PREDICTED: activin receptor type-2B [Drosophila miranda]
>gi|1037081769|ref|XP_017143340.1| PREDICTED: histone H4 [Drosophila miranda]
>gi|1037081727|ref|XP_017143337.1| PREDICTED: protein Cep78 homolog [Drosophila miranda]
>gi|1037081741|ref|XP_017143338.1| PREDICTED: protein Cep78 homolog [Drosophila miranda]
>gi|1037081755|ref|XP_017143339.1| PREDICTED: protein Cep78 homolog [Drosophila miranda]
>gi|1037081699|ref|XP_017143334.1| PREDICTED: cadherin-23 [Drosophila miranda]
>gi|1037081711|ref|XP_017143335.1| PREDICTED: cadherin-23 [Drosophila miranda]
>gi|1037075249|ref|XP_017142877.1| PREDICTED: uncharacterized protein LOC108156099 [Drosophila
>gi|1037075263|ref|XP_017142878.1| PREDICTED: uncharacterized protein LOC108156099 [Drosophila
>gi|1037075278|ref|XP_017142879.1| PREDICTED: uncharacterized protein LOC108156099 [Drosophila
>gi|1037075310|ref|XP_017142882.1| PREDICTED: serine protease easter isoform X2 [Drosophila mir
>gi|1037075294|ref|XP_017142881.1| PREDICTED: serine protease easter isoform X1 [Drosophila mir
>gi|1037100900|ref|XP_017144831.1| PREDICTED: protein HEXIM1 [Drosophila miranda]
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>gi|1037081107|ref|XP_017143288.1| PREDICTED: disintegrin and metalloproteinase domain-contain
>gi|1037093869|ref|XP_017144217.1| PREDICTED: autophagy-related protein 13 homolog [Drosophila
>gi|1037082611|ref|XP_017143402.1| PREDICTED: trichohyalin-like [Drosophila miranda]
>gi|1037082583|ref|XP_017143399.1| PREDICTED: dipeptidyl peptidase 3 isoform X1 [Drosophila mi
>gi|1037082597|ref|XP_017143401.1| PREDICTED: dipeptidyl peptidase 3 isoform X2 [Drosophila mi
>gi|1037086975|ref|XP_017143731.1| PREDICTED: E3 ubiquitin-protein ligase HRD1-like [Drosophila
>gi|1037086989|ref|XP_017143732.1| PREDICTED: E3 ubiquitin-protein ligase HRD1-like [Drosophila
>gi|1037087009|ref|XP_017143734.1| PREDICTED: JNK1/MAPK8-associated membrane protein-like [Dros
>gi|1037033801|ref|XP_017140119.1| PREDICTED: nuclear hormone receptor HR96-like [Drosophila mi
>gi|1037101548|ref|XP_017144883.1| PREDICTED: T-cell leukemia homeobox protein 3 [Drosophila mi
>gi|1037083713|ref|XP_017143485.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037038932|ref|XP_017140417.1| PREDICTED: uncharacterized protein LOC108154612 [Drosophila
>gi|1037106532|ref|XP_017145510.1| PREDICTED: acyl-CoA Delta(11) desaturase-like [Drosophila mi
>gi|1037037665|ref|XP_017140340.1| PREDICTED: LOW QUALITY PROTEIN: homeobox protein Hmx [Drosop
>gi|1037104963|ref|XP_017145288.1| PREDICTED: homeobox protein unplugged [Drosophila miranda]
>gi|1037063669|ref|XP_017142094.1| PREDICTED: monocarboxylate transporter 10 isoform X2 [Droso
>gi|1037063653|ref|XP_017142093.1| PREDICTED: monocarboxylate transporter 10 isoform X1 [Droso
>gi|1037063699|ref|XP_017142096.1| PREDICTED: suppressor of fused homolog isoform X1 [Drosophi
>gi|1037063715|ref|XP_017142097.1| PREDICTED: suppressor of fused homolog isoform X2 [Drosophi
>gi|1037063729|ref|XP_017142098.1| PREDICTED: actin-related protein 1 [Drosophila miranda]
>gi|1037063763|ref|XP_017142100.1| PREDICTED: ninjurin-1 isoform X2 [Drosophila miranda]
>gi|1037063797|ref|XP_017142103.1| PREDICTED: ninjurin-1 isoform X4 [Drosophila miranda]
>gi|1037063747|ref|XP_017142099.1| PREDICTED: ninjurin-1 isoform X1 [Drosophila miranda]
>gi|1037063779|ref|XP_017142101.1| PREDICTED: ninjurin-2 isoform X3 [Drosophila miranda]
>gi|1037063844|ref|XP_017142106.1| PREDICTED: ninjurin-1 isoform X7 [Drosophila miranda]
>gi|1037063829|ref|XP_017142105.1| PREDICTED: ninjurin-2 isoform X6 [Drosophila miranda]
>gi|1037063813|ref|XP_017142104.1| PREDICTED: ninjurin-2 isoform X5 [Drosophila miranda]
>gi|1037063685|ref|XP_017142095.1| PREDICTED: EH domain-containing protein 1 [Drosophila miran
>gi|1037065373|ref|XP_017142198.1| PREDICTED: uncharacterized protein LOC108155713 isoform X1
>gi|1037065389|ref|XP_017142199.1| PREDICTED: uncharacterized protein LOC108155713 isoform X2
>gi|1037065309|ref|XP_017142193.1| PREDICTED: DNA polymerase theta isoform X1 [Drosophila mira
>gi|1037065343|ref|XP_017142196.1| PREDICTED: DNA polymerase theta isoform X3 [Drosophila mira
>gi|1037065407|ref|XP_017142200.1| PREDICTED: heat shock protein 67B2 [Drosophila miranda]
>gi|1037065325|ref|XP_017142195.1| PREDICTED: DNA polymerase theta isoform X2 [Drosophila mira
>gi|1037065357|ref|XP_017142197.1| PREDICTED: kanadaptin [Drosophila miranda]
>gi|1037104404|ref|XP_017145209.1| PREDICTED: high-affinity choline transporter 1 [Drosophila
>gi|1037104414|ref|XP_017145210.1| PREDICTED: high-affinity choline transporter 1 [Drosophila
>gi|1037024375|ref|XP_017139545.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1037037151|ref|XP_017140310.1| PREDICTED: uncharacterized protein LOC108154520 [Drosophila
>gi|1037037312|ref|XP_017140319.1| PREDICTED: uncharacterized protein LOC108154527 [Drosophila
>gi|1037024393|ref|XP_017139546.1| PREDICTED: choline O-acetyltransferase, partial [Drosophila
>gi|1037098459|ref|XP_017144617.1| PREDICTED: vesicular acetylcholine transporter [Drosophila
>gi|1037098483|ref|XP_017144619.1| PREDICTED: peritrophin-55 [Drosophila miranda]
>gi|1037098471|ref|XP_017144618.1| PREDICTED: uncharacterized protein LOC108157197 [Drosophila
>gi|1037080680|ref|XP_017143254.1| PREDICTED: uncharacterized protein LOC108156362 [Drosophila
>gi|1037080635|ref|XP_017143251.1| PREDICTED: mitogen-activated protein kinase kinase kinase 4
>gi|1037080622|ref|XP_017143250.1| PREDICTED: mitogen-activated protein kinase kinase kinase 4
>gi|1037080649|ref|XP_017143252.1| PREDICTED: mitogen-activated protein kinase kinase kinase 4
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>gi|1037080662|ref|XP_017143253.1| PREDICTED: LOW QUALITY PROTEIN: CDP-diacylglycerol--glycerol
>gi|1037024409|ref|XP_017139548.1| PREDICTED: WD repeat-containing protein 20, partial [Drosoph
>gi|1037089279|ref|XP_017143879.1| PREDICTED: protein YIF1B-A isoform X4 [Drosophila miranda]
>gi|1037089265|ref|XP_017143878.1| PREDICTED: protein YIF1B-A isoform X3 [Drosophila miranda]
>gi|1037089251|ref|XP_017143877.1| PREDICTED: protein YIF1B-A isoform X2 [Drosophila miranda]
>gi|1037089237|ref|XP_017143876.1| PREDICTED: protein YIF1B-A isoform X1 [Drosophila miranda]
>gi|1037100660|ref|XP_017144803.1| PREDICTED: uncharacterized protein LOC108157310 [Drosophila
>gi|1037103858|ref|XP_017145137.1| PREDICTED: neuroendocrine convertase 2 [Drosophila miranda]
>gi|1037039371|ref|XP_017140444.1| PREDICTED: basic-leucine zipper transcription factor A [Dros
>gi|1037103607|ref|XP_017145103.1| PREDICTED: uncharacterized protein LOC108157514 [Drosophila
>gi|1037022509|ref|XP_017139425.1| PREDICTED: keratin, type II cytoskeletal 1 [Drosophila mira
>gi|1037038034|ref|XP_017140357.1| PREDICTED: uncharacterized protein LOC108154570 [Drosophila
>gi|1037024425|ref|XP_017139549.1| PREDICTED: uncharacterized protein LOC108153963 [Drosophila
>gi|1037024443|ref|XP_017139550.1| PREDICTED: uncharacterized protein LOC108153964 [Drosophila
>gi|1037024461|ref|XP_017139551.1| PREDICTED: uncharacterized protein LOC108153965 [Drosophila
>gi|1037037698|ref|XP_017140342.1| PREDICTED: glycine, alanine and asparagine-rich protein [Dro
>gi|1037038104|ref|XP_017140363.1| PREDICTED: uncharacterized protein LOC108154574 [Drosophila
>gi|1037024479|ref|XP_017139552.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037038285|ref|XP_017140376.1| PREDICTED: uncharacterized protein LOC108154584 [Drosophila
>gi|1037037502|ref|XP_017140331.1| PREDICTED: uncharacterized PPE family protein PPE54-like iso
>gi|1037037522|ref|XP_017140332.1| PREDICTED: glycine, alanine and asparagine-rich protein-like
>gi|1037037013|ref|XP_017140301.1| PREDICTED: LOW QUALITY PROTEIN: glycine, alanine and asparag
>gi|1037036852|ref|XP_017140291.1| PREDICTED: uncharacterized protein LOC108154503 [Drosophila
>gi|1037034646|ref|XP_017140173.1| PREDICTED: LOW QUALITY PROTEIN: glycine, alanine and asparag
>gi|1037050200|ref|XP_017141170.1| PREDICTED: protein lap4 isoform X9 [Drosophila miranda]
>gi|1037050149|ref|XP_017141166.1| PREDICTED: protein lap4 isoform X5 [Drosophila miranda]
>gi|1037050109|ref|XP_017141163.1| PREDICTED: protein lap4 isoform X2 [Drosophila miranda]
>gi|1037050135|ref|XP_017141165.1| PREDICTED: protein lap4 isoform X4 [Drosophila miranda]
>gi|1037050186|ref|XP_017141169.1| PREDICTED: protein lap4 isoform X8 [Drosophila miranda]
>gi|1037050174|ref|XP_017141168.1| PREDICTED: protein lap4 isoform X7 [Drosophila miranda]
>gi|1037050162|ref|XP_017141167.1| PREDICTED: protein lap4 isoform X6 [Drosophila miranda]
>gi|1037050123|ref|XP_017141164.1| PREDICTED: protein lap4 isoform X3 [Drosophila miranda]
>gi|1037050066|ref|XP_017141161.1| PREDICTED: protein lap4 isoform X1 [Drosophila miranda]
>gi|1037050080|ref|XP_017141162.1| PREDICTED: protein lap4 isoform X1 [Drosophila miranda]
>gi|1037050227|ref|XP_017141173.1| PREDICTED: uncharacterized protein LOC108155077 isoform X11
>gi|1037050213|ref|XP_017141172.1| PREDICTED: uncharacterized protein LOC108155077 isoform X10
>gi|1037050257|ref|XP_017141175.1| PREDICTED: FK506-binding protein 5 isoform X13 [Drosophila m
>gi|1037050241|ref|XP_017141174.1| PREDICTED: protein lap4 isoform X12 [Drosophila miranda]
>gi|1037106424|ref|XP_017145495.1| PREDICTED: uncharacterized protein LOC108157809 [Drosophila
>gi|1037024495|ref|XP_017139553.1| PREDICTED: putative odorant receptor 98b [Drosophila miranda
>gi|1037105446|ref|XP_017145360.1| PREDICTED: uncharacterized protein LOC108157701 [Drosophila
>gi|1037105454|ref|XP_017145361.1| PREDICTED: uncharacterized protein LOC108157701 [Drosophila
>gi|1037098961|ref|XP_017144663.1| PREDICTED: dynein heavy chain 10, axonemal [Drosophila mira
>gi|1037098973|ref|XP_017144664.1| PREDICTED: uncharacterized protein LOC108157218 [Drosophila
>gi|1037098985|ref|XP_017144665.1| PREDICTED: uncharacterized protein LOC108157218 [Drosophila
>gi|1037085336|ref|XP_017143608.1| PREDICTED: papilin isoform X5 [Drosophila miranda]
>gi|1037085320|ref|XP_017143607.1| PREDICTED: papilin isoform X4 [Drosophila miranda]
>gi|1037085294|ref|XP_017143605.1| PREDICTED: papilin isoform X2 [Drosophila miranda]
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>gi|1037085306|ref|XP_017143606.1| PREDICTED: papilin isoform X3 [Drosophila miranda]
>gi|1037085282|ref|XP_017143604.1| PREDICTED: papilin isoform X1 [Drosophila miranda]
>gi|1037092132|ref|XP_017144101.1| PREDICTED: LOW QUALITY PROTEIN: translation initiation facto
>gi|1037092120|ref|XP_017144100.1| PREDICTED: sarcalumenin isoform X2 [Drosophila miranda]
>gi|1037092092|ref|XP_017144098.1| PREDICTED: sarcalumenin isoform X1 [Drosophila miranda]
>gi|1037092106|ref|XP_017144099.1| PREDICTED: sarcalumenin isoform X1 [Drosophila miranda]
>gi|1037057061|ref|XP_017141634.1| PREDICTED: endochitinase [Drosophila miranda]
>gi|1037057075|ref|XP_017141635.1| PREDICTED: uncharacterized protein LOC108155374 [Drosophila
>gi|1037057016|ref|XP_017141630.1| PREDICTED: uncharacterized protein LOC108155371 isoform X6
>gi|1037056969|ref|XP_017141627.1| PREDICTED: uncharacterized protein LOC108155371 isoform X3
>gi|1037056951|ref|XP_017141626.1| PREDICTED: uncharacterized protein LOC108155371 isoform X2
>gi|1037057032|ref|XP_017141632.1| PREDICTED: uncharacterized protein LOC108155371 isoform X7
>gi|1037057000|ref|XP_017141629.1| PREDICTED: uncharacterized protein LOC108155371 isoform X5
>gi|1037056985|ref|XP_017141628.1| PREDICTED: uncharacterized protein LOC108155371 isoform X4
>gi|1037057046|ref|XP_017141633.1| PREDICTED: uncharacterized protein LOC108155371 isoform X8
>gi|1037056933|ref|XP_017141625.1| PREDICTED: uncharacterized protein LOC108155371 isoform X1
>gi|1037057089|ref|XP_017141636.1| PREDICTED: ubiquitin-conjugating enzyme E2 G1 [Drosophila m
>gi|1037102223|ref|XP_017144955.1| PREDICTED: LOW QUALITY PROTEIN: low molecular weight phospho
>gi|1037102213|ref|XP_017144954.1| PREDICTED: low molecular weight phosphotyrosine protein phos
>gi|1037102205|ref|XP_017144953.1| PREDICTED: low molecular weight phosphotyrosine protein phos
>gi|1037102195|ref|XP_017144951.1| PREDICTED: adenosine deaminase CECR1 isoform X2 [Drosophila
>gi|1037102185|ref|XP_017144950.1| PREDICTED: adenosine deaminase CECR1 isoform X1 [Drosophila
>gi|1037091196|ref|XP_017144030.1| PREDICTED: adenosine deaminase CECR1 [Drosophila miranda]
>gi|1037091182|ref|XP_017144029.1| PREDICTED: carotenoid isomerooxygenase [Drosophila miranda]
>gi|1037091061|ref|XP_017144021.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091075|ref|XP_017144022.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091103|ref|XP_017144024.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091143|ref|XP_017144026.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091156|ref|XP_017144027.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091089|ref|XP_017144023.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091130|ref|XP_017144025.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037091168|ref|XP_017144028.1| PREDICTED: leucine-rich repeat-containing protein 40 isoform
>gi|1037034542|ref|XP_017140166.1| PREDICTED: cilia- and flagella-associated protein 36 isoform
>gi|1037034526|ref|XP_017140165.1| PREDICTED: cilia- and flagella-associated protein 36 isoform
>gi|1037040949|ref|XP_017140555.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1037041258|ref|XP_017140577.1| PREDICTED: uncharacterized protein LOC108154725 [Drosophila
>gi|1037041274|ref|XP_017140578.1| PREDICTED: uncharacterized protein LOC108154726 [Drosophila
>gi|1037040990|ref|XP_017140559.1| PREDICTED: transmembrane protein 245 isoform X1 [Drosophila
>gi|1037041004|ref|XP_017140560.1| PREDICTED: transmembrane protein 245 isoform X2 [Drosophila
>gi|1037041018|ref|XP_017140561.1| PREDICTED: transmembrane protein 245 isoform X3 [Drosophila
>gi|1037041213|ref|XP_017140574.1| PREDICTED: alcohol dehydrogenase [NADP(+)] [Drosophila mira
>gi|1037041242|ref|XP_017140575.1| PREDICTED: uncharacterized protein LOC108154724 [Drosophila
>gi|1037041200|ref|XP_017140573.1| PREDICTED: tRNA dimethylallyltransferase, mitochondrial [Dro
>gi|1037041163|ref|XP_017140570.1| PREDICTED: bifunctional purine biosynthesis protein PURH iso
>gi|1037041179|ref|XP_017140571.1| PREDICTED: bifunctional purine biosynthesis protein PURH iso
>gi|1037040963|ref|XP_017140556.1| PREDICTED: ATP-binding cassette sub-family G member 8 isofor
>gi|1037040975|ref|XP_017140557.1| PREDICTED: ATP-binding cassette sub-family G member 8 isofor
>gi|1037041117|ref|XP_017140566.1| PREDICTED: ATP-binding cassette sub-family G member 5 [Droso
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>gi|1037041132|ref|XP_017140567.1| PREDICTED: ATP-binding cassette sub-family G member 5 [Dros
>gi|1037041147|ref|XP_017140569.1| PREDICTED: ATP-binding cassette sub-family G member 5 [Dros
>gi|1037041082|ref|XP_017140565.1| PREDICTED: mucin-5AC isoform X4 [Drosophila miranda]
>gi|1037041066|ref|XP_017140564.1| PREDICTED: mucin-5AC isoform X3 [Drosophila miranda]
>gi|1037041048|ref|XP_017140563.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1037041034|ref|XP_017140562.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037041187|ref|XP_017140572.1| PREDICTED: uncharacterized protein LOC108154721 [Drosophila
>gi|1037089855|ref|XP_017143925.1| PREDICTED: GPI transamidase component PIG-S-like isoform X2
>gi|1037089844|ref|XP_017143924.1| PREDICTED: prolyl 3-hydroxylase sudestada1-like isoform X1
>gi|1037104396|ref|XP_017145208.1| PREDICTED: stearoyl-CoA desaturase 5 [Drosophila miranda]
>gi|1037024516|ref|XP_017139554.1| PREDICTED: uncharacterized protein LOC108153968 [Drosophila
>gi|1037039974|ref|XP_017140488.1| PREDICTED: uncharacterized protein LOC108154662 [Drosophila
>gi|1037034993|ref|XP_017140191.1| PREDICTED: uncharacterized protein LOC108154432 [Drosophila
>gi|1037096182|ref|XP_017144402.1| PREDICTED: wnt inhibitor of Dorsal protein [Drosophila mirar
>gi|1037096194|ref|XP_017144403.1| PREDICTED: glutamyl aminopeptidase [Drosophila miranda]
>gi|1037096170|ref|XP_017144401.1| PREDICTED: glutamyl aminopeptidase [Drosophila miranda]
>gi|1037084607|ref|XP_017143556.1| PREDICTED: glutamyl aminopeptidase isoform X2 [Drosophila mi
>gi|1037084621|ref|XP_017143557.1| PREDICTED: uncharacterized protein LOC108156549 [Drosophila
>gi|1037084591|ref|XP_017143555.1| PREDICTED: glutamyl aminopeptidase isoform X1 [Drosophila mi
>gi|1037024533|ref|XP_017139555.1| PREDICTED: beta-1,3-galactosyltransferase 1-like [Drosophila
>gi|1037035906|ref|XP_017140237.1| PREDICTED: uncharacterized protein LOC108154474 [Drosophila
>gi|1037104784|ref|XP_017145262.1| PREDICTED: neuromedin-U receptor 1 [Drosophila miranda]
>gi|1037104792|ref|XP_017145263.1| PREDICTED: neuromedin-U receptor 1 [Drosophila miranda]
>gi|1037105802|ref|XP_017145409.1| PREDICTED: neurotensin receptor type 1 isoform X1 [Drosophil
>gi|1037105808|ref|XP_017145410.1| PREDICTED: neuromedin-U receptor 2 isoform X2 [Drosophila mi
>gi|1037105816|ref|XP_017145411.1| PREDICTED: neuromedin-U receptor 2 isoform X3 [Drosophila mi
>gi|1037105826|ref|XP_017145412.1| PREDICTED: neuromedin-U receptor 2 isoform X4 [Drosophila mi
>gi|1037094910|ref|XP_017144296.1| PREDICTED: uncharacterized protein LOC108157007 [Drosophila
>gi|1037094923|ref|XP_017144297.1| PREDICTED: uncharacterized protein LOC108157007 [Drosophila
>gi|1037094936|ref|XP_017144298.1| PREDICTED: uncharacterized protein LOC108157007 [Drosophila
>gi|1037094883|ref|XP_017144294.1| PREDICTED: mitochondrial dicarboxylate carrier [Drosophila n
>gi|1037094897|ref|XP_017144295.1| PREDICTED: mitochondrial dicarboxylate carrier [Drosophila n
>gi|1037033576|ref|XP_017140105.1| PREDICTED: uncharacterized protein LOC108154351 [Drosophila
>gi|1037033559|ref|XP_017140104.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1037039396|ref|XP_017140445.1| PREDICTED: uncharacterized protein LOC108154633 [Drosophila
>gi|1037101113|ref|XP_017144851.1| PREDICTED: uncharacterized protein LOC108157342 isoform X3
>gi|1037101103|ref|XP_017144850.1| PREDICTED: probable replication factor C subunit 1 isoform X
>gi|1037101095|ref|XP_017144849.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037030939|ref|XP_017139949.1| PREDICTED: 28S ribosomal protein S21, mitochondrial [Drosoph
>gi|1037078305|ref|XP_017143094.1| PREDICTED: dnaJ homolog subfamily A member 4 [Drosophila mir
>gi|1037078291|ref|XP_017143093.1| PREDICTED: WD repeat-containing protein 36 [Drosophila mirar
>gi|1037104592|ref|XP_017145235.1| PREDICTED: neuropeptide CCHamide-2 [Drosophila miranda]
>gi|1037104600|ref|XP_017145236.1| PREDICTED: neuropeptide CCHamide-2 [Drosophila miranda]
>gi|1037104606|ref|XP_017145237.1| PREDICTED: neuropeptide CCHamide-2 [Drosophila miranda]
>gi|1037104614|ref|XP_017145238.1| PREDICTED: neuropeptide CCHamide-2 [Drosophila miranda]
>gi|1037104626|ref|XP_017145239.1| PREDICTED: neuropeptide CCHamide-2 [Drosophila miranda]
>gi|1037097027|ref|XP_017144477.1| PREDICTED: GILT-like protein C02D5.2 isoform X2 [Drosophila
>gi|1037097015|ref|XP_017144476.1| PREDICTED: GILT-like protein C02D5.2 isoform X1 [Drosophila
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>gi|1037030906|ref|XP_017139947.1| PREDICTED: dolichol-phosphate mannosyltransferase subunit 3
>gi|1037084157|ref|XP_017143521.1| PREDICTED: uncharacterized protein LOC108156527 isoform X1
>gi|1037084171|ref|XP_017143522.1| PREDICTED: coiled-coil domain-containing protein 149 isoform
>gi|1037024551|ref|XP_017139556.1| PREDICTED: copper homeostasis protein cutC homolog, partial
>gi|1037084187|ref|XP_017143523.1| PREDICTED: programmed cell death protein 10 [Drosophila mira
>gi|1037096414|ref|XP_017144423.1| PREDICTED: uncharacterized protein LOC108157085 isoform X1
>gi|1037096426|ref|XP_017144424.1| PREDICTED: uncharacterized protein LOC108157085 isoform X2
>gi|1037096513|ref|XP_017144432.1| PREDICTED: uncharacterized protein LOC108157085 isoform X9
>gi|1037096525|ref|XP_017144433.1| PREDICTED: uncharacterized protein LOC108157085 isoform X10
>gi|1037096491|ref|XP_017144430.1| PREDICTED: uncharacterized protein LOC108157085 isoform X7
>gi|1037096442|ref|XP_017144425.1| PREDICTED: uncharacterized protein LOC108157085 isoform X3
>gi|1037096466|ref|XP_017144427.1| PREDICTED: uncharacterized protein LOC108157085 isoform X5
>gi|1037096548|ref|XP_017144435.1| PREDICTED: uncharacterized protein LOC108157085 isoform X12
>gi|1037096479|ref|XP_017144429.1| PREDICTED: uncharacterized protein LOC108157085 isoform X6
>gi|1037096560|ref|XP_017144436.1| PREDICTED: uncharacterized protein LOC108157085 isoform X13
>gi|1037096536|ref|XP_017144434.1| PREDICTED: uncharacterized protein LOC108157085 isoform X11
>gi|1037096454|ref|XP_017144426.1| PREDICTED: uncharacterized protein LOC108157085 isoform X4
>gi|1037096503|ref|XP_017144431.1| PREDICTED: uncharacterized protein LOC108157085 isoform X8
>gi|1037096570|ref|XP_017144437.1| PREDICTED: uncharacterized protein LOC108157086 [Drosophila
>gi|1037096582|ref|XP_017144440.1| PREDICTED: uncharacterized protein LOC108157087 [Drosophila
>gi|1037104297|ref|XP_017145194.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1037104289|ref|XP_017145193.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1037076489|ref|XP_017142960.1| PREDICTED: ataxin-2 homolog isoform X1 [Drosophila miranda]
>gi|1037076505|ref|XP_017142961.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1037076521|ref|XP_017142963.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1037076537|ref|XP_017142964.1| PREDICTED: broad-complex core protein [Drosophila miranda]
>gi|1037048747|ref|XP_017141070.1| PREDICTED: LOW QUALITY PROTEIN: protein split ends [Drosophi
>gi|1037048731|ref|XP_017141069.1| PREDICTED: endoplasmic reticulum mannosyl-oligosaccharide 1
>gi|1037065897|ref|XP_017142233.1| PREDICTED: RUN and FYVE domain-containing protein 2 isoform
>gi|1037065833|ref|XP_017142229.1| PREDICTED: RUN and FYVE domain-containing protein 2 isoform
>gi|1037065849|ref|XP_017142230.1| PREDICTED: RUN and FYVE domain-containing protein 2 isoform
>gi|1037065911|ref|XP_017142234.1| PREDICTED: protein RUFY3 isoform X5 [Drosophila miranda]
>gi|1037065881|ref|XP_017142232.1| PREDICTED: protein RUFY3 isoform X3 [Drosophila miranda]
>gi|1037065865|ref|XP_017142231.1| PREDICTED: uncharacterized protein LOC108155731 isoform X2
>gi|1037065943|ref|XP_017142236.1| PREDICTED: protein RUFY3 isoform X7 [Drosophila miranda]
>gi|1037065927|ref|XP_017142235.1| PREDICTED: protein RUFY3 isoform X6 [Drosophila miranda]
>gi|1037065957|ref|XP_017142237.1| PREDICTED: uncharacterized protein LOC108155733 isoform X1
>gi|1037065971|ref|XP_017142238.1| PREDICTED: uncharacterized protein LOC108155733 isoform X2
>gi|1037065986|ref|XP_017142239.1| PREDICTED: DNA polymerase alpha subunit B [Drosophila miran
>gi|1037066001|ref|XP_017142240.1| PREDICTED: uncharacterized protein LOC108155735 [Drosophila
>gi|1037066016|ref|XP_017142241.1| PREDICTED: uncharacterized protein LOC108155735 [Drosophila
>gi|1037034796|ref|XP_017140181.1| PREDICTED: filamin-B [Drosophila miranda]
>gi|1037087634|ref|XP_017143750.1| PREDICTED: tubulin polyglutamylase TTLL13 [Drosophila miran
>gi|1037087648|ref|XP_017143752.1| PREDICTED: serine/threonine-protein kinase pelle [Drosophila
>gi|1037087662|ref|XP_017143753.1| PREDICTED: transcription initiation factor IIA subunit 1 iso
>gi|1037087675|ref|XP_017143754.1| PREDICTED: transcription initiation factor IIA subunit 1 iso
>gi|1037087689|ref|XP_017143755.1| PREDICTED: transcription initiation factor IIA subunit 1 iso
>gi|1037041847|ref|XP_017140614.1| PREDICTED: zinc finger MYM-type protein 4 isoform X2 [Drosop
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>gi|1037041817|ref|XP_017140612.1| PREDICTED: zinc finger MYM-type protein 4 isoform X1 [Drosop
>gi|1037041831|ref|XP_017140613.1| PREDICTED: zinc finger MYM-type protein 4 isoform X1 [Drosop
>gi|1037041969|ref|XP_017140621.1| PREDICTED: uncharacterized protein LOC108154758 [Drosophila
>gi|1037041910|ref|XP_017140618.1| PREDICTED: uncharacterized protein LOC108154754 [Drosophila
>gi|1037041893|ref|XP_017140617.1| PREDICTED: uncharacterized protein LOC108154753 [Drosophila
>gi|1037041801|ref|XP_017140611.1| PREDICTED: uncharacterized protein LOC108154750 [Drosophila
>gi|1037041940|ref|XP_017140620.1| PREDICTED: short coiled-coil protein B [Drosophila miranda]
>gi|1037041924|ref|XP_017140619.1| PREDICTED: cysteine-rich hydrophobic domain-containing prote
>gi|1037041861|ref|XP_017140615.1| PREDICTED: formin-like protein 13 isoform X1 [Drosophila mi
>gi|1037041879|ref|XP_017140616.1| PREDICTED: splicing factor, arginine/serine-rich 15 isoform
>gi|1037032920|ref|XP_017140067.1| PREDICTED: probable V-type proton ATPase subunit d 2 [Drosop
>gi|1037093105|ref|XP_017144165.1| PREDICTED: uncharacterized protein LOC108156922 [Drosophila
>gi|1037093032|ref|XP_017144160.1| PREDICTED: G protein-activated inward rectifier potassium cl
>gi|1037093046|ref|XP_017144161.1| PREDICTED: G protein-activated inward rectifier potassium cl
>gi|1037093089|ref|XP_017144164.1| PREDICTED: G protein-activated inward rectifier potassium cl
>gi|1037093060|ref|XP_017144162.1| PREDICTED: G protein-activated inward rectifier potassium cl
>gi|1037093075|ref|XP_017144163.1| PREDICTED: G protein-activated inward rectifier potassium cl
>gi|1037071089|ref|XP_017142591.1| PREDICTED: segmentation protein cap'n'collar isoform X2 [Dr
>gi|1037071073|ref|XP_017142590.1| PREDICTED: segmentation protein cap'n'collar isoform X1 [Dr
>gi|1037071107|ref|XP_017142592.1| PREDICTED: segmentation protein cap'n'collar isoform X3 [Dr
>gi|1037071123|ref|XP_017142594.1| PREDICTED: segmentation protein cap'n'collar isoform X4 [Dr
>gi|1037099494|ref|XP_017144707.1| PREDICTED: endoplasmic reticulum aminopeptidase 2 isoform X2
>gi|1037099482|ref|XP_017144706.1| PREDICTED: endoplasmic reticulum aminopeptidase 2 isoform X1
>gi|1037099505|ref|XP_017144708.1| PREDICTED: endoplasmic reticulum aminopeptidase 2 isoform X3
>gi|1037099517|ref|XP_017144709.1| PREDICTED: endoplasmic reticulum aminopeptidase 2 isoform X3
>gi|1037099530|ref|XP_017144710.1| PREDICTED: endoplasmic reticulum aminopeptidase 2 isoform X3
>gi|1037100338|ref|XP_017144774.1| PREDICTED: uncharacterized protein CG4449 [Drosophila mirand
>gi|1037077132|ref|XP_017143002.1| PREDICTED: transcription elongation factor B polypeptide 3
>gi|1037077117|ref|XP_017143001.1| PREDICTED: TAF5-like RNA polymerase II p300/CBP-associated i
>gi|1037101919|ref|XP_017144923.1| PREDICTED: peroxisomal membrane protein 11C [Drosophila mira
>gi|1037090363|ref|XP_017143967.1| PREDICTED: uncharacterized protein LOC108156798 isoform X2
>gi|1037090377|ref|XP_017143968.1| PREDICTED: uncharacterized protein LOC108156798 isoform X2
>gi|1037090349|ref|XP_017143966.1| PREDICTED: uncharacterized protein LOC108156798 isoform X1
>gi|1037086863|ref|XP_017143722.1| PREDICTED: cell division cycle protein 16 homolog [Drosophil
>gi|1037086877|ref|XP_017143723.1| PREDICTED: low choriolytic enzyme [Drosophila miranda]
>gi|1037069992|ref|XP_017142516.1| PREDICTED: uncharacterized protein LOC108155887 [Drosophila
>gi|1037070008|ref|XP_017142517.1| PREDICTED: glutamate dehydrogenase, mitochondrial [Drosophil
>gi|1037069926|ref|XP_017142512.1| PREDICTED: ETS-like protein pointed isoform X1 [Drosophila m
>gi|1037069976|ref|XP_017142515.1| PREDICTED: ETS-like protein pointed isoform X4 [Drosophila m
>gi|1037069942|ref|XP_017142513.1| PREDICTED: ETS-like protein pointed isoform X2 [Drosophila m
>gi|1037069958|ref|XP_017142514.1| PREDICTED: ETS-like protein pointed isoform X3 [Drosophila m
>gi|1037080224|ref|XP_017143220.1| PREDICTED: phosphatidylinositide phosphatase SAC2 isoform X1
>gi|1037080251|ref|XP_017143222.1| PREDICTED: phosphatidylinositide phosphatase SAC2 isoform X3
>gi|1037080238|ref|XP_017143221.1| PREDICTED: phosphatidylinositide phosphatase SAC2 isoform X2
>gi|1037080210|ref|XP_017143219.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1037035806|ref|XP_017140230.1| PREDICTED: ficolin-1-like [Drosophila miranda]
>gi|1037102231|ref|XP_017144956.1| PREDICTED: uncharacterized protein LOC108157416 [Drosophila
>gi|1037033734|ref|XP_017140114.1| PREDICTED: homeobox protein unc-4-like [Drosophila miranda]
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>gi|1037101732|ref|XP_017144903.1| PREDICTED: uncharacterized protein LOC108157383 [Drosophila
>gi|1037101722|ref|XP_017144901.1| PREDICTED: anaphase-promoting complex subunit 4-like [Drosop
>gi|1037052530|ref|XP_017141331.1| PREDICTED: transcription factor IIIB 90 kDa subunit [Drosoph
>gi|1037052484|ref|XP_017141328.1| PREDICTED: BTB/POZ domain-containing protein 6 isoform X1 [D
>gi|1037052499|ref|XP_017141329.1| PREDICTED: BTB/POZ domain-containing protein 6 isoform X1 [D
>gi|1037052514|ref|XP_017141330.1| PREDICTED: BTB/POZ domain-containing protein 6 isoform X2 [D
>gi|1037052559|ref|XP_017141334.1| PREDICTED: protein phosphatase 1 regulatory subunit 7 [Droso
>gi|1037052575|ref|XP_017141335.1| PREDICTED: protein CREG1 [Drosophila miranda]
>gi|1037052545|ref|XP_017141333.1| PREDICTED: leucine-rich repeat protein soc-2 homolog [Drosop
>gi|1037034613|ref|XP_017140171.1| PREDICTED: uncharacterized protein LOC108154410 [Drosophila
>gi|1037034630|ref|XP_017140172.1| PREDICTED: uncharacterized protein LOC108154411 [Drosophila
>gi|1037104213|ref|XP_017145183.1| PREDICTED: salivary glue protein Sgs-5-like [Drosophila mira
>gi|1037104229|ref|XP_017145185.1| PREDICTED: uncharacterized protein LOC108157575 [Drosophila
>gi|1037104221|ref|XP_017145184.1| PREDICTED: salivary glue protein Sgs-5-like [Drosophila mira
>gi|1037104205|ref|XP_017145182.1| PREDICTED: salivary glue protein Sgs-5-like [Drosophila mira
>gi|1037104237|ref|XP_017145186.1| PREDICTED: vasotab [Drosophila miranda]
>gi|1037105346|ref|XP_017145344.1| PREDICTED: lysosomal aspartic protease [Drosophila miranda]
>gi|1037105354|ref|XP_017145345.1| PREDICTED: aspartic proteinase A1 [Drosophila miranda]
>gi|1037106752|ref|XP_017145539.1| PREDICTED: lysosomal aspartic protease [Drosophila miranda]
>gi|1037106759|ref|XP_017145541.1| PREDICTED: uncharacterized protein LOC108157848 [Drosophila
>gi|1037037937|ref|XP_017140354.1| PREDICTED: uncharacterized protein LOC108154563 isoform X2
>gi|1037037907|ref|XP_017140353.1| PREDICTED: uncharacterized protein LOC108154563 isoform X1
>gi|1037037466|ref|XP_017140329.1| PREDICTED: uncharacterized protein LOC108154537 [Drosophila
>gi|1037105925|ref|XP_017145425.1| PREDICTED: uncharacterized protein LOC108157756 [Drosophila
>gi|1037105917|ref|XP_017145424.1| PREDICTED: uncharacterized protein LOC108157755 [Drosophila
>gi|1037037748|ref|XP_017140345.1| PREDICTED: uncharacterized protein LOC108154553 [Drosophila
>gi|1037037344|ref|XP_017140321.1| PREDICTED: uncharacterized protein LOC108154529 [Drosophila
>gi|1037102582|ref|XP_017144989.1| PREDICTED: peroxidase [Drosophila miranda]
>gi|1037102626|ref|XP_017144994.1| PREDICTED: turripeptide Ici9.2 [Drosophila miranda]
>gi|1037102614|ref|XP_017144993.1| PREDICTED: uncharacterized protein LOC108157447 [Drosophila
>gi|1037102604|ref|XP_017144992.1| PREDICTED: uncharacterized protein LOC108157446 isoform X2
>gi|1037102594|ref|XP_017144991.1| PREDICTED: uncharacterized protein LOC108157446 isoform X1
>gi|1037038120|ref|XP_017140364.1| PREDICTED: eclosion hormone [Drosophila miranda]
>gi|1037038805|ref|XP_017140409.1| PREDICTED: uncharacterized protein LOC108154603 [Drosophila
>gi|1037104047|ref|XP_017145160.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104023|ref|XP_017145157.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104039|ref|XP_017145159.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037103997|ref|XP_017145154.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104005|ref|XP_017145155.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104013|ref|XP_017145156.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104063|ref|XP_017145162.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104055|ref|XP_017145161.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037104031|ref|XP_017145158.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1037106697|ref|XP_017145532.1| PREDICTED: uncharacterized protein LOC108157840 [Drosophila
>gi|1037097893|ref|XP_017144559.1| PREDICTED: ubiquitin-conjugating enzyme E2 J2 [Drosophila mi
>gi|1037097905|ref|XP_017144560.1| PREDICTED: ubiquitin-conjugating enzyme E2 J2 [Drosophila mi
>gi|1037030816|ref|XP_017139941.1| PREDICTED: uncharacterized protein LOC108154226 [Drosophila
>gi|1037062073|ref|XP_017141983.1| PREDICTED: pyrroline-5-carboxylate reductase 3 [Drosophila m
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>gi|1037062010|ref|XP_017141978.1| PREDICTED: nucleolar protein 14 homolog [Drosophila miranda]
>gi|1037062026|ref|XP_017141979.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037062042|ref|XP_017141980.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037062057|ref|XP_017141981.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037032851|ref|XP_017140063.1| PREDICTED: solute carrier family 25 member 45 [Drosophila mi
>gi|1037079181|ref|XP_017143153.1| PREDICTED: kinesin-like protein Klp98A isoform X1 [Drosophil
>gi|1037079195|ref|XP_017143154.1| PREDICTED: kinesin-like protein Klp98A isoform X2 [Drosophil
>gi|1037024569|ref|XP_017139557.1| PREDICTED: putative gustatory receptor 98b [Drosophila mirar
>gi|1037105300|ref|XP_017145337.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1037024585|ref|XP_017139559.1| PREDICTED: IQ domain-containing protein G [Drosophila mirand
>gi|1037034953|ref|XP_017140188.1| PREDICTED: uncharacterized protein LOC108154431 isoform X1
>gi|1037034969|ref|XP_017140189.1| PREDICTED: uncharacterized protein LOC108154431 isoform X2
>gi|1037024601|ref|XP_017139560.1| PREDICTED: putative gustatory receptor 98a [Drosophila mirar
>gi|1037024617|ref|XP_017139561.1| PREDICTED: kelch-like protein 2 [Drosophila miranda]
>gi|1037105308|ref|XP_017145338.1| PREDICTED: GPI ethanolamine phosphate transferase 1 [Drosop]
>gi|1037094965|ref|XP_017144300.1| PREDICTED: uncharacterized protein LOC108157009 isoform X2
>gi|1037094951|ref|XP_017144299.1| PREDICTED: uncharacterized protein LOC108157009 isoform X1
>gi|1037084291|ref|XP_017143531.1| PREDICTED: uncharacterized protein LOC108156534 [Drosophila
>gi|1037084305|ref|XP_017143532.1| PREDICTED: uncharacterized protein LOC108156535 [Drosophila
>gi|1037084277|ref|XP_017143529.1| PREDICTED: dynein beta chain, ciliary [Drosophila miranda]
>gi|1037084253|ref|XP_017143528.1| PREDICTED: dynein beta chain, ciliary [Drosophila miranda]
>gi|1037092038|ref|XP_017144093.1| PREDICTED: uncharacterized protein LOC108156875 [Drosophila
>gi|1037092024|ref|XP_017144092.1| PREDICTED: adipocyte plasma membrane-associated protein [Dro
>gi|1037089030|ref|XP_017143859.1| PREDICTED: max-like protein X [Drosophila miranda]
>gi|1037089017|ref|XP_017143858.1| PREDICTED: uncharacterized protein LOC108156718 [Drosophila
>gi|1037079209|ref|XP_017143156.1| PREDICTED: RING finger protein unkempt isoform X1 [Drosophil
>gi|1037079223|ref|XP_017143157.1| PREDICTED: RING finger protein unkempt isoform X2 [Drosophil
>gi|1037079237|ref|XP_017143158.1| PREDICTED: RING finger protein unkempt isoform X3 [Drosophil
>gi|1037079443|ref|XP_017143159.1| PREDICTED: RING finger protein unkempt isoform X4 [Drosophil
>gi|1037098106|ref|XP_017144579.1| PREDICTED: protein hedgehog [Drosophila miranda]
>gi|1037034124|ref|XP_017140138.1| PREDICTED: uncharacterized protein LOC108154385 [Drosophila
>gi|1037041474|ref|XP_017140591.1| PREDICTED: histone RNA hairpin-binding protein isoform X1 [l
>gi|1037041488|ref|XP_017140592.1| PREDICTED: histone RNA hairpin-binding protein isoform X2 [l
>gi|1037041430|ref|XP_017140587.1| PREDICTED: eukaryotic translation initiation factor 2D [Dros
>gi|1037041504|ref|XP_017140593.1| PREDICTED: phosphoglycerate mutase 1 [Drosophila miranda]
>gi|1037041460|ref|XP_017140590.1| PREDICTED: uncharacterized protein LOC108154734 [Drosophila
>gi|1037041446|ref|XP_017140589.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1037041290|ref|XP_017140579.1| PREDICTED: aminopeptidase N [Drosophila miranda]
>gi|1037041516|ref|XP_017140594.1| PREDICTED: UDP-N-acetylglucosamine transferase subunit ALG13
>gi|1037041336|ref|XP_017140582.1| PREDICTED: V-type proton ATPase 116 kDa subunit a isoform X3
>gi|1037041320|ref|XP_017140581.1| PREDICTED: V-type proton ATPase 116 kDa subunit a isoform X2
>gi|1037041352|ref|XP_017140583.1| PREDICTED: V-type proton ATPase 116 kDa subunit a isoform X4
>gi|1037041304|ref|XP_017140580.1| PREDICTED: V-type proton ATPase 116 kDa subunit a isoform X1
>gi|1037041398|ref|XP_017140585.1| PREDICTED: uncharacterized protein LOC108154731 isoform X1
>gi|1037041414|ref|XP_017140586.1| PREDICTED: uncharacterized protein LOC108154731 isoform X2
>gi|1037041382|ref|XP_017140584.1| PREDICTED: CTL-like protein 2 [Drosophila miranda]
>gi|1037063561|ref|XP_017142086.1| PREDICTED: yemanuclein-alpha [Drosophila miranda]
>gi|1037063577|ref|XP_017142087.1| PREDICTED: beclin 1-associated autophagy-related key regulat
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>gi|1037063593|ref|XP_017142088.1| PREDICTED: pyruvate dehydrogenase E1 component subunit beta
>gi|1037063609|ref|XP_017142090.1| PREDICTED: uncharacterized protein LOC108155650 isoform X2
>gi|1037063625|ref|XP_017142091.1| PREDICTED: uridine diphosphate glucose pyrophosphatase-like
>gi|1037063639|ref|XP_017142092.1| PREDICTED: uridine diphosphate glucose pyrophosphatase [Dros
>gi|1037024637|ref|XP_017139562.1| PREDICTED: uncharacterized protein LOC108153976 [Drosophila
>gi|1037093911|ref|XP_017144220.1| PREDICTED: protein serrate isoform X1 [Drosophila miranda]
>gi|1037093924|ref|XP_017144221.1| PREDICTED: protein serrate isoform X2 [Drosophila miranda]
>gi|1037104197|ref|XP_017145181.1| PREDICTED: DNA repair protein RAD51 homolog 3 [Drosophila mi
>gi|1037104340|ref|XP_017145200.1| PREDICTED: serine palmitoyltransferase small subunit B [Dros
>gi|1037104346|ref|XP_017145201.1| PREDICTED: degenerin unc-8 [Drosophila miranda]
>gi|1037095694|ref|XP_017144357.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1037095707|ref|XP_017144359.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1037095661|ref|XP_017144355.1| PREDICTED: NADP-dependent malic enzyme isoform X1 [Drosophil
>gi|1037095684|ref|XP_017144356.1| PREDICTED: NADP-dependent malic enzyme isoform X2 [Drosophil
>gi|1037062471|ref|XP_017142010.1| PREDICTED: uncharacterized protein LOC108155599 isoform X1
>gi|1037062486|ref|XP_017142011.1| PREDICTED: uncharacterized protein LOC108155599 isoform X2
>gi|1037062548|ref|XP_017142016.1| PREDICTED: lateral signaling target protein 2 homolog [Dros
>gi|1037062502|ref|XP_017142012.1| PREDICTED: cilia- and flagella-associated protein 58 [Drosop
>gi|1037062518|ref|XP_017142013.1| PREDICTED: cilia- and flagella-associated protein 58 [Drosop
>gi|1037062532|ref|XP_017142014.1| PREDICTED: probable palmitoyltransferase ZDHHC16 [Drosophila
>gi|1037083699|ref|XP_017143484.1| PREDICTED: UPF0396 protein CG6066 [Drosophila miranda]
>gi|1037083685|ref|XP_017143483.1| PREDICTED: large neutral amino acids transporter small subu
>gi|1037104745|ref|XP_017145256.1| PREDICTED: carbonic anhydrase 2 isoform X2 [Drosophila mira
>gi|1037104737|ref|XP_017145255.1| PREDICTED: carbonic anhydrase 2 isoform X1 [Drosophila mira
>gi|1037104753|ref|XP_017145257.1| PREDICTED: uncharacterized protein LOC108157629 [Drosophila
>gi|1037103512|ref|XP_017145091.1| PREDICTED: multiple epidermal growth factor-like domains pr
>gi|1037103520|ref|XP_017145092.1| PREDICTED: uncharacterized protein LOC108157504 [Drosophila
>gi|1037103526|ref|XP_017145093.1| PREDICTED: uncharacterized protein LOC108157505 [Drosophila
>gi|1037032536|ref|XP_017140043.1| PREDICTED: DNA translocase FtsK [Drosophila miranda]
>gi|1037033037|ref|XP_017140074.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 2
>gi|1037033023|ref|XP_017140073.1| PREDICTED: loricrin isoform X3 [Drosophila miranda]
>gi|1037032987|ref|XP_017140071.1| PREDICTED: uncharacterized protein LOC108154323 isoform X1
>gi|1037033005|ref|XP_017140072.1| PREDICTED: uncharacterized protein LOC108154323 isoform X2
>gi|1037098541|ref|XP_017144624.1| PREDICTED: protein spaetzle isoform X1 [Drosophila miranda]
>gi|1037098553|ref|XP_017144626.1| PREDICTED: protein spaetzle isoform X2 [Drosophila miranda]
>gi|1037098565|ref|XP_017144627.1| PREDICTED: protein spaetzle isoform X3 [Drosophila miranda]
>gi|1037098577|ref|XP_017144628.1| PREDICTED: protein spaetzle isoform X4 [Drosophila miranda]
>gi|1037098601|ref|XP_017144630.1| PREDICTED: protein spaetzle isoform X6 [Drosophila miranda]
>gi|1037098613|ref|XP_017144631.1| PREDICTED: protein spaetzle isoform X7 [Drosophila miranda]
>gi|1037098625|ref|XP_017144632.1| PREDICTED: protein spaetzle isoform X8 [Drosophila miranda]
>gi|1037098652|ref|XP_017144634.1| PREDICTED: protein spaetzle isoform X10 [Drosophila miranda]
>gi|1037098664|ref|XP_017144635.1| PREDICTED: protein spaetzle isoform X11 [Drosophila miranda]
>gi|1037098589|ref|XP_017144629.1| PREDICTED: protein spaetzle isoform X5 [Drosophila miranda]
>gi|1037098640|ref|XP_017144633.1| PREDICTED: protein spaetzle isoform X9 [Drosophila miranda]
>gi|1037098676|ref|XP_017144637.1| PREDICTED: protein spaetzle isoform X12 [Drosophila miranda]
>gi|1037098699|ref|XP_017144639.1| PREDICTED: protein spaetzle isoform X14 [Drosophila miranda]
>gi|1037098709|ref|XP_017144640.1| PREDICTED: protein spaetzle isoform X15 [Drosophila miranda]
>gi|1037098687|ref|XP_017144638.1| PREDICTED: protein spaetzle isoform X13 [Drosophila miranda]
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>gi|1037098721|ref|XP_017144641.1| PREDICTED: uncharacterized protein LOC108157204 [Drosophila
>gi|1037033598|ref|XP_017140106.1| PREDICTED: endothelin-converting enzyme 1 [Drosophila mirano
>gi|1037033632|ref|XP_017140109.1| PREDICTED: uncharacterized protein LOC108154353 isoform X2
>gi|1037033616|ref|XP_017140107.1| PREDICTED: uncharacterized protein LOC108154353 isoform X1
>gi|1037095481|ref|XP_017144341.1| PREDICTED: neuropeptide Y receptor isoform X1 [Drosophila m:
>gi|1037095513|ref|XP_017144343.1| PREDICTED: neuropeptide Y receptor isoform X3 [Drosophila m:
>gi|1037095497|ref|XP_017144342.1| PREDICTED: neuropeptide Y receptor isoform X2 [Drosophila m:
>gi|1037095621|ref|XP_017144352.1| PREDICTED: pupal cuticle protein 36a [Drosophila miranda]
>gi|1037095635|ref|XP_017144353.1| PREDICTED: pupal cuticle protein 36a [Drosophila miranda]
>gi|1037095552|ref|XP_017144346.1| PREDICTED: pancreatic triacylglycerol lipase-like [Drosophil
>gi|1037095540|ref|XP_017144345.1| PREDICTED: phospholipase A1-like [Drosophila miranda]
>gi|1037095593|ref|XP_017144349.1| PREDICTED: phospholipase A1-like [Drosophila miranda]
>gi|1037095579|ref|XP_017144348.1| PREDICTED: phospholipase A1-like [Drosophila miranda]
>gi|1037095607|ref|XP_017144351.1| PREDICTED: phospholipase A1-like [Drosophila miranda]
>gi|1037095565|ref|XP_017144347.1| PREDICTED: phospholipase A1 [Drosophila miranda]
>gi|1037095455|ref|XP_017144339.1| PREDICTED: polysialoglycoprotein isoform X1 [Drosophila mira
>gi|1037095467|ref|XP_017144340.1| PREDICTED: uncharacterized protein LOC108157037 isoform X2
>gi|1037095527|ref|XP_017144344.1| PREDICTED: tenascin [Drosophila miranda]
>gi|1037024655|ref|XP_017139563.1| PREDICTED: extensin [Drosophila miranda]
>gi|1037034104|ref|XP_017140137.1| PREDICTED: leucine-rich repeat-containing protein C10orf11 l
>gi|1037074321|ref|XP_017142814.1| PREDICTED: aldehyde dehydrogenase X, mitochondrial [Drosophi
>gi|1037074337|ref|XP_017142815.1| PREDICTED: uncharacterized protein LOC108156053 isoform X1
>gi|1037074349|ref|XP_017142816.1| PREDICTED: uncharacterized protein LOC108156053 isoform X2
>gi|1037074363|ref|XP_017142817.1| PREDICTED: uncharacterized protein LOC108156053 isoform X2
>gi|1037074306|ref|XP_017142813.1| PREDICTED: probable Rho GTPase-activating protein CG5521 isc
>gi|1037074291|ref|XP_017142812.1| PREDICTED: probable Rho GTPase-activating protein CG5521 isc
>gi|1037042518|ref|XP_017140659.1| PREDICTED: putative gustatory receptor 97a [Drosophila mirar
>gi|1037042420|ref|XP_017140652.1| PREDICTED: coatomer subunit gamma [Drosophila miranda]
>gi|1037042436|ref|XP_017140653.1| PREDICTED: protein pygopus [Drosophila miranda]
>gi|1037042406|ref|XP_017140651.1| PREDICTED: uncharacterized protein LOC108154777 [Drosophila
>gi|1037042533|ref|XP_017140660.1| PREDICTED: putative tricarboxylate transport protein, mitoch
>gi|1037042483|ref|XP_017140657.1| PREDICTED: acyl-coenzyme A thioesterase 9, mitochondrial-lil
>gi|1037042500|ref|XP_017140658.1| PREDICTED: acyl-coenzyme A thioesterase 9, mitochondrial-lil
>gi|1037042452|ref|XP_017140654.1| PREDICTED: acyl-coenzyme A thioesterase 9, mitochondrial [D:
>gi|1037042468|ref|XP_017140656.1| PREDICTED: acyl-coenzyme A thioesterase 9, mitochondrial [D:
>gi|1037042543|ref|XP_017140661.1| PREDICTED: 60S ribosomal protein L6 [Drosophila miranda]
>gi|1037062564|ref|XP_017142017.1| PREDICTED: cyclin G [Drosophila miranda]
>gi|1037062580|ref|XP_017142018.1| PREDICTED: cyclin G [Drosophila miranda]
>gi|1037083263|ref|XP_017143448.1| PREDICTED: mothers against decapentaplegic homolog 4 isoforn
>gi|1037083277|ref|XP_017143449.1| PREDICTED: mothers against decapentaplegic homolog 4 isoforn
>gi|1037083305|ref|XP_017143451.1| PREDICTED: N-acetyltransferase 9-like protein [Drosophila m:
>gi|1037083291|ref|XP_017143450.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037050697|ref|XP_017141206.1| PREDICTED: colorectal mutant cancer protein isoform X1 [Dros
>gi|1037050713|ref|XP_017141207.1| PREDICTED: colorectal mutant cancer protein isoform X2 [Dros
>gi|1037050743|ref|XP_017141209.1| PREDICTED: pre-mRNA-splicing factor SPF27 [Drosophila mirano
>gi|1037050727|ref|XP_017141208.1| PREDICTED: 60S ribosomal protein L4 [Drosophila miranda]
>gi|1037050665|ref|XP_017141203.1| PREDICTED: glycerol-3-phosphate acyltransferase 1, mitochono
>gi|1037050757|ref|XP_017141210.1| PREDICTED: RING finger protein 212B [Drosophila miranda]
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>gi|1037050681|ref|XP_017141204.1| PREDICTED: glycerol-3-phosphate acyltransferase 1, mitochond
>gi|1037050651|ref|XP_017141202.1| PREDICTED: probable histone-lysine N-methyltransferase Mes-4
>gi|1037075577|ref|XP_017142901.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1037075593|ref|XP_017142902.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1037075607|ref|XP_017142903.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1037075623|ref|XP_017142904.1| PREDICTED: mitoferrin [Drosophila miranda]
>gi|1037045256|ref|XP_017140838.1| PREDICTED: endoplasmin homolog [Drosophila miranda]
>gi|1037045317|ref|XP_017140843.1| PREDICTED: uncharacterized protein CG4951 isoform X3 [Drosop
>gi|1037045302|ref|XP_017140842.1| PREDICTED: uncharacterized protein CG4951 isoform X2 [Drosop
>gi|1037045288|ref|XP_017140841.1| PREDICTED: uncharacterized protein CG4951 isoform X1 [Drosop
>gi|1037045272|ref|XP_017140840.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1037045426|ref|XP_017140850.1| PREDICTED: uncharacterized protein C6orf203 homolog [Drosoph
>gi|1037045367|ref|XP_017140846.1| PREDICTED: uncharacterized protein LOC108154905 [Drosophila
>gi|1037045379|ref|XP_017140847.1| PREDICTED: uncharacterized protein LOC108154905 [Drosophila
>gi|1037045349|ref|XP_017140845.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1037045240|ref|XP_017140837.1| PREDICTED: 116 kDa U5 small nuclear ribonucleoprotein compon
>gi|1037045410|ref|XP_017140849.1| PREDICTED: uncharacterized protein LOC108154908 [Drosophila
>gi|1037045333|ref|XP_017140844.1| PREDICTED: tubulin beta chain [Drosophila miranda]
>gi|1037045394|ref|XP_017140848.1| PREDICTED: trypsin [Drosophila miranda]
>gi|1037099436|ref|XP_017144702.1| PREDICTED: nucleolysin TIA-1 isoform X1 [Drosophila miranda]
>gi|1037099448|ref|XP_017144703.1| PREDICTED: nucleolysin TIA-1 isoform X2 [Drosophila miranda]
>gi|1037104871|ref|XP_017145276.1| PREDICTED: nucleolysin TIA-1 isoform X1 [Drosophila miranda]
>gi|1037104879|ref|XP_017145277.1| PREDICTED: nucleolysin TIA-1 isoform X1 [Drosophila miranda]
>gi|1037104897|ref|XP_017145278.1| PREDICTED: nucleolysin TIA-1 isoform X2 [Drosophila miranda]
>gi|1037105009|ref|XP_017145295.1| PREDICTED: LOW QUALITY PROTEIN: DBF4-type zinc finger-contai
>gi|1037105017|ref|XP_017145296.1| PREDICTED: DBF4-type zinc finger-containing protein 2 homolo
>gi|1037033398|ref|XP_017140094.1| PREDICTED: uncharacterized protein LOC108154343 [Drosophila
>gi|1037034661|ref|XP_017140174.1| PREDICTED: cytochrome c-type heme lyase-like [Drosophila mir
>gi|1037068950|ref|XP_017142442.1| PREDICTED: LOW QUALITY PROTEIN: la-related protein 1 [Drosop
>gi|1037096135|ref|XP_017144397.1| PREDICTED: LOW QUALITY PROTEIN: glutamine--fructose-6-phosph
>gi|1037077272|ref|XP_017143012.1| PREDICTED: LOW QUALITY PROTEIN: peptidyl-prolyl cis-trans is
>gi|1037077256|ref|XP_017143011.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037096616|ref|XP_017144443.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037037877|ref|XP_017140351.1| PREDICTED: transmembrane protein 26 [Drosophila miranda]
>gi|1037099912|ref|XP_017144745.1| PREDICTED: ribosomal RNA processing protein 36 homolog [Dros
>gi|1037099922|ref|XP_017144746.1| PREDICTED: 39S ribosomal protein L11, mitochondrial [Drosoph
>gi|1037037398|ref|XP_017140325.1| PREDICTED: uncharacterized protein LOC108154532 [Drosophila
>gi|1037032869|ref|XP_017140064.1| PREDICTED: ctenidin-3 [Drosophila miranda]
>gi|1037035577|ref|XP_017140217.1| PREDICTED: uncharacterized protein LOC108154457 isoform X1
>gi|1037035595|ref|XP_017140218.1| PREDICTED: uncharacterized protein LOC108154457 isoform X2
>gi|1037094307|ref|XP_017144250.1| PREDICTED: uncharacterized protein LOC108156971 [Drosophila
>gi|1037094321|ref|XP_017144251.1| PREDICTED: dynein light chain Tctex-type [Drosophila miranda
>gi|1037094335|ref|XP_017144252.1| PREDICTED: dynein light chain Tctex-type [Drosophila miranda
>gi|1037034173|ref|XP_017140141.1| PREDICTED: uncharacterized protein LOC108154389 [Drosophila
>gi|1037103898|ref|XP_017145142.1| PREDICTED: CD209 antigen-like [Drosophila miranda]
>gi|1037081258|ref|XP_017143299.1| PREDICTED: protein no-on-transient A-like isoform X2 [Drosop
>gi|1037081229|ref|XP_017143298.1| PREDICTED: protein no-on-transient A-like isoform X1 [Drosop
>gi|1037057121|ref|XP_017141638.1| PREDICTED: diphthamide biosynthesis protein 2 [Drosophila mi
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>gi|1037057137|ref|XP_017141640.1| PREDICTED: mitochondrial-processing peptidase subunit beta
>gi|1037057228|ref|XP_017141646.1| PREDICTED: cyclin-C [Drosophila miranda]
>gi|1037057183|ref|XP_017141643.1| PREDICTED: GPALPP motifs-containing protein 1 [Drosophila mi
>gi|1037057197|ref|XP_017141644.1| PREDICTED: GPALPP motifs-containing protein 1 [Drosophila mi
>gi|1037057213|ref|XP_017141645.1| PREDICTED: phosphatidate cytidylyltransferase, mitochondrial
>gi|1037057106|ref|XP_017141637.1| PREDICTED: exosome component 10 [Drosophila miranda]
>gi|1037057153|ref|XP_017141641.1| PREDICTED: glycosaminoglycan xylosylkinase homolog [Drosophi
>gi|1037057167|ref|XP_017141642.1| PREDICTED: glycosaminoglycan xylosylkinase homolog [Drosophi
>gi|1037055839|ref|XP_017141552.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037055823|ref|XP_017141551.1| PREDICTED: solute carrier family 22 member 6 [Drosophila mi
>gi|1037055787|ref|XP_017141548.1| PREDICTED: LOW QUALITY PROTEIN: organic cation transporter-1
>gi|1037055805|ref|XP_017141549.1| PREDICTED: LOW QUALITY PROTEIN: organic cation transporter p
>gi|1037045824|ref|XP_017140875.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037045840|ref|XP_017140876.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1037045856|ref|XP_017140877.1| PREDICTED: mucin-5AC isoform X3 [Drosophila miranda]
>gi|1037045872|ref|XP_017140878.1| PREDICTED: regucalcin [Drosophila miranda]
>gi|1037045906|ref|XP_017140880.1| PREDICTED: ubiquitin-conjugating enzyme E2-17 kDa isoform X2
>gi|1037045951|ref|XP_017140882.1| PREDICTED: ubiquitin-conjugating enzyme E2-17 kDa isoform X4
>gi|1037045935|ref|XP_017140881.1| PREDICTED: ubiquitin-conjugating enzyme E2-17 kDa isoform X3
>gi|1037045888|ref|XP_017140879.1| PREDICTED: ubiquitin-conjugating enzyme E2-17 kDa isoform X1
>gi|1037103219|ref|XP_017145059.1| PREDICTED: progestin and adipoQ receptor family member 3 iso
>gi|1037103197|ref|XP_017145057.1| PREDICTED: progestin and adipoQ receptor family member 3 iso
>gi|1037103229|ref|XP_017145060.1| PREDICTED: progestin and adipoQ receptor family member 3 iso
>gi|1037103209|ref|XP_017145058.1| PREDICTED: progestin and adipoQ receptor family member 3 iso
>gi|1037044964|ref|XP_017140820.1| PREDICTED: LOW QUALITY PROTEIN: protein kibra [Drosophila mi
>gi|1037045014|ref|XP_017140823.1| PREDICTED: uncharacterized protein LOC108154891 isoform X1
>gi|1037045029|ref|XP_017140824.1| PREDICTED: uncharacterized protein LOC108154891 isoform X2
>gi|1037045001|ref|XP_017140822.1| PREDICTED: helicase sen1-like [Drosophila miranda]
>gi|1037044979|ref|XP_017140821.1| PREDICTED: TATA box-binding protein-associated factor RNA po
>gi|1037045044|ref|XP_017140825.1| PREDICTED: protein phosphatase PP2A 55 kDa regulatory subuni
>gi|1037024671|ref|XP_017139564.1| PREDICTED: protein phosphatase PP2A 55 kDa regulatory subuni
>gi|1037103080|ref|XP_017145043.1| PREDICTED: LOW QUALITY PROTEIN: protein Skeletor, isoforms I
>gi|1037080692|ref|XP_017143255.1| PREDICTED: protein fem-1 homolog A [Drosophila miranda]
>gi|1037080744|ref|XP_017143260.1| PREDICTED: baculoviral IAP repeat-containing protein 5 [Dros
>gi|1037080717|ref|XP_017143258.1| PREDICTED: single-strand selective monofunctional uracil DNA
>gi|1037080730|ref|XP_017143259.1| PREDICTED: tRNA-specific adenosine deaminase 2 [Drosophila m
>gi|1037080704|ref|XP_017143256.1| PREDICTED: unconventional myosin-XVIIIa-like [Drosophila mi
>gi|1037081846|ref|XP_017143347.1| PREDICTED: leishmanolysin-like peptidase isoform X1 [Drosoph
>gi|1037081860|ref|XP_017143348.1| PREDICTED: leishmanolysin-like peptidase isoform X1 [Drosoph
>gi|1037081872|ref|XP_017143350.1| PREDICTED: leishmanolysin-like peptidase isoform X2 [Drosoph
>gi|1037022524|ref|XP_017139427.1| PREDICTED: uncharacterized protein LOC108153839 [Drosophila
>gi|1037081886|ref|XP_017143351.1| PREDICTED: zinc finger protein weckle [Drosophila miranda]
>gi|1037091319|ref|XP_017144040.1| PREDICTED: bestrophin-2 isoform X4 [Drosophila miranda]
>gi|1037091305|ref|XP_017144039.1| PREDICTED: bestrophin-2 isoform X3 [Drosophila miranda]
>gi|1037091291|ref|XP_017144038.1| PREDICTED: bestrophin-4 isoform X2 [Drosophila miranda]
>gi|1037091263|ref|XP_017144036.1| PREDICTED: bestrophin-4 isoform X1 [Drosophila miranda]
>gi|1037091275|ref|XP_017144037.1| PREDICTED: bestrophin-4 isoform X1 [Drosophila miranda]
>gi|1037102088|ref|XP_017144939.1| PREDICTED: uncharacterized protein LOC108157405 isoform X4
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>gi|1037102078|ref|XP_017144938.1| PREDICTED: uncharacterized protein LOC108157405 isoform X3
>gi|1037102068|ref|XP_017144937.1| PREDICTED: uncharacterized protein LOC108157405 isoform X2
>gi|1037102058|ref|XP_017144936.1| PREDICTED: uncharacterized protein LOC108157405 isoform X1
>gi|1037100963|ref|XP_017144836.1| PREDICTED: LOW QUALITY PROTEIN: synapsin [Drosophila miranda
>gi|1037100953|ref|XP_017144835.1| PREDICTED: tissue inhibitor of metalloproteases [Drosophila
>gi|1037103780|ref|XP_017145127.1| PREDICTED: uncharacterized protein LOC108157535 [Drosophila
>gi|1037103788|ref|XP_017145128.1| PREDICTED: multiple coagulation factor deficiency protein 2
>gi|1037103798|ref|XP_017145129.1| PREDICTED: UPF0691 protein C9orf116 homolog [Drosophila mira
>gi|1037085009|ref|XP_017143587.1| PREDICTED: titin isoform X7 [Drosophila miranda]
>gi|1037085023|ref|XP_017143588.1| PREDICTED: titin isoform X8 [Drosophila miranda]
>gi|1037084929|ref|XP_017143581.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1037084942|ref|XP_017143582.1| PREDICTED: titin isoform X2 [Drosophila miranda]
>gi|1037084956|ref|XP_017143583.1| PREDICTED: titin isoform X3 [Drosophila miranda]
>gi|1037085037|ref|XP_017143589.1| PREDICTED: titin isoform X9 [Drosophila miranda]
>gi|1037084969|ref|XP_017143584.1| PREDICTED: titin isoform X4 [Drosophila miranda]
>gi|1037084983|ref|XP_017143585.1| PREDICTED: titin isoform X5 [Drosophila miranda]
>gi|1037084997|ref|XP_017143586.1| PREDICTED: titin isoform X6 [Drosophila miranda]
>gi|1037085049|ref|XP_017143591.1| PREDICTED: glycine dehydrogenase (decarboxylating), mitochon
>gi|1037066344|ref|XP_017142265.1| PREDICTED: ruvB-like helicase 1 [Drosophila miranda]
>gi|1037066178|ref|XP_017142253.1| PREDICTED: uncharacterized protein LOC108155744 isoform X1
>gi|1037066192|ref|XP_017142254.1| PREDICTED: uncharacterized protein LOC108155744 isoform X1
>gi|1037066204|ref|XP_017142255.1| PREDICTED: uncharacterized protein LOC108155744 isoform X1
>gi|1037066220|ref|XP_017142256.1| PREDICTED: uncharacterized protein LOC108155744 isoform X2
>gi|1037066244|ref|XP_017142257.1| PREDICTED: mucin-5AC isoform X3 [Drosophila miranda]
>gi|1037066258|ref|XP_017142258.1| PREDICTED: mucin-5AC isoform X4 [Drosophila miranda]
>gi|1037066272|ref|XP_017142260.1| PREDICTED: mucin-5AC isoform X5 [Drosophila miranda]
>gi|1037066284|ref|XP_017142261.1| PREDICTED: flocculation protein FLO11 isoform X6 [Drosophila
>gi|1037066300|ref|XP_017142262.1| PREDICTED: flocculation protein FLO11 isoform X7 [Drosophila
>gi|1037066315|ref|XP_017142263.1| PREDICTED: mucin-5AC isoform X8 [Drosophila miranda]
>gi|1037066330|ref|XP_017142264.1| PREDICTED: solute carrier family 23 member 1 [Drosophila mir
>gi|1037043657|ref|XP_017140734.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043641|ref|XP_017140733.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043611|ref|XP_017140731.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043596|ref|XP_017140730.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043524|ref|XP_017140725.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043580|ref|XP_017140729.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043538|ref|XP_017140726.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043564|ref|XP_017140728.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043552|ref|XP_017140727.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043512|ref|XP_017140724.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043627|ref|XP_017140732.1| PREDICTED: baculoviral IAP repeat-containing protein 6 isofo
>gi|1037043708|ref|XP_017140737.1| PREDICTED: leukocyte receptor cluster member 1 [Drosophila m
>gi|1037043723|ref|XP_017140738.1| PREDICTED: leukocyte receptor cluster member 1 [Drosophila m
>gi|1037043673|ref|XP_017140735.1| PREDICTED: protein slender lobes isoform X1 [Drosophila mira
>gi|1037043696|ref|XP_017140736.1| PREDICTED: protein slender lobes isoform X2 [Drosophila mira
>gi|1037102537|ref|XP_017144987.1| PREDICTED: putative leucine-rich repeat-containing protein l
>gi|1037076280|ref|XP_017142947.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037024689|ref|XP_017139565.1| PREDICTED: uncharacterized protein LOC108153981 [Drosophila
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>gi|1037024711|ref|XP_017139566.1| PREDICTED: pancreatic triacylglycerol lipase-like [Drosophil
>gi|1037024729|ref|XP_017139568.1| PREDICTED: solute carrier family 22 member 4-like [Drosophil
>gi|1037089554|ref|XP_017143897.1| PREDICTED: DNA-binding protein RFX2 [Drosophila miranda]
>gi|1037086891|ref|XP_017143725.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037024742|ref|XP_017139569.1| PREDICTED: probable G-protein coupled receptor Mth-like 11
>gi|1037089307|ref|XP_017143881.1| PREDICTED: probable G-protein coupled receptor Mth-like 11
>gi|1037089293|ref|XP_017143880.1| PREDICTED: uncharacterized protein LOC108156732 [Drosophila
>gi|1037094252|ref|XP_017144246.1| PREDICTED: cytoplasmic aconitate hydratase isoform X2 [Droso
>gi|1037094239|ref|XP_017144245.1| PREDICTED: cytoplasmic aconitate hydratase isoform X1 [Droso
>gi|1037024756|ref|XP_017139570.1| PREDICTED: exosome complex component RRP46 [Drosophila miran
>gi|1037097813|ref|XP_017144551.1| PREDICTED: CDK5RAP1-like protein [Drosophila miranda]
>gi|1037102117|ref|XP_017144943.1| PREDICTED: probable cytochrome P450 12e1, mitochondrial isof
>gi|1037102126|ref|XP_017144944.1| PREDICTED: probable cytochrome P450 12e1, mitochondrial isof
>gi|1037046926|ref|XP_017140950.1| PREDICTED: homeobox protein homothorax isoform X2 [Drosophil
>gi|1037046910|ref|XP_017140949.1| PREDICTED: homeobox protein homothorax isoform X1 [Drosophil
>gi|1037046939|ref|XP_017140951.1| PREDICTED: homeobox protein homothorax isoform X3 [Drosophil
>gi|1037046955|ref|XP_017140952.1| PREDICTED: homeobox protein homothorax isoform X4 [Drosophil
>gi|1037094993|ref|XP_017144303.1| PREDICTED: aminoacylase-1B-like [Drosophila miranda]
>gi|1037094979|ref|XP_017144301.1| PREDICTED: aminoacylase-1-like [Drosophila miranda]
>gi|1037095007|ref|XP_017144304.1| PREDICTED: phytanoyl-CoA dioxygenase domain-containing prote
>gi|1037102175|ref|XP_017144949.1| PREDICTED: protein Skeletor, isoforms D/E [Drosophila mirand
>gi|1037078086|ref|XP_017143075.1| PREDICTED: probable 28S rRNA (cytosine-C(5))-methyltransfera
>gi|1037078128|ref|XP_017143078.1| PREDICTED: uncharacterized protein LOC108156241 [Drosophila
>gi|1037078114|ref|XP_017143077.1| PREDICTED: protein nanos [Drosophila miranda]
>gi|1037078100|ref|XP_017143076.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037103154|ref|XP_017145052.1| PREDICTED: uncharacterized protein LOC108157479 [Drosophila
>gi|1037087063|ref|XP_017143738.1| PREDICTED: phosphopantothenate--cysteine ligase [Drosophila
>gi|1037024775|ref|XP_017139571.1| PREDICTED: probable cytochrome P450 12a4, mitochondrial [Dro
>gi|1037087050|ref|XP_017143737.1| PREDICTED: LOW QUALITY PROTEIN: probable cytochrome P450 12a
>gi|1037024793|ref|XP_017139572.1| PREDICTED: LOW QUALITY PROTEIN: protein regulator of cytokin
>gi|1037094474|ref|XP_017144263.1| PREDICTED: probable cytochrome P450 12a4, mitochondrial [Dro
>gi|1037092500|ref|XP_017144124.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1037038317|ref|XP_017140379.1| PREDICTED: uncharacterized protein LOC108154586 [Drosophila
>gi|1037038203|ref|XP_017140371.1| PREDICTED: uncharacterized protein LOC108154580 [Drosophila
>gi|1037083852|ref|XP_017143497.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037105420|ref|XP_017145355.1| PREDICTED: LOW QUALITY PROTEIN: mediator of RNA polymerase
>gi|1037103052|ref|XP_017145040.1| PREDICTED: coiled-coil domain-containing protein 124 [Drosop
>gi|1037105524|ref|XP_017145371.1| PREDICTED: uncharacterized protein LOC108157711 [Drosophila
>gi|1037070441|ref|XP_017142546.1| PREDICTED: integrator complex subunit 11 [Drosophila miranda
>gi|1037070457|ref|XP_017142547.1| PREDICTED: ATP synthase subunit gamma, mitochondrial [Drosop
>gi|1037070470|ref|XP_017142548.1| PREDICTED: ATP synthase subunit gamma, mitochondrial [Drosop
>gi|1037070425|ref|XP_017142545.1| PREDICTED: N-terminal kinase-like protein [Drosophila miran
>gi|1037082692|ref|XP_017143408.1| PREDICTED: transcription factor kayak isoform X1 [Drosophila
>gi|1037082705|ref|XP_017143409.1| PREDICTED: transcription factor kayak isoform X2 [Drosophila
>gi|1037082733|ref|XP_017143411.1| PREDICTED: protein phosphatase PTC7 homolog fig [Drosophila
>gi|1037082719|ref|XP_017143410.1| PREDICTED: transcription factor kayak isoform X3 [Drosophila
>gi|1037089618|ref|XP_017143904.1| PREDICTED: D-beta-hydroxybutyrate dehydrogenase, mitochondri
>gi|1037089604|ref|XP_017143903.1| PREDICTED: uncharacterized protein LOC108156753 [Drosophila
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>gi|1037075168|ref|XP_017142872.1| PREDICTED: fumarylacetoacetate hydrolase domain-containing p
>gi|1037075183|ref|XP_017142873.1| PREDICTED: 39S ribosomal protein L35, mitochondrial [Drosoph
>gi|1037075108|ref|XP_017142867.1| PREDICTED: MICOS complex subunit Mic60 [Drosophila miranda]
>gi|1037075123|ref|XP_017142869.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit beta, mito
>gi|1037075153|ref|XP_017142871.1| PREDICTED: uncharacterized protein LOC108156093 [Drosophila
>gi|1037075138|ref|XP_017142870.1| PREDICTED: uncharacterized protein LOC108156092 [Drosophila
>gi|1037024809|ref|XP_017139573.1| PREDICTED: uncharacterized protein LOC108153991 [Drosophila
>gi|1037101538|ref|XP_017144882.1| PREDICTED: uncharacterized protein LOC108157367 [Drosophila
>gi|1037101516|ref|XP_017144880.1| PREDICTED: ATP synthase subunit d, mitochondrial [Drosophila
>gi|1037101526|ref|XP_017144881.1| PREDICTED: ATP synthase subunit d, mitochondrial [Drosophila
>gi|1037031435|ref|XP_017139980.1| PREDICTED: 39S ribosomal protein L55, mitochondrial [Drosoph
>gi|1037072207|ref|XP_017142665.1| PREDICTED: mushroom body large-type Kenyon cell-specific pro
>gi|1037072223|ref|XP_017142666.1| PREDICTED: mushroom body large-type Kenyon cell-specific pro
>gi|1037072396|ref|XP_017142678.1| PREDICTED: ras-related protein Ral-a-like [Drosophila mirand
>gi|1037072255|ref|XP_017142668.1| PREDICTED: RCC1 domain-containing protein 1 isoform X2 [Dros
>gi|1037072412|ref|XP_017142679.1| PREDICTED: Golgi-associated plant pathogenesis-related prote
>gi|1037072303|ref|XP_017142671.1| PREDICTED: ADP-ribosylation factor-like protein 3 isoform X
>gi|1037072334|ref|XP_017142674.1| PREDICTED: ADP-ribosylation factor-like protein 3 isoform X3
>gi|1037072239|ref|XP_017142667.1| PREDICTED: RCC1 domain-containing protein 1 isoform X1 [Dros
>gi|1037072271|ref|XP_017142669.1| PREDICTED: uncharacterized protein LOC108155986 isoform X3
>gi|1037072289|ref|XP_017142670.1| PREDICTED: uncharacterized protein LOC108155986 isoform X3
>gi|1037072319|ref|XP_017142673.1| PREDICTED: ADP-ribosylation factor-like protein 3 isoform X2
>gi|1037072380|ref|XP_017142677.1| PREDICTED: ADP-ribosylation factor-like protein 3 isoform X3
>gi|1037072350|ref|XP_017142675.1| PREDICTED: ADP-ribosylation factor-like protein 3 isoform X
>gi|1037072366|ref|XP_017142676.1| PREDICTED: ADP-ribosylation factor-like protein 3 isoform X2
>gi|1037072426|ref|XP_017142680.1| PREDICTED: uncharacterized protein LOC108155993 [Drosophila
>gi|1037072442|ref|XP_017142681.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037077747|ref|XP_017143048.1| PREDICTED: delta-1-pyrroline-5-carboxylate dehydrogenase, m
>gi|1037077859|ref|XP_017143057.1| PREDICTED: uncharacterized protein C15orf61 homolog [Drosoph
>gi|1037077817|ref|XP_017143054.1| PREDICTED: lysosomal acid phosphatase [Drosophila miranda]
>gi|1037077831|ref|XP_017143055.1| PREDICTED: uncharacterized protein LOC108156227 [Drosophila
>gi|1037077789|ref|XP_017143052.1| PREDICTED: fibroblast growth factor receptor-like 1 [Drosoph
>gi|1037077845|ref|XP_017143056.1| PREDICTED: protein chibby homolog 1 [Drosophila miranda]
>gi|1037077761|ref|XP_017143049.1| PREDICTED: myosin-2 heavy chain, non muscle isoform X1 [Dros
>gi|1037077775|ref|XP_017143051.1| PREDICTED: myosin-2 heavy chain, non muscle isoform X2 [Dros
>gi|1037077803|ref|XP_017143053.1| PREDICTED: vacuolar-sorting protein SNF8 [Drosophila miranda
>gi|1037092486|ref|XP_017144123.1| PREDICTED: gliolectin [Drosophila miranda]
>gi|1037037119|ref|XP_017140308.1| PREDICTED: putative gustatory receptor 93c [Drosophila miran
>gi|1037037133|ref|XP_017140309.1| PREDICTED: putative gustatory receptor 93c [Drosophila miran
>gi|1037039414|ref|XP_017140446.1| PREDICTED: LOW QUALITY PROTEIN: putative gustatory receptor
>gi|1037103599|ref|XP_017145102.1| PREDICTED: gustatory receptor for bitter taste 93a [Drosophi
>gi|1037073283|ref|XP_017142742.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073297|ref|XP_017142743.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073313|ref|XP_017142744.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073363|ref|XP_017142747.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073379|ref|XP_017142748.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073329|ref|XP_017142745.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073347|ref|XP_017142746.1| PREDICTED: peripheral plasma membrane protein CASK isoform
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>gi|1037073392|ref|XP_017142749.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073433|ref|XP_017142752.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073449|ref|XP_017142753.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073406|ref|XP_017142750.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073420|ref|XP_017142751.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037073465|ref|XP_017142755.1| PREDICTED: peripheral plasma membrane protein CASK isoform
>gi|1037061009|ref|XP_017141913.1| PREDICTED: torso-like protein [Drosophila miranda]
>gi|1037061024|ref|XP_017141914.1| PREDICTED: torso-like protein [Drosophila miranda]
>gi|1037060994|ref|XP_017141912.1| PREDICTED: uridine phosphorylase 1-like [Drosophila miranda]
>gi|1037061072|ref|XP_017141917.1| PREDICTED: DNA-directed RNA polymerase I subunit RPA12 [Dros
>gi|1037061088|ref|XP_017141919.1| PREDICTED: cyclin-dependent kinase 20-like [Drosophila mira
>gi|1037060801|ref|XP_017141898.1| PREDICTED: gamma-aminobutyric acid type B receptor subunit
>gi|1037060817|ref|XP_017141899.1| PREDICTED: gamma-aminobutyric acid type B receptor subunit
>gi|1037060833|ref|XP_017141900.1| PREDICTED: gamma-aminobutyric acid type B receptor subunit
>gi|1037060847|ref|XP_017141901.1| PREDICTED: gamma-aminobutyric acid type B receptor subunit
>gi|1037061106|ref|XP_017141920.1| PREDICTED: cyclin-dependent kinase 20-like [Drosophila mira
>gi|1037060863|ref|XP_017141903.1| PREDICTED: gamma-aminobutyric acid type B receptor subunit
>gi|1037060881|ref|XP_017141904.1| PREDICTED: gamma-aminobutyric acid type B receptor subunit
>gi|1037061039|ref|XP_017141915.1| PREDICTED: bursicon [Drosophila miranda]
>gi|1037060978|ref|XP_017141911.1| PREDICTED: aminopeptidase N [Drosophila miranda]
>gi|1037060946|ref|XP_017141908.1| PREDICTED: membrane alanyl aminopeptidase [Drosophila miran
>gi|1037060962|ref|XP_017141909.1| PREDICTED: membrane alanyl aminopeptidase [Drosophila miran
>gi|1037060895|ref|XP_017141905.1| PREDICTED: aminopeptidase N [Drosophila miranda]
>gi|1037060928|ref|XP_017141907.1| PREDICTED: aminopeptidase N [Drosophila miranda]
>gi|1037060914|ref|XP_017141906.1| PREDICTED: aminopeptidase N [Drosophila miranda]
>gi|1037061056|ref|XP_017141916.1| PREDICTED: uncharacterized protein LOC108155548 [Drosophila
>gi|1037033950|ref|XP_017140127.1| PREDICTED: uncharacterized protein LOC108154375 [Drosophila
>gi|1037105712|ref|XP_017145396.1| PREDICTED: carbonic anhydrase 1-like [Drosophila miranda]
>gi|1037105696|ref|XP_017145394.1| PREDICTED: uncharacterized protein LOC108157731 isoform X1
>gi|1037105704|ref|XP_017145395.1| PREDICTED: muscarinic acetylcholine receptor M4 isoform X2
>gi|1037105492|ref|XP_017145366.1| PREDICTED: fatty-acid amide hydrolase 2-B [Drosophila miran
>gi|1037097553|ref|XP_017144526.1| PREDICTED: LOW QUALITY PROTEIN: fatty-acid amide hydrolase
>gi|1037104507|ref|XP_017145223.1| PREDICTED: importin-4-like [Drosophila miranda]
>gi|1037104517|ref|XP_017145224.1| PREDICTED: LOW QUALITY PROTEIN: importin-4-like [Drosophila
>gi|1037083573|ref|XP_017143475.1| PREDICTED: actin, indirect flight muscle [Drosophila mirand
>gi|1037083531|ref|XP_017143471.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1037083545|ref|XP_017143473.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1037083559|ref|XP_017143474.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1037098411|ref|XP_017144612.1| PREDICTED: B9 domain-containing protein 1 [Drosophila miran
>gi|1037098399|ref|XP_017144611.1| PREDICTED: opsin Rh6 [Drosophila miranda]
>gi|1037037765|ref|XP_017140346.1| PREDICTED: uncharacterized protein LOC108154554 [Drosophila
>gi|1037083405|ref|XP_017143461.1| PREDICTED: activating signal cointegrator 1 complex subunit
>gi|1037083419|ref|XP_017143463.1| PREDICTED: probable cytochrome P450 6d4 [Drosophila miranda]
>gi|1037077417|ref|XP_017143023.1| PREDICTED: cardioactive peptide [Drosophila miranda]
>gi|1037077403|ref|XP_017143022.1| PREDICTED: uncharacterized protein LOC108156199 [Drosophila
>gi|1037077433|ref|XP_017143024.1| PREDICTED: uncharacterized protein LOC108156202 [Drosophila
>gi|1037077390|ref|XP_017143021.1| PREDICTED: LOW QUALITY PROTEIN: protein winged eye [Drosoph
>gi|1037082481|ref|XP_017143394.1| PREDICTED: cytoplasmic aconitate hydratase [Drosophila mira
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>gi|1037082527|ref|XP_017143395.1| PREDICTED: E3 ubiquitin-protein ligase Rnf220 [Drosophila mi
>gi|1037088625|ref|XP_017143827.1| PREDICTED: transmembrane protein 47 isoform X1 [Drosophila r
>gi|1037088636|ref|XP_017143829.1| PREDICTED: transmembrane protein 47 isoform X2 [Drosophila r
>gi|1037088649|ref|XP_017143830.1| PREDICTED: transmembrane protein 47 isoform X2 [Drosophila r
>gi|1037088611|ref|XP_017143826.1| PREDICTED: mitogen-activated protein kinase kinase kinase 7
>gi|1037075561|ref|XP_017142899.1| PREDICTED: endoribonuclease Dcr-1 [Drosophila miranda]
>gi|1037092934|ref|XP_017144152.1| PREDICTED: uncharacterized protein LOC108156915 [Drosophila
>gi|1037092920|ref|XP_017144151.1| PREDICTED: protein vreteno [Drosophila miranda]
>gi|1037087773|ref|XP_017143761.1| PREDICTED: LOW QUALITY PROTEIN: chromobox protein homolog 3
>gi|1037087759|ref|XP_017143760.1| PREDICTED: mitochondrial GTPase 1 [Drosophila miranda]
>gi|1037087731|ref|XP_017143758.1| PREDICTED: O-glucosyltransferase rumi [Drosophila miranda]
>gi|1037087745|ref|XP_017143759.1| PREDICTED: O-glucosyltransferase rumi [Drosophila miranda]
>gi|1037058479|ref|XP_017141735.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1037058495|ref|XP_017141736.1| PREDICTED: membrane metallo-endopeptidase-like 1 [Drosophila
>gi|1037058511|ref|XP_017141737.1| PREDICTED: phosphate-regulating neutral endopeptidase [Droso
>gi|1037058560|ref|XP_017141741.1| PREDICTED: uncharacterized protein LOC108155439 [Drosophila
>gi|1037058544|ref|XP_017141739.1| PREDICTED: flocculation protein FLO11-like [Drosophila miran
>gi|1037058526|ref|XP_017141738.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 12A [Drosop
>gi|1037058463|ref|XP_017141734.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037035305|ref|XP_017140206.1| PREDICTED: uncharacterized protein LOC108154446 [Drosophila
>gi|1037106612|ref|XP_017145520.1| PREDICTED: DNA N6-methyl adenine demethylase isoform X1 [Dro
>gi|1037106620|ref|XP_017145522.1| PREDICTED: DNA N6-methyl adenine demethylase isoform X2 [Dro
>gi|1037024827|ref|XP_017139574.1| PREDICTED: uncharacterized protein LOC108153992 [Drosophila
>gi|1037104525|ref|XP_017145226.1| PREDICTED: lon protease homolog, mitochondrial-like [Drosopl
>gi|1037024844|ref|XP_017139575.1| PREDICTED: ras-related protein Ral-a-like [Drosophila miranc
>gi|1037024860|ref|XP_017139577.1| PREDICTED: uncharacterized protein LOC108153994 [Drosophila
>gi|1037024875|ref|XP_017139578.1| PREDICTED: calcium uptake protein 1 homolog, mitochondrial
>gi|1037097342|ref|XP_017144506.1| PREDICTED: neuronal growth regulator 1 [Drosophila miranda]
>gi|1037097352|ref|XP_017144507.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037104128|ref|XP_017145171.1| PREDICTED: sodium/hydrogen exchanger 9B1 isoform X1 [Drosopl
>gi|1037104136|ref|XP_017145172.1| PREDICTED: sodium/hydrogen exchanger 9B1 isoform X2 [Drosopl
>gi|1037104155|ref|XP_017145174.1| PREDICTED: odorant receptor 94a [Drosophila miranda]
>gi|1037104146|ref|XP_017145173.1| PREDICTED: odorant receptor 94b [Drosophila miranda]
>gi|1037029950|ref|XP_017139886.1| PREDICTED: uncharacterized protein LOC108154190 [Drosophila
>gi|1037029968|ref|XP_017139888.1| PREDICTED: uncharacterized protein LOC108154190 [Drosophila
>gi|1037029985|ref|XP_017139889.1| PREDICTED: uncharacterized protein LOC108154190 [Drosophila
>gi|1037030003|ref|XP_017139890.1| PREDICTED: uncharacterized protein LOC108154190 [Drosophila
>gi|1037030019|ref|XP_017139891.1| PREDICTED: uncharacterized protein LOC108154190 [Drosophila
>gi|1037030449|ref|XP_017139918.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037029325|ref|XP_017139849.1| PREDICTED: uncharacterized protein LOC108154185 [Drosophila
>gi|1037030220|ref|XP_017139903.1| PREDICTED: estradiol 17-beta-dehydrogenase 11 [Drosophila mi
>gi|1037030429|ref|XP_017139916.1| PREDICTED: uncharacterized protein LOC108154206 [Drosophila
>gi|1037030341|ref|XP_017139910.1| PREDICTED: uncharacterized protein LOC108154202 [Drosophila
>gi|1037030202|ref|XP_017139902.1| PREDICTED: estradiol 17-beta-dehydrogenase 11 [Drosophila mi
>gi|1037030035|ref|XP_017139892.1| PREDICTED: uncharacterized protein LOC108154191 isoform X1
>gi|1037030053|ref|XP_017139893.1| PREDICTED: uncharacterized protein LOC108154191 isoform X2
>gi|1037030071|ref|XP_017139894.1| PREDICTED: cellular tumor antigen p53 isoform X3 [Drosophila
>gi|1037030089|ref|XP_017139895.1| PREDICTED: cellular tumor antigen p53 isoform X4 [Drosophila
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>gi|1037030144|ref|XP_017139899.1| PREDICTED: putative gustatory receptor 94a [Drosophila mira
>gi|1037030166|ref|XP_017139900.1| PREDICTED: cystinosin homolog [Drosophila miranda]
>gi|1037030105|ref|XP_017139896.1| PREDICTED: scaffold protein salvador [Drosophila miranda]
>gi|1037030307|ref|XP_017139908.1| PREDICTED: G-box-binding factor isoform X2 [Drosophila mira
>gi|1037030289|ref|XP_017139907.1| PREDICTED: nuclear transcription factor Y subunit beta isof
>gi|1037030325|ref|XP_017139909.1| PREDICTED: nuclear transcription factor Y subunit beta isof
>gi|1037030359|ref|XP_017139911.1| PREDICTED: uncharacterized protein LOC108154203 isoform X1
>gi|1037030375|ref|XP_017139912.1| PREDICTED: uncharacterized protein LOC108154203 isoform X2
>gi|1037029313|ref|XP_017139848.1| PREDICTED: Fanconi anemia group D2 protein homolog [Drosoph
>gi|1037029708|ref|XP_017139873.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029724|ref|XP_017139874.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029690|ref|XP_017139872.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029742|ref|XP_017139875.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029672|ref|XP_017139871.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029656|ref|XP_017139870.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029413|ref|XP_017139854.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029395|ref|XP_017139853.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029642|ref|XP_017139869.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029624|ref|XP_017139868.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029588|ref|XP_017139865.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029553|ref|XP_017139863.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029483|ref|XP_017139859.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029568|ref|XP_017139864.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029433|ref|XP_017139855.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029778|ref|XP_017139877.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029465|ref|XP_017139858.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029377|ref|XP_017139852.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029760|ref|XP_017139876.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029450|ref|XP_017139857.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037029535|ref|XP_017139862.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029359|ref|XP_017139851.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029499|ref|XP_017139860.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029517|ref|XP_017139861.1| PREDICTED: heterogeneous nuclear ribonucleoprotein Q isofor
>gi|1037029606|ref|XP_017139866.1| PREDICTED: heterogeneous nuclear ribonucleoprotein R isofor
>gi|1037030127|ref|XP_017139898.1| PREDICTED: putative mitogen-activated protein kinase kinase
>gi|1037029343|ref|XP_017139850.1| PREDICTED: uncharacterized protein LOC108154186 [Drosophila
>gi|1037029796|ref|XP_017139879.1| PREDICTED: uncharacterized protein LOC108154189 isoform X1
>gi|1037029813|ref|XP_017139880.1| PREDICTED: uncharacterized protein LOC108154189 isoform X1
>gi|1037029932|ref|XP_017139885.1| PREDICTED: uncharacterized protein LOC108154189 isoform X6
>gi|1037029896|ref|XP_017139883.1| PREDICTED: uncharacterized protein LOC108154189 isoform X4
>gi|1037029831|ref|XP_017139881.1| PREDICTED: uncharacterized protein LOC108154189 isoform X2
>gi|1037029914|ref|XP_017139884.1| PREDICTED: uncharacterized protein LOC108154189 isoform X5
>gi|1037029849|ref|XP_017139882.1| PREDICTED: uncharacterized protein LOC108154189 isoform X3
>gi|1037030271|ref|XP_017139906.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1037030411|ref|XP_017139915.1| PREDICTED: transmembrane protease serine 9-like [Drosophila
>gi|1037030184|ref|XP_017139901.1| PREDICTED: uncharacterized protein LOC108154196 [Drosophila
>gi|1037030253|ref|XP_017139905.1| PREDICTED: cyclin-dependent kinase 2 [Drosophila miranda]
>gi|1037030393|ref|XP_017139914.1| PREDICTED: low-density lipoprotein receptor-related protein
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>gi|1037095714|ref|XP_017144360.1| PREDICTED: uncharacterized protein LOC108157052 [Drosophila
>gi|1037095726|ref|XP_017144361.1| PREDICTED: uncharacterized protein LOC108157052 [Drosophila
>gi|1037095738|ref|XP_017144362.1| PREDICTED: uncharacterized protein LOC108157052 [Drosophila
>gi|1037105650|ref|XP_017145387.1| PREDICTED: venom peptide isomerase heavy chain-like isoform
>gi|1037105656|ref|XP_017145388.1| PREDICTED: uncharacterized protein LOC108157727 isoform X2
>gi|1037024890|ref|XP_017139579.1| PREDICTED: vitellin-degrading protease-like [Drosophila mira
>gi|1037104729|ref|XP_017145254.1| PREDICTED: uncharacterized protein LOC108157627 [Drosophila
>gi|1037033917|ref|XP_017140126.1| PREDICTED: serine protease easter-like [Drosophila miranda]
>gi|1037103866|ref|XP_017145138.1| PREDICTED: octopamine receptor Oamb isoform X1 [Drosophila r
>gi|1037103874|ref|XP_017145139.1| PREDICTED: octopamine receptor Oamb isoform X2 [Drosophila r
>gi|1037032466|ref|XP_017140038.1| PREDICTED: glycine-rich cell wall structural protein 1.0 [Dr
>gi|1037105532|ref|XP_017145372.1| PREDICTED: histidine-rich protein PFHRP-II [Drosophila mirar
>gi|1037068227|ref|XP_017142391.1| PREDICTED: endothelin-converting enzyme 1 isoform X2 [Drosop
>gi|1037068211|ref|XP_017142390.1| PREDICTED: endothelin-converting enzyme 1 isoform X1 [Drosop
>gi|1037068243|ref|XP_017142393.1| PREDICTED: glutamate receptor ionotropic, kainate 2 [Drosopl
>gi|1037068256|ref|XP_017142394.1| PREDICTED: glutamate receptor ionotropic, kainate 2 [Drosopl
>gi|1037068181|ref|XP_017142388.1| PREDICTED: leucine-rich repeat serine/threonine-protein kina
>gi|1037068163|ref|XP_017142387.1| PREDICTED: leucine-rich repeat serine/threonine-protein kina
>gi|1037068131|ref|XP_017142385.1| PREDICTED: leucine-rich repeat serine/threonine-protein kina
>gi|1037068115|ref|XP_017142384.1| PREDICTED: leucine-rich repeat serine/threonine-protein kina
>gi|1037068197|ref|XP_017142389.1| PREDICTED: leucine-rich repeat serine/threonine-protein kina
>gi|1037068147|ref|XP_017142386.1| PREDICTED: leucine-rich repeat serine/threonine-protein kina
>gi|1037062218|ref|XP_017141993.1| PREDICTED: phosphatidylinositol 4,5-bisphosphate 3-kinase ca
>gi|1037062236|ref|XP_017141994.1| PREDICTED: phosphatidylinositol 4,5-bisphosphate 3-kinase ca
>gi|1037062172|ref|XP_017141990.1| PREDICTED: protein hairless isoform X1 [Drosophila miranda]
>gi|1037062186|ref|XP_017141991.1| PREDICTED: protein hairless isoform X2 [Drosophila miranda]
>gi|1037062204|ref|XP_017141992.1| PREDICTED: protein hairless isoform X2 [Drosophila miranda]
>gi|1037101027|ref|XP_017144842.1| PREDICTED: ras guanine nucleotide exchange factor B isoform
>gi|1037101003|ref|XP_017144840.1| PREDICTED: ras guanine nucleotide exchange factor B isoform
>gi|1037101017|ref|XP_017144841.1| PREDICTED: ras guanine nucleotide exchange factor B isoform
>gi|1037080166|ref|XP_017143216.1| PREDICTED: lipase 3-like [Drosophila miranda]
>gi|1037080152|ref|XP_017143215.1| PREDICTED: lipase 3-like [Drosophila miranda]
>gi|1037080180|ref|XP_017143217.1| PREDICTED: lipase 3-like isoform X1 [Drosophila miranda]
>gi|1037080194|ref|XP_017143218.1| PREDICTED: lipase 3-like isoform X2 [Drosophila miranda]
>gi|1037080138|ref|XP_017143214.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1037080124|ref|XP_017143213.1| PREDICTED: translation initiation factor IF-2 [Drosophila mi
>gi|1037104818|ref|XP_017145268.1| PREDICTED: xaa-Pro aminopeptidase 2 [Drosophila miranda]
>gi|1037037226|ref|XP_017140313.1| PREDICTED: uncharacterized protein LOC108154525 isoform X1
>gi|1037037262|ref|XP_017140316.1| PREDICTED: uncharacterized protein LOC108154525 isoform X3
>gi|1037037244|ref|XP_017140314.1| PREDICTED: uncharacterized protein LOC108154525 isoform X2
>gi|1037037276|ref|XP_017140317.1| PREDICTED: uncharacterized protein LOC108154525 isoform X4
>gi|1037090141|ref|XP_017143949.1| PREDICTED: glycine N-methyltransferase [Drosophila miranda]
>gi|1037090155|ref|XP_017143950.1| PREDICTED: uncharacterized protein LOC108156786 [Drosophila
>gi|1037090127|ref|XP_017143947.1| PREDICTED: nuclear pore complex protein Nup88 [Drosophila mi
>gi|1037024908|ref|XP_017139580.1| PREDICTED: uncharacterized protein LOC108153998 [Drosophila
>gi|1037024936|ref|XP_017139581.1| PREDICTED: uncharacterized protein LOC108153999 [Drosophila
>gi|1037032245|ref|XP_017140024.1| PREDICTED: homeobox protein bagpipe [Drosophila miranda]
>gi|1037024953|ref|XP_017139582.1| PREDICTED: uncharacterized protein LOC108154000, partial [Dr
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>gi|1037104533|ref|XP_017145227.1| PREDICTED: muscle-specific homeobox protein tinman [Drosophi
>gi|1037036290|ref|XP_017140260.1| PREDICTED: modifier of mdg4-like isoform X17 [Drosophila mir
>gi|1037036420|ref|XP_017140267.1| PREDICTED: modifier of mdg4-like isoform X24 [Drosophila mir
>gi|1037036404|ref|XP_017140266.1| PREDICTED: modifier of mdg4-like isoform X23 [Drosophila mir
>gi|1037036387|ref|XP_017140265.1| PREDICTED: modifier of mdg4-like isoform X22 [Drosophila mir
>gi|1037036167|ref|XP_017140251.1| PREDICTED: modifier of mdg4-like isoform X10 [Drosophila mir
>gi|1037036200|ref|XP_017140253.1| PREDICTED: modifier of mdg4-like isoform X12 [Drosophila mir
>gi|1037036045|ref|XP_017140244.1| PREDICTED: modifier of mdg4-like isoform X4 [Drosophila mira
>gi|1037036253|ref|XP_017140256.1| PREDICTED: modifier of mdg4-like isoform X15 [Drosophila mir
>gi|1037036317|ref|XP_017140261.1| PREDICTED: modifier of mdg4-like isoform X18 [Drosophila mir
>gi|1037036333|ref|XP_017140262.1| PREDICTED: modifier of mdg4-like isoform X19 [Drosophila mir
>gi|1037036027|ref|XP_017140243.1| PREDICTED: modifier of mdg4-like isoform X3 [Drosophila mira
>gi|1037036131|ref|XP_017140249.1| PREDICTED: modifier of mdg4-like isoform X8 [Drosophila mira
>gi|1037036149|ref|XP_017140250.1| PREDICTED: modifier of mdg4-like isoform X9 [Drosophila mira
>gi|1037036369|ref|XP_017140264.1| PREDICTED: modifier of mdg4-like isoform X21 [Drosophila mir
>gi|1037036216|ref|XP_017140254.1| PREDICTED: modifier of mdg4-like isoform X13 [Drosophila mir
>gi|1037036081|ref|XP_017140246.1| PREDICTED: modifier of mdg4-like isoform X5 [Drosophila mira
>gi|1037036271|ref|XP_017140259.1| PREDICTED: modifier of mdg4-like isoform X16 [Drosophila mir
>gi|1037035993|ref|XP_017140240.1| PREDICTED: modifier of mdg4-like isoform X1 [Drosophila mira
>gi|1037036063|ref|XP_017140245.1| PREDICTED: modifier of mdg4-like isoform X1 [Drosophila mira
>gi|1037036233|ref|XP_017140255.1| PREDICTED: modifier of mdg4-like isoform X14 [Drosophila mir
>gi|1037036113|ref|XP_017140248.1| PREDICTED: modifier of mdg4-like isoform X7 [Drosophila mira
>gi|1037036184|ref|XP_017140252.1| PREDICTED: modifier of mdg4-like isoform X11 [Drosophila mir
>gi|1037036437|ref|XP_017140268.1| PREDICTED: modifier of mdg4-like isoform X25 [Drosophila mir
>gi|1037036351|ref|XP_017140263.1| PREDICTED: modifier of mdg4-like isoform X20 [Drosophila mir
>gi|1037036099|ref|XP_017140247.1| PREDICTED: modifier of mdg4-like isoform X6 [Drosophila mira
>gi|1037036009|ref|XP_017140242.1| PREDICTED: modifier of mdg4-like isoform X2 [Drosophila mira
>gi|1037022540|ref|XP_017139428.1| PREDICTED: TPR-containing protein DDB_G0280363, partial [Dro
>gi|1037036538|ref|XP_017140274.1| PREDICTED: uncharacterized protein LOC108154483 [Drosophila
>gi|1037036455|ref|XP_017140269.1| PREDICTED: uncharacterized protein LOC108154481 isoform X1
>gi|1037036473|ref|XP_017140270.1| PREDICTED: uncharacterized protein LOC108154481 isoform X2
>gi|1037036491|ref|XP_017140271.1| PREDICTED: uncharacterized protein LOC108154481 isoform X3
>gi|1037036553|ref|XP_017140275.1| PREDICTED: uncharacterized protein LOC108154484 [Drosophila
>gi|1037036509|ref|XP_017140272.1| PREDICTED: uncharacterized protein LOC108154482 [Drosophila
>gi|1037024971|ref|XP_017139584.1| PREDICTED: uncharacterized protein LOC108154001 [Drosophila
>gi|1037052853|ref|XP_017141354.1| PREDICTED: neuropeptide SIFamide receptor isoform X2 [Drosop
>gi|1037052868|ref|XP_017141355.1| PREDICTED: neuropeptide SIFamide receptor isoform X3 [Drosop
>gi|1037052840|ref|XP_017141353.1| PREDICTED: neuropeptide SIFamide receptor isoform X1 [Drosop
>gi|1037052883|ref|XP_017141356.1| PREDICTED: uncharacterized protein LOC108155193 [Drosophila
>gi|1037097915|ref|XP_017144561.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1037075051|ref|XP_017142863.1| PREDICTED: 3'-5' ssDNA/RNA exonuclease TatD [Drosophila mira
>gi|1037075066|ref|XP_017142864.1| PREDICTED: isopentenyl-diphosphate Delta-isomerase 1 [Drosop
>gi|1037075092|ref|XP_017142866.1| PREDICTED: AP-2 complex subunit sigma [Drosophila miranda]
>gi|1037075080|ref|XP_017142865.1| PREDICTED: ras-related protein Rab-1A [Drosophila miranda]
>gi|1037075019|ref|XP_017142861.1| PREDICTED: growth hormone secretagogue receptor type 1 isofo
>gi|1037075035|ref|XP_017142862.1| PREDICTED: growth hormone secretagogue receptor type 1 isofo
>gi|1037105122|ref|XP_017145312.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1037080567|ref|XP_017143247.1| PREDICTED: mycosubtilin synthase subunit C [Drosophila mirar
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>gi|1037080553|ref|XP_017143245.1| PREDICTED: negative elongation factor A [Drosophila miranda]
>gi|1037031366|ref|XP_017139975.1| PREDICTED: protein FAM173B [Drosophila miranda]
>gi|1037031009|ref|XP_017139953.1| PREDICTED: protein O-GlcNAcase [Drosophila miranda]
>gi|1037031328|ref|XP_017139972.1| PREDICTED: uncharacterized protein LOC108154248 [Drosophila
>gi|1037031348|ref|XP_017139974.1| PREDICTED: uncharacterized protein LOC108154248 [Drosophila
>gi|1037031099|ref|XP_017139958.1| PREDICTED: actin-binding LIM protein 3 isoform X5 [Drosophil
>gi|1037031065|ref|XP_017139956.1| PREDICTED: actin-binding LIM protein 3 isoform X3 [Drosophil
>gi|1037031083|ref|XP_017139957.1| PREDICTED: actin-binding LIM protein 3 isoform X4 [Drosophil
>gi|1037031027|ref|XP_017139954.1| PREDICTED: actin-binding LIM protein 3 isoform X1 [Drosophil
>gi|1037031045|ref|XP_017139955.1| PREDICTED: actin-binding LIM protein 3 isoform X2 [Drosophil
>gi|1037031115|ref|XP_017139959.1| PREDICTED: actin-binding LIM protein 3 isoform X6 [Drosophil
>gi|1037031169|ref|XP_017139962.1| PREDICTED: ubiquitin thioesterase trabid [Drosophila miranda
>gi|1037031185|ref|XP_017139963.1| PREDICTED: ubiquitin thioesterase trabid [Drosophila miranda
>gi|1037031151|ref|XP_017139961.1| PREDICTED: protein dalmatian isoform X2 [Drosophila miranda]
>gi|1037031133|ref|XP_017139960.1| PREDICTED: protein dalmatian isoform X1 [Drosophila miranda]
>gi|1037031221|ref|XP_017139966.1| PREDICTED: LOW QUALITY PROTEIN: major heat shock 70 kDa prot
>gi|1037024989|ref|XP_017139585.1| PREDICTED: major heat shock 70 kDa protein Bbb-like [Drosoph
>gi|1037031203|ref|XP_017139964.1| PREDICTED: protein VAC14 homolog [Drosophila miranda]
>gi|1037031239|ref|XP_017139967.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC3 [Dro
>gi|1037031293|ref|XP_017139970.1| PREDICTED: F-box only protein 9 [Drosophila miranda]
>gi|1037031255|ref|XP_017139968.1| PREDICTED: programmed cell death protein 2-like isoform X1
>gi|1037031273|ref|XP_017139969.1| PREDICTED: programmed cell death protein 2-like isoform X2
>gi|1037031382|ref|XP_017139976.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosoph
>gi|1037031310|ref|XP_017139971.1| PREDICTED: UNC93-like protein MFSD11 [Drosophila miranda]
>gi|1037062906|ref|XP_017142042.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1037062916|ref|XP_017142043.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1037062892|ref|XP_017142041.1| PREDICTED: uncharacterized protein LOC108155622 [Drosophila
>gi|1037062930|ref|XP_017142044.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1037062944|ref|XP_017142045.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1037062962|ref|XP_017142046.1| PREDICTED: probable enoyl-CoA hydratase echA8 [Drosophila m
>gi|1037062977|ref|XP_017142047.1| PREDICTED: SUMO-activating enzyme subunit 1 [Drosophila mira
>gi|1037062994|ref|XP_017142048.1| PREDICTED: proteasome subunit alpha type-2 [Drosophila miran
>gi|1037025005|ref|XP_017139586.1| PREDICTED: interleukin enhancer-binding factor 2 homolog, pa
>gi|1037092906|ref|XP_017144150.1| PREDICTED: sialic acid synthase [Drosophila miranda]
>gi|1037091515|ref|XP_017144055.1| PREDICTED: serine/threonine-protein phosphatase alpha-2 iso
>gi|1037091525|ref|XP_017144057.1| PREDICTED: serine/threonine-protein phosphatase alpha-2 iso
>gi|1037091539|ref|XP_017144058.1| PREDICTED: serine/threonine-protein phosphatase alpha-2 iso
>gi|1037091555|ref|XP_017144059.1| PREDICTED: serine/threonine-protein phosphatase alpha-2 iso
>gi|1037091569|ref|XP_017144060.1| PREDICTED: C-Myc-binding protein homolog [Drosophila miranda
>gi|1037043450|ref|XP_017140721.1| PREDICTED: beta-sarcoglycan [Drosophila miranda]
>gi|1037043296|ref|XP_017140710.1| PREDICTED: uncharacterized protein LOC108154820 isoform X2
>gi|1037043280|ref|XP_017140709.1| PREDICTED: uncharacterized protein LOC108154820 isoform X1
>gi|1037043384|ref|XP_017140716.1| PREDICTED: xaa-Pro dipeptidase [Drosophila miranda]
>gi|1037043432|ref|XP_017140720.1| PREDICTED: palmitoyltransferase ZDHHC6 [Drosophila miranda]
>gi|1037043400|ref|XP_017140718.1| PREDICTED: saccharopine dehydrogenase-like oxidoreductase [
>gi|1037043416|ref|XP_017140719.1| PREDICTED: saccharopine dehydrogenase-like oxidoreductase [
>gi|1037043494|ref|XP_017140723.1| PREDICTED: suppressor of lurcher protein 1-like [Drosophila
>gi|1037043314|ref|XP_017140711.1| PREDICTED: DNA ligase 3 isoform X1 [Drosophila miranda]
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>gi|1037043326|ref|XP_017140712.1| PREDICTED: DNA ligase 3 isoform X2 [Drosophila miranda]
>gi|1037043340|ref|XP_017140713.1| PREDICTED: probable cytochrome P450 9f2 [Drosophila miranda]
>gi|1037043354|ref|XP_017140714.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037043369|ref|XP_017140715.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037025023|ref|XP_017139587.1| PREDICTED: uncharacterized protein LOC108154004 [Drosophila
>gi|1037043480|ref|XP_017140722.1| PREDICTED: calmodulin [Drosophila miranda]
>gi|1037099554|ref|XP_017144712.1| PREDICTED: glutathione S-transferase D7 isoform X1 [Drosoph:
>gi|1037099566|ref|XP_017144713.1| PREDICTED: glutathione S-transferase D7 isoform X2 [Drosoph:
>gi|1037099542|ref|XP_017144711.1| PREDICTED: uncharacterized protein LOC108157248 [Drosophila
>gi|1037104798|ref|XP_017145264.1| PREDICTED: glutathione S-transferase 1-1 [Drosophila miranda
>gi|1037086086|ref|XP_017143663.1| PREDICTED: glutathione S-transferase D4-like [Drosophila mi
>gi|1037086128|ref|XP_017143666.1| PREDICTED: glutathione S-transferase D7-like [Drosophila mi
>gi|1037086100|ref|XP_017143664.1| PREDICTED: glutathione S-transferase 1-1 [Drosophila miranda
>gi|1037086114|ref|XP_017143665.1| PREDICTED: glutathione S-transferase 1-1-like [Drosophila m:
>gi|1037104981|ref|XP_017145290.1| PREDICTED: glutathione S-transferase 1-1-like [Drosophila m:
>gi|1037025041|ref|XP_017139588.1| PREDICTED: cadherin-89D [Drosophila miranda]
>gi|1037025059|ref|XP_017139589.1| PREDICTED: putative gustatory receptor 89a [Drosophila mirar
>gi|1037105734|ref|XP_017145399.1| PREDICTED: caspase-3 [Drosophila miranda]
>gi|1037080085|ref|XP_017143208.1| PREDICTED: collagen alpha chain CG42342 [Drosophila miranda]
>gi|1037080071|ref|XP_017143207.1| PREDICTED: putative peptidyl-tRNA hydrolase PTRHD1 [Drosoph:
>gi|1037104987|ref|XP_017145292.1| PREDICTED: calexcitin-2 [Drosophila miranda]
>gi|1037082364|ref|XP_017143388.1| PREDICTED: transmembrane protease serine 7 [Drosophila mirar
>gi|1037082283|ref|XP_017143381.1| PREDICTED: lysosome membrane protein 2 [Drosophila miranda]
>gi|1037082390|ref|XP_017143390.1| PREDICTED: neuropeptide F isoform X1 [Drosophila miranda]
>gi|1037082400|ref|XP_017143391.1| PREDICTED: neuropeptide F isoform X2 [Drosophila miranda]
>gi|1037082337|ref|XP_017143385.1| PREDICTED: synaptic vesicle glycoprotein 2C isoform X2 [Dros
>gi|1037082324|ref|XP_017143384.1| PREDICTED: 4-hydroxybenzoate transporter PcaK isoform X1 [D:
>gi|1037082376|ref|XP_017143389.1| PREDICTED: ATP synthase mitochondrial F1 complex assembly fa
>gi|1037082350|ref|XP_017143386.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037082311|ref|XP_017143383.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037082297|ref|XP_017143382.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037032903|ref|XP_017140066.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037053853|ref|XP_017141420.1| PREDICTED: uncharacterized protein LOC108155242 [Drosophila
>gi|1037053825|ref|XP_017141418.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1037053837|ref|XP_017141419.1| PREDICTED: histone-arginine methyltransferase CARMER [Drosop
>gi|1037025077|ref|XP_017139590.1| PREDICTED: putative gustatory receptor 85a [Drosophila mirar
>gi|1037025095|ref|XP_017139591.1| PREDICTED: uncharacterized protein LOC108154009 [Drosophila
>gi|1037098997|ref|XP_017144666.1| PREDICTED: tctex1 domain-containing protein 2 [Drosophila m:
>gi|1037060462|ref|XP_017141874.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060416|ref|XP_017141871.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060382|ref|XP_017141869.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060366|ref|XP_017141868.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060478|ref|XP_017141875.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060398|ref|XP_017141870.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060320|ref|XP_017141864.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060335|ref|XP_017141865.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060351|ref|XP_017141867.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
>gi|1037060446|ref|XP_017141873.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo:
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>gi|1037060430|ref|XP_017141872.1| PREDICTED: protein-methionine sulfoxide oxidase Mical isofo
>gi|1037060492|ref|XP_017141877.1| PREDICTED: uncharacterized protein LOC108155527 [Drosophila
>gi|1037025113|ref|XP_017139592.1| PREDICTED: uncharacterized protein LOC108154010, partial [Dr
>gi|1037089881|ref|XP_017143928.1| PREDICTED: WD repeat-containing protein 61 [Drosophila mira
>gi|1037089894|ref|XP_017143929.1| PREDICTED: protein NDUFAF4 homolog [Drosophila miranda]
>gi|1037089867|ref|XP_017143926.1| PREDICTED: dimethyladenosine transferase 2, mitochondrial [D
>gi|1037089905|ref|XP_017143930.1| PREDICTED: uncharacterized protein LOC108156774 [Drosophila
>gi|1037089156|ref|XP_017143869.1| PREDICTED: protein cereblon homolog [Drosophila miranda]
>gi|1037089168|ref|XP_017143870.1| PREDICTED: uncharacterized protein LOC108156727 [Drosophila
>gi|1037089181|ref|XP_017143872.1| PREDICTED: mite group 2 allergen Lep d 2 [Drosophila miranda
>gi|1037034288|ref|XP_017140149.1| PREDICTED: protein NPC2 homolog [Drosophila miranda]
>gi|1037036679|ref|XP_017140284.1| PREDICTED: uncharacterized protein LOC108154494 [Drosophila
>gi|1037025130|ref|XP_017139593.1| PREDICTED: uncharacterized protein LOC108154011 [Drosophila
>gi|1037025148|ref|XP_017139595.1| PREDICTED: uncharacterized protein LOC108154012 [Drosophila
>gi|1037039682|ref|XP_017140467.1| PREDICTED: uncharacterized protein LOC108154647 [Drosophila
>gi|1037040053|ref|XP_017140492.1| PREDICTED: uncharacterized protein LOC108154666 [Drosophila
>gi|1037025168|ref|XP_017139596.1| PREDICTED: uncharacterized protein LOC108154013 [Drosophila
>gi|1037025186|ref|XP_017139597.1| PREDICTED: uncharacterized protein LOC108154014 [Drosophila
>gi|1037038301|ref|XP_017140378.1| PREDICTED: angiomotin [Drosophila miranda]
>gi|1037098079|ref|XP_017144576.1| PREDICTED: uncharacterized protein LOC108157168 [Drosophila
>gi|1037086793|ref|XP_017143717.1| PREDICTED: fragile X mental retardation syndrome-related pro
>gi|1037086779|ref|XP_017143716.1| PREDICTED: fragile X mental retardation syndrome-related pro
>gi|1037086765|ref|XP_017143715.1| PREDICTED: fragile X mental retardation syndrome-related pro
>gi|1037097541|ref|XP_017144525.1| PREDICTED: putative carbonic anhydrase 3 isoform X2 [Drosop
>gi|1037097530|ref|XP_017144524.1| PREDICTED: putative carbonic anhydrase 3 isoform X1 [Drosop
>gi|1037034255|ref|XP_017140147.1| PREDICTED: carbonic anhydrase 6-like isoform X1 [Drosophila
>gi|1037034273|ref|XP_017140148.1| PREDICTED: carbonic anhydrase 6-like isoform X2 [Drosophila
>gi|1037100233|ref|XP_017144764.1| PREDICTED: SWI/SNF chromatin-remodeling complex subunit SNF5
>gi|1037100267|ref|XP_017144767.1| PREDICTED: chymotrypsin-2-like [Drosophila miranda]
>gi|1037100257|ref|XP_017144766.1| PREDICTED: chymotrypsin-2-like [Drosophila miranda]
>gi|1037100245|ref|XP_017144765.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037100278|ref|XP_017144768.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037100314|ref|XP_017144772.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037100290|ref|XP_017144770.1| PREDICTED: chymotrypsin-2 isoform X1 [Drosophila miranda]
>gi|1037100300|ref|XP_017144771.1| PREDICTED: chymotrypsin-1 isoform X2 [Drosophila miranda]
>gi|1037044657|ref|XP_017140798.1| PREDICTED: chymotrypsin-2-like isoform X2 [Drosophila miran
>gi|1037022557|ref|XP_017139429.1| PREDICTED: chymotrypsin-2, partial [Drosophila miranda]
>gi|1037044715|ref|XP_017140802.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037044671|ref|XP_017140799.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037044699|ref|XP_017140801.1| PREDICTED: chymotrypsin-2 isoform X2 [Drosophila miranda]
>gi|1037044683|ref|XP_017140800.1| PREDICTED: chymotrypsin-2 isoform X1 [Drosophila miranda]
>gi|1037044469|ref|XP_017140785.1| PREDICTED: filamin-A isoform X1 [Drosophila miranda]
>gi|1037044536|ref|XP_017140789.1| PREDICTED: filamin-A isoform X5 [Drosophila miranda]
>gi|1037044487|ref|XP_017140786.1| PREDICTED: filamin-A isoform X2 [Drosophila miranda]
>gi|1037044614|ref|XP_017140795.1| PREDICTED: rac GTPase-activating protein 1-like [Drosophila
>gi|1037044628|ref|XP_017140796.1| PREDICTED: chymotrypsin-2-like isoform X1 [Drosophila miran
>gi|1037044641|ref|XP_017140797.1| PREDICTED: chymotrypsin-2-like isoform X1 [Drosophila miran
>gi|1037044731|ref|XP_017140804.1| PREDICTED: uncharacterized protein LOC108154876 [Drosophila
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>gi|1037044502|ref|XP_017140787.1| PREDICTED: filamin-A isoform X3 [Drosophila miranda]
>gi|1037044520|ref|XP_017140788.1| PREDICTED: filamin-A isoform X4 [Drosophila miranda]
>gi|1037044568|ref|XP_017140792.1| PREDICTED: filamin-A isoform X7 [Drosophila miranda]
>gi|1037044552|ref|XP_017140790.1| PREDICTED: filamin-A isoform X6 [Drosophila miranda]
>gi|1037044584|ref|XP_017140793.1| PREDICTED: filamin-A isoform X8 [Drosophila miranda]
>gi|1037044600|ref|XP_017140794.1| PREDICTED: filamin-A isoform X9 [Drosophila miranda]
>gi|1037025204|ref|XP_017139598.1| PREDICTED: uncharacterized protein LOC108154016 [Drosophila
>gi|1037044747|ref|XP_017140805.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein ki
>gi|1037025218|ref|XP_017139599.1| PREDICTED: protein maelstrom 2-like [Drosophila miranda]
>gi|1037039358|ref|XP_017140442.1| PREDICTED: uncharacterized protein LOC108154630 [Drosophila
>gi|1037033292|ref|XP_017140089.1| PREDICTED: uncharacterized protein LOC108154339 [Drosophila
>gi|1037033308|ref|XP_017140090.1| PREDICTED: uncharacterized protein LOC108154340 [Drosophila
>gi|1037033326|ref|XP_017140091.1| PREDICTED: uncharacterized protein LOC108154340 [Drosophila
>gi|1037033344|ref|XP_017140092.1| PREDICTED: uncharacterized protein LOC108154340 [Drosophila
>gi|1037037433|ref|XP_017140327.1| PREDICTED: uncharacterized protein LOC108154535 [Drosophila
>gi|1037035234|ref|XP_017140203.1| PREDICTED: uncharacterized protein LOC108154442 [Drosophila
>gi|1037104309|ref|XP_017145196.1| PREDICTED: KAT8 regulatory NSL complex subunit 2-like isofo
>gi|1037104301|ref|XP_017145195.1| PREDICTED: KAT8 regulatory NSL complex subunit 2-like isofo
>gi|1037078699|ref|XP_017143115.1| PREDICTED: protein toll [Drosophila miranda]
>gi|1037070236|ref|XP_017142533.1| PREDICTED: cation-independent mannose-6-phosphate receptor
>gi|1037070220|ref|XP_017142532.1| PREDICTED: cation-independent mannose-6-phosphate receptor
>gi|1037070330|ref|XP_017142539.1| PREDICTED: uncharacterized protein LOC108155906 isoform X1
>gi|1037070345|ref|XP_017142540.1| PREDICTED: uncharacterized protein LOC108155906 isoform X2
>gi|1037070286|ref|XP_017142536.1| PREDICTED: uncharacterized protein DDB_G0271670 [Drosophila
>gi|1037070252|ref|XP_017142534.1| PREDICTED: nucleosomal histone kinase 1 [Drosophila miranda]
>gi|1037070268|ref|XP_017142535.1| PREDICTED: nucleosomal histone kinase 1 [Drosophila miranda]
>gi|1037070314|ref|XP_017142538.1| PREDICTED: uncharacterized protein LOC108155904 isoform X2
>gi|1037070302|ref|XP_017142537.1| PREDICTED: uncharacterized protein LOC108155904 isoform X1
>gi|1037079975|ref|XP_017143202.1| PREDICTED: histone H2A.v [Drosophila miranda]
>gi|1037079961|ref|XP_017143201.1| PREDICTED: SPARC [Drosophila miranda]
>gi|1037079947|ref|XP_017143200.1| PREDICTED: ribonucleoprotein RB97D isoform X2 [Drosophila m
>gi|1037079919|ref|XP_017143197.1| PREDICTED: ribonucleoprotein RB97D isoform X1 [Drosophila m
>gi|1037079933|ref|XP_017143199.1| PREDICTED: ribonucleoprotein RB97D isoform X1 [Drosophila m
>gi|1037074263|ref|XP_017142810.1| PREDICTED: LOW QUALITY PROTEIN: homeobox protein rough [Dro
>gi|1037074275|ref|XP_017142811.1| PREDICTED: oxidoreductase-like domain-containing protein 1
>gi|1037074249|ref|XP_017142809.1| PREDICTED: uncharacterized protein LOC108156048 [Drosophila
>gi|1037074234|ref|XP_017142808.1| PREDICTED: DNA-binding protein Ets97D [Drosophila miranda]
>gi|1037083059|ref|XP_017143432.1| PREDICTED: aminopeptidase N isoform X1 [Drosophila miranda]
>gi|1037083073|ref|XP_017143433.1| PREDICTED: aminopeptidase N isoform X1 [Drosophila miranda]
>gi|1037083087|ref|XP_017143434.1| PREDICTED: aminopeptidase N isoform X1 [Drosophila miranda]
>gi|1037083113|ref|XP_017143436.1| PREDICTED: aminopeptidase N isoform X3 [Drosophila miranda]
>gi|1037083099|ref|XP_017143435.1| PREDICTED: aminopeptidase N isoform X2 [Drosophila miranda]
>gi|1037083127|ref|XP_017143437.1| PREDICTED: aminopeptidase N isoform X4 [Drosophila miranda]
>gi|1037083139|ref|XP_017143438.1| PREDICTED: uncharacterized protein LOC108156476 [Drosophila
>gi|1037083152|ref|XP_017143440.1| PREDICTED: sperm-associated antigen 1 [Drosophila miranda]
>gi|1037083165|ref|XP_017143441.1| PREDICTED: sperm-associated antigen 1 [Drosophila miranda]
>gi|1037083179|ref|XP_017143442.1| PREDICTED: uridine phosphorylase 1 isoform X1 [Drosophila m
>gi|1037083193|ref|XP_017143443.1| PREDICTED: uridine phosphorylase 1 isoform X2 [Drosophila m
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>gi|1037083207|ref|XP_017143444.1| PREDICTED: uridine phosphorylase 1 isoform X2 [Drosophila mi
>gi|1037102108|ref|XP_017144942.1| PREDICTED: uridine phosphorylase 1 isoform X2 [Drosophila mi
>gi|1037102098|ref|XP_017144941.1| PREDICTED: uridine phosphorylase 1 isoform X1 [Drosophila mi
>gi|1037035341|ref|XP_017140208.1| PREDICTED: probable proteasome subunit beta type-2 [Drosophi
>gi|1037095272|ref|XP_017144325.1| PREDICTED: tetraspanin-31-A isoform X1 [Drosophila miranda]
>gi|1037095286|ref|XP_017144326.1| PREDICTED: tetraspanin-13 isoform X2 [Drosophila miranda]
>gi|1037102446|ref|XP_017144978.1| PREDICTED: uncharacterized protein LOC108157435 [Drosophila
>gi|1037036870|ref|XP_017140292.1| PREDICTED: uncharacterized protein LOC108154504 [Drosophila
>gi|1037104247|ref|XP_017145187.1| PREDICTED: uncharacterized protein LOC108157577 [Drosophila
>gi|1037035099|ref|XP_017140197.1| PREDICTED: bromodomain testis-specific protein-like [Drosoph
>gi|1037035115|ref|XP_017140198.1| PREDICTED: bromodomain testis-specific protein-like [Drosoph
>gi|1037035133|ref|XP_017140199.1| PREDICTED: bromodomain testis-specific protein-like [Drosoph
>gi|1037033967|ref|XP_017140129.1| PREDICTED: isochorismatase domain-containing protein 1 [Dros
>gi|1037103125|ref|XP_017145049.1| PREDICTED: methylthioribose-1-phosphate isomerase [Drosophi
>gi|1037103135|ref|XP_017145050.1| PREDICTED: methylthioribose-1-phosphate isomerase [Drosophi
>gi|1037103145|ref|XP_017145051.1| PREDICTED: methylthioribose-1-phosphate isomerase [Drosophi
>gi|1037103113|ref|XP_017145048.1| PREDICTED: lysyl oxidase homolog 2A [Drosophila miranda]
>gi|1037066434|ref|XP_017142271.1| PREDICTED: G protein-coupled receptor kinase 2 [Drosophila m
>gi|1037081037|ref|XP_017143283.1| PREDICTED: polyribonucleotide nucleotidyltransferase 1, mito
>gi|1037081009|ref|XP_017143281.1| PREDICTED: eIF-2-alpha kinase GCN2 [Drosophila miranda]
>gi|1037081023|ref|XP_017143282.1| PREDICTED: eIF-2-alpha kinase GCN2 [Drosophila miranda]
>gi|1037078072|ref|XP_017143074.1| PREDICTED: uncharacterized protein LOC108156237 [Drosophila
>gi|1037078045|ref|XP_017143071.1| PREDICTED: Y+L amino acid transporter 2 [Drosophila miranda]
>gi|1037078058|ref|XP_017143073.1| PREDICTED: putative glycogen synthase kinase-3 homolog [Dros
>gi|1037078002|ref|XP_017143068.1| PREDICTED: band 4.1-like protein 4A isoform X5 [Drosophila m
>gi|1037077989|ref|XP_017143067.1| PREDICTED: hornerin isoform X4 [Drosophila miranda]
>gi|1037077961|ref|XP_017143065.1| PREDICTED: filaggrin isoform X2 [Drosophila miranda]
>gi|1037077947|ref|XP_017143064.1| PREDICTED: filaggrin isoform X1 [Drosophila miranda]
>gi|1037077975|ref|XP_017143066.1| PREDICTED: filaggrin isoform X3 [Drosophila miranda]
>gi|1037078017|ref|XP_017143069.1| PREDICTED: glycine receptor subunit alpha-4 isoform X1 [Dros
>gi|1037078031|ref|XP_017143070.1| PREDICTED: glycine receptor subunit alpha-4 isoform X2 [Dros
>gi|1037025238|ref|XP_017139600.1| PREDICTED: acid-sensing ion channel 5 [Drosophila miranda]
>gi|1037025256|ref|XP_017139601.1| PREDICTED: carbonic anhydrase 14 [Drosophila miranda]
>gi|1037037611|ref|XP_017140337.1| PREDICTED: carbonic anhydrase 2-like [Drosophila miranda]
>gi|1037100209|ref|XP_017144762.1| PREDICTED: uncharacterized protein LOC108157275 [Drosophila
>gi|1037089483|ref|XP_017143894.1| PREDICTED: methionyl-tRNA formyltransferase, mitochondrial
>gi|1037089497|ref|XP_017143895.1| PREDICTED: methionyl-tRNA formyltransferase, mitochondrial
>gi|1037089546|ref|XP_017143896.1| PREDICTED: 39S ribosomal protein L32, mitochondrial [Drosoph
>gi|1037089469|ref|XP_017143893.1| PREDICTED: decaprenyl-diphosphate synthase subunit 1 [Drosop
>gi|1037040317|ref|XP_017140510.1| PREDICTED: uncharacterized protein LOC108154680 [Drosophila
>gi|1037025274|ref|XP_017139602.1| PREDICTED: uncharacterized protein LOC108154021, partial [Dr
>gi|1037106285|ref|XP_017145476.1| PREDICTED: uncharacterized protein LOC108157793 [Drosophila
>gi|1037106293|ref|XP_017145477.1| PREDICTED: uncharacterized protein LOC108157793 [Drosophila
>gi|1037095995|ref|XP_017144382.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037095973|ref|XP_017144380.1| PREDICTED: serine/threonine-protein kinase ULK3 isoform X1
>gi|1037095983|ref|XP_017144381.1| PREDICTED: serine/threonine-protein kinase ULK3 isoform X2
>gi|1037049890|ref|XP_017141148.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila m
>gi|1037049934|ref|XP_017141150.1| PREDICTED: putative uncharacterized protein DDB_G0293878 iso
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>gi|1037049948|ref|XP_017141151.1| PREDICTED: serine-rich adhesin for platelets isoform X5 [Dro
>gi|1037049876|ref|XP_017141147.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila n
>gi|1037049920|ref|XP_017141149.1| PREDICTED: putative uncharacterized protein DDB_G0293878 iso
>gi|1037050006|ref|XP_017141156.1| PREDICTED: S1 RNA-binding domain-containing protein 1 [Dros
>gi|1037050054|ref|XP_017141159.1| PREDICTED: uncharacterized protein LOC108155076 [Drosophila
>gi|1037050022|ref|XP_017141157.1| PREDICTED: ribosome-releasing factor 2, mitochondrial isofor
>gi|1037050038|ref|XP_017141158.1| PREDICTED: ribosome-releasing factor 2, mitochondrial isofor
>gi|1037049962|ref|XP_017141153.1| PREDICTED: uncharacterized protein PFB0145c isoform X1 [Dros
>gi|1037049974|ref|XP_017141154.1| PREDICTED: uncharacterized protein PFB0145c isoform X1 [Dros
>gi|1037049990|ref|XP_017141155.1| PREDICTED: keratin, type I cytoskeletal 9 isoform X2 [Drosop
>gi|1037063425|ref|XP_017142076.1| PREDICTED: probable 39S ribosomal protein L45, mitochondrial
>gi|1037063454|ref|XP_017142079.1| PREDICTED: U-actitoxin-Avd3s-like [Drosophila miranda]
>gi|1037063438|ref|XP_017142078.1| PREDICTED: kappaPI-actitoxin-Avd3c-like [Drosophila miranda]
>gi|1037063314|ref|XP_017142069.1| PREDICTED: regulator of G-protein signaling loco isoform X1
>gi|1037063380|ref|XP_017142073.1| PREDICTED: regulator of G-protein signaling loco isoform X4
>gi|1037063350|ref|XP_017142071.1| PREDICTED: regulator of G-protein signaling loco isoform X3
>gi|1037063366|ref|XP_017142072.1| PREDICTED: regulator of G-protein signaling loco isoform X3
>gi|1037063395|ref|XP_017142074.1| PREDICTED: regulator of G-protein signaling loco isoform X5
>gi|1037063411|ref|XP_017142075.1| PREDICTED: regulator of G-protein signaling loco isoform X6
>gi|1037063332|ref|XP_017142070.1| PREDICTED: regulator of G-protein signaling loco isoform X2
>gi|1037051401|ref|XP_017141255.1| PREDICTED: LOW QUALITY PROTEIN: aprataxin and PNK-like facto
>gi|1037051417|ref|XP_017141256.1| PREDICTED: translin-associated protein X [Drosophila miranda
>gi|1037051320|ref|XP_017141250.1| PREDICTED: conserved oligomeric Golgi complex subunit 2 [Dro
>gi|1037051365|ref|XP_017141253.1| PREDICTED: 3-hydroxyisobutyryl-CoA hydrolase, mitochondrial
>gi|1037051383|ref|XP_017141254.1| PREDICTED: 3-hydroxyisobutyryl-CoA hydrolase, mitochondrial
>gi|1037051352|ref|XP_017141252.1| PREDICTED: cysteine protease ATG4C [Drosophila miranda]
>gi|1037051336|ref|XP_017141251.1| PREDICTED: transmembrane and TPR repeat-containing protein C
>gi|1037051505|ref|XP_017141262.1| PREDICTED: trafficking protein particle complex subunit 6B
>gi|1037051482|ref|XP_017141261.1| PREDICTED: uncharacterized protein LOC108155128 [Drosophila
>gi|1037051464|ref|XP_017141259.1| PREDICTED: zinc finger protein 28 [Drosophila miranda]
>gi|1037051449|ref|XP_017141258.1| PREDICTED: NF-kappa-B-repressing factor isoform X2 [Drosophi
>gi|1037051433|ref|XP_017141257.1| PREDICTED: NF-kappa-B-repressing factor isoform X1 [Drosophi
>gi|1037081571|ref|XP_017143324.1| PREDICTED: uncharacterized protein LOC108156398 [Drosophila
>gi|1037081599|ref|XP_017143326.1| PREDICTED: surfeit locus protein 4 homolog isoform X2 [Dros
>gi|1037081585|ref|XP_017143325.1| PREDICTED: surfeit locus protein 4 homolog isoform X1 [Dros
>gi|1037081613|ref|XP_017143327.1| PREDICTED: protein BUD31 homolog [Drosophila miranda]
>gi|1037078667|ref|XP_017143113.1| PREDICTED: gastrula zinc finger protein XlCGF46.1 [Drosophil
>gi|1037078653|ref|XP_017143112.1| PREDICTED: 28S ribosomal protein S22, mitochondrial [Drosoph
>gi|1037078639|ref|XP_017143111.1| PREDICTED: uncharacterized protein LOC108156259 [Drosophila
>gi|1037078685|ref|XP_017143114.1| PREDICTED: nucleosome assembly protein 1-like 1 [Drosophila
>gi|1037100762|ref|XP_017144815.1| PREDICTED: uncharacterized protein LOC108157317 [Drosophila
>gi|1037025292|ref|XP_017139603.1| PREDICTED: odorant receptor 98a [Drosophila miranda]
>gi|1037100716|ref|XP_017144809.1| PREDICTED: opioid-binding protein/cell adhesion molecule iso
>gi|1037100726|ref|XP_017144810.1| PREDICTED: opioid-binding protein/cell adhesion molecule iso
>gi|1037100738|ref|XP_017144811.1| PREDICTED: opioid-binding protein/cell adhesion molecule iso
>gi|1037100748|ref|XP_017144814.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1037102905|ref|XP_017145023.1| PREDICTED: DNA-binding protein D-ETS-4 [Drosophila miranda]
>gi|1037102547|ref|XP_017144988.1| PREDICTED: eukaryotic translation initiation factor 2 subuni
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>gi|1037054571|ref|XP_017141471.1| PREDICTED: RNA exonuclease 1 homolog [Drosophila miranda]
>gi|1037054621|ref|XP_017141474.1| PREDICTED: peroxisomal carnitine O-octanoyltransferase [Dros
>gi|1037054631|ref|XP_017141475.1| PREDICTED: peroxisomal carnitine O-octanoyltransferase [Dros
>gi|1037054647|ref|XP_017141476.1| PREDICTED: peroxisomal carnitine O-octanoyltransferase [Dros
>gi|1037054665|ref|XP_017141477.1| PREDICTED: peroxisomal carnitine O-octanoyltransferase [Dros
>gi|1037054605|ref|XP_017141473.1| PREDICTED: protein CASC3 [Drosophila miranda]
>gi|1037054681|ref|XP_017141478.1| PREDICTED: HSPB1-associated protein 1 [Drosophila miranda]
>gi|1037054589|ref|XP_017141472.1| PREDICTED: programmed cell death 6-interacting protein [Dros
>gi|1037101937|ref|XP_017144925.1| PREDICTED: uncharacterized protein LOC108157396 [Drosophila
>gi|1037088663|ref|XP_017143831.1| PREDICTED: E3 ubiquitin-protein ligase RING1 isoform X1 [Dro
>gi|1037088677|ref|XP_017143832.1| PREDICTED: E3 ubiquitin-protein ligase RING1 isoform X2 [Dro
>gi|1037088705|ref|XP_017143834.1| PREDICTED: uncharacterized protein LOC108156698 [Drosophila
>gi|1037088691|ref|XP_017143833.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 [Dros
>gi|1037089455|ref|XP_017143892.1| PREDICTED: vacuolar protein sorting-associated protein 72 ho
>gi|1037089441|ref|XP_017143891.1| PREDICTED: protein unc-80 homolog isoform X3 [Drosophila mir
>gi|1037089414|ref|XP_017143889.1| PREDICTED: protein unc-80 homolog isoform X1 [Drosophila mir
>gi|1037089427|ref|XP_017143890.1| PREDICTED: protein unc-80 homolog isoform X2 [Drosophila mir
>gi|1037104834|ref|XP_017145270.1| PREDICTED: myosin light chain alkali isoform X2 [Drosophila
>gi|1037104826|ref|XP_017145269.1| PREDICTED: myosin light chain alkali isoform X1 [Drosophila
>gi|1037087503|ref|XP_017143740.1| PREDICTED: microtubule-associated protein 2 isoform X2 [Dros
>gi|1037087526|ref|XP_017143742.1| PREDICTED: microtubule-associated protein 2 isoform X3 [Dros
>gi|1037087567|ref|XP_017143745.1| PREDICTED: microtubule-associated protein 2 isoform X6 [Dros
>gi|1037087581|ref|XP_017143746.1| PREDICTED: microtubule-associated protein 2 isoform X7 [Dros
>gi|1037087607|ref|XP_017143748.1| PREDICTED: microtubule-associated protein tau isoform X9 [D
>gi|1037087539|ref|XP_017143743.1| PREDICTED: microtubule-associated protein 2 isoform X4 [Dros
>gi|1037087553|ref|XP_017143744.1| PREDICTED: microtubule-associated protein 2 isoform X5 [Dros
>gi|1037087595|ref|XP_017143747.1| PREDICTED: microtubule-associated protein tau isoform X8 [D
>gi|1037087489|ref|XP_017143739.1| PREDICTED: extensin isoform X1 [Drosophila miranda]
>gi|1037087620|ref|XP_017143749.1| PREDICTED: ras-related C3 botulinum toxin substrate 1 [Dros
>gi|1037089694|ref|XP_017143912.1| PREDICTED: probable enoyl-CoA hydratase [Drosophila miranda]
>gi|1037089680|ref|XP_017143909.1| PREDICTED: tubby-related protein 4 isoform X4 [Drosophila m
>gi|1037089660|ref|XP_017143907.1| PREDICTED: tubby-related protein 4 isoform X2 [Drosophila m
>gi|1037089672|ref|XP_017143908.1| PREDICTED: tubby-related protein 4 isoform X3 [Drosophila m
>gi|1037089648|ref|XP_017143906.1| PREDICTED: tubby-related protein 4 isoform X1 [Drosophila m
>gi|1037089688|ref|XP_017143910.1| PREDICTED: uncharacterized protein LOC108156759 [Drosophila
>gi|1037084522|ref|XP_017143549.1| PREDICTED: attractin-like protein 1 [Drosophila miranda]
>gi|1037087023|ref|XP_017143735.1| PREDICTED: uncharacterized protein LOC108156644 [Drosophila
>gi|1037087037|ref|XP_017143736.1| PREDICTED: G-protein-signaling modulator 2 [Drosophila mira
>gi|1037083880|ref|XP_017143499.1| PREDICTED: splicing factor 1 isoform X1 [Drosophila miranda]
>gi|1037083910|ref|XP_017143501.1| PREDICTED: pre-mRNA-splicing factor 38B isoform X3 [Drosophi
>gi|1037083896|ref|XP_017143500.1| PREDICTED: splicing factor 1 isoform X2 [Drosophila miranda]
>gi|1037083924|ref|XP_017143502.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1037103496|ref|XP_017145089.1| PREDICTED: homeobox protein Hmx [Drosophila miranda]
>gi|1037103504|ref|XP_017145090.1| PREDICTED: uncharacterized protein LOC108157501 [Drosophila
>gi|1037034877|ref|XP_017140184.1| PREDICTED: salivary glue protein Sgs-5 [Drosophila miranda]
>gi|1037036619|ref|XP_017140279.1| PREDICTED: chorion class high-cysteine HCB protein 13 [Dros
>gi|1037025307|ref|XP_017139605.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5-like [Drosophi
>gi|1037105983|ref|XP_017145434.1| PREDICTED: homeobox protein slou [Drosophila miranda]
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>gi|1037076949|ref|XP_017142991.1| PREDICTED: DNA polymerase epsilon catalytic subunit A [Dros
>gi|1037076964|ref|XP_017142992.1| PREDICTED: ATP synthase-coupling factor 6, mitochondrial [D
>gi|1037091940|ref|XP_017144086.1| PREDICTED: uncharacterized protein LOC108156869 [Drosophila
>gi|1037091926|ref|XP_017144085.1| PREDICTED: protein SEC13 homolog [Drosophila miranda]
>gi|1037025321|ref|XP_017139606.1| PREDICTED: uncharacterized protein LOC108154024 [Drosophila
>gi|1037106766|ref|XP_017145542.1| PREDICTED: uncharacterized protein LOC108157849 [Drosophila
>gi|1037091748|ref|XP_017144073.1| PREDICTED: sodium-independent sulfate anion transporter [Dro
>gi|1037050961|ref|XP_017141226.1| PREDICTED: heterogeneous nuclear ribonucleoprotein A1 isofo
>gi|1037050972|ref|XP_017141227.1| PREDICTED: heterogeneous nuclear ribonucleoprotein A1 isofo
>gi|1037050988|ref|XP_017141228.1| PREDICTED: heterogeneous nuclear ribonucleoprotein A1 isofo
>gi|1037050947|ref|XP_017141225.1| PREDICTED: protein C12orf4 homolog [Drosophila miranda]
>gi|1037050933|ref|XP_017141224.1| PREDICTED: ankyrin repeat domain-containing protein 50 isof
>gi|1037050917|ref|XP_017141223.1| PREDICTED: ankyrin repeat domain-containing protein 50 isof
>gi|1037106508|ref|XP_017145506.1| PREDICTED: uncharacterized protein LOC108157818 [Drosophila
>gi|1037032551|ref|XP_017140044.1| PREDICTED: uncharacterized protein LOC108154301 [Drosophila
>gi|1037025339|ref|XP_017139607.1| PREDICTED: uncharacterized protein LOC108154025 [Drosophila
>gi|1037025357|ref|XP_017139608.1| PREDICTED: uncharacterized protein LOC108154026 [Drosophila
>gi|1037101841|ref|XP_017144914.1| PREDICTED: transcription initiation factor TFIID subunit 5-
>gi|1037101831|ref|XP_017144913.1| PREDICTED: transcription initiation factor TFIID subunit 5-
>gi|1037025375|ref|XP_017139609.1| PREDICTED: uncharacterized protein LOC108154027 [Drosophila
>gi|1037100562|ref|XP_017144794.1| PREDICTED: ABC transporter G family member 23 isoform X1 [D
>gi|1037100572|ref|XP_017144795.1| PREDICTED: ABC transporter G family member 20 isoform X2 [D
>gi|1037100584|ref|XP_017144796.1| PREDICTED: ABC transporter G family member 23 isoform X3 [D
>gi|1037100594|ref|XP_017144797.1| PREDICTED: ABC transporter G family member 20 isoform X4 [D
>gi|1037071139|ref|XP_017142595.1| PREDICTED: uncharacterized protein LOC108155943 isoform X1
>gi|1037071155|ref|XP_017142596.1| PREDICTED: uncharacterized protein LOC108155943 isoform X2
>gi|1037105374|ref|XP_017145349.1| PREDICTED: allatostatin-A receptor isoform X2 [Drosophila m
>gi|1037105366|ref|XP_017145348.1| PREDICTED: allatostatin-A receptor isoform X1 [Drosophila m
>gi|1037105884|ref|XP_017145420.1| PREDICTED: putative polypeptide N-acetylgalactosaminyltrans
>gi|1037025393|ref|XP_017139610.1| PREDICTED: uncharacterized protein LOC108154028 [Drosophila
>gi|1037025411|ref|XP_017139611.1| PREDICTED: uncharacterized protein LOC108154029 [Drosophila
>gi|1037105787|ref|XP_017145407.1| PREDICTED: trypsin [Drosophila miranda]
>gi|1037106324|ref|XP_017145480.1| PREDICTED: uncharacterized protein LOC108157798 [Drosophila
>gi|1037106316|ref|XP_017145479.1| PREDICTED: uncharacterized protein LOC108157797 [Drosophila
>gi|1037084810|ref|XP_017143572.1| PREDICTED: uncharacterized protein LOC108156558 [Drosophila
>gi|1037084796|ref|XP_017143571.1| PREDICTED: serendipity locus protein H-1 [Drosophila miranda
>gi|1037052202|ref|XP_017141309.1| PREDICTED: pre-mRNA-processing factor 39 [Drosophila miranda
>gi|1037052233|ref|XP_017141311.1| PREDICTED: DDB1- and CUL4-associated factor 10 homolog isof
>gi|1037052217|ref|XP_017141310.1| PREDICTED: DDB1- and CUL4-associated factor 10 homolog isof
>gi|1037052186|ref|XP_017141308.1| PREDICTED: RE1-silencing transcription factor [Drosophila m
>gi|1037052249|ref|XP_017141312.1| PREDICTED: neural Wiskott-Aldrich syndrome protein isoform
>gi|1037052265|ref|XP_017141313.1| PREDICTED: neural Wiskott-Aldrich syndrome protein isoform
>gi|1037052281|ref|XP_017141314.1| PREDICTED: neural Wiskott-Aldrich syndrome protein isoform
>gi|1037030889|ref|XP_017139946.1| PREDICTED: uncharacterized protein CG13380-like [Drosophila
>gi|1037105947|ref|XP_017145429.1| PREDICTED: octopamine receptor beta-1R isoform X1 [Drosophi
>gi|1037105953|ref|XP_017145430.1| PREDICTED: octopamine receptor beta-1R isoform X1 [Drosophi
>gi|1037105961|ref|XP_017145431.1| PREDICTED: octopamine receptor beta-1R isoform X2 [Drosophi
>gi|1037090197|ref|XP_017143954.1| PREDICTED: gibberellin 20 oxidase 1-like [Drosophila miranda
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>gi|1037090211|ref|XP_017143955.1| PREDICTED: gibberellin 20 oxidase 1-like [Drosophila miranda
>gi|1037103824|ref|XP_017145132.1| PREDICTED: UPF0676 protein C1494.01-like [Drosophila miranda
>gi|1037105412|ref|XP_017145354.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037092146|ref|XP_017144102.1| PREDICTED: adiponectin receptor protein isoform X1 [Drosophi
>gi|1037092162|ref|XP_017144103.1| PREDICTED: adiponectin receptor protein isoform X1 [Drosophi
>gi|1037092189|ref|XP_017144104.1| PREDICTED: adiponectin receptor protein isoform X2 [Drosophi
>gi|1037104183|ref|XP_017145178.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037104189|ref|XP_017145179.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037099270|ref|XP_017144684.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037099284|ref|XP_017144685.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037099235|ref|XP_017144681.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037099248|ref|XP_017144682.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037099259|ref|XP_017144683.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037089321|ref|XP_017143882.1| PREDICTED: COP9 signalosome complex subunit 6 [Drosophila m
>gi|1037089334|ref|XP_017143883.1| PREDICTED: diphthine methyl ester synthase [Drosophila miran
>gi|1037089347|ref|XP_017143884.1| PREDICTED: uncharacterized protein LOC108156736 [Drosophila
>gi|1037098328|ref|XP_017144604.1| PREDICTED: MKI67 FHA domain-interacting nucleolar phosphopro
>gi|1037098340|ref|XP_017144606.1| PREDICTED: homeobox protein MOX-1 [Drosophila miranda]
>gi|1037078921|ref|XP_017143132.1| PREDICTED: uncharacterized protein LOC108156274 [Drosophila
>gi|1037078935|ref|XP_017143133.1| PREDICTED: uncharacterized protein LOC108156274 [Drosophila
>gi|1037078949|ref|XP_017143134.1| PREDICTED: AN1-type zinc finger protein 6 isoform X1 [Drosop
>gi|1037078903|ref|XP_017143131.1| PREDICTED: uncharacterized protein LOC108156273 [Drosophila
>gi|1037078963|ref|XP_017143136.1| PREDICTED: AN1-type zinc finger protein 6 isoform X2 [Drosop
>gi|1037096605|ref|XP_017144442.1| PREDICTED: phosphotriesterase-related protein [Drosophila m
>gi|1037067165|ref|XP_017142318.1| PREDICTED: L-threonine ammonia-lyase [Drosophila miranda]
>gi|1037067149|ref|XP_017142317.1| PREDICTED: tyrosine-protein kinase Fer isoform X4 [Drosophil
>gi|1037067135|ref|XP_017142316.1| PREDICTED: tyrosine-protein kinase Fer isoform X3 [Drosophil
>gi|1037067087|ref|XP_017142312.1| PREDICTED: tyrosine-protein kinase Fer isoform X1 [Drosophil
>gi|1037067103|ref|XP_017142313.1| PREDICTED: tyrosine-protein kinase Fer isoform X1 [Drosophil
>gi|1037067119|ref|XP_017142314.1| PREDICTED: tyrosine-protein kinase Fer isoform X2 [Drosophil
>gi|1037067296|ref|XP_017142326.1| PREDICTED: uncharacterized protein LOC108155778 [Drosophila
>gi|1037067201|ref|XP_017142320.1| PREDICTED: odorant receptor 85f isoform X1 [Drosophila miran
>gi|1037067216|ref|XP_017142321.1| PREDICTED: odorant receptor 85f isoform X1 [Drosophila miran
>gi|1037067232|ref|XP_017142322.1| PREDICTED: odorant receptor 85f isoform X2 [Drosophila miran
>gi|1037067312|ref|XP_017142327.1| PREDICTED: UPF0562 protein C7orf55 homolog [Drosophila miran
>gi|1037067280|ref|XP_017142325.1| PREDICTED: 60S ribosomal protein L34 [Drosophila miranda]
>gi|1037067246|ref|XP_017142323.1| PREDICTED: omega-amidase NIT2 isoform X1 [Drosophila miranda
>gi|1037067262|ref|XP_017142324.1| PREDICTED: omega-amidase NIT2 isoform X2 [Drosophila miranda
>gi|1037067185|ref|XP_017142319.1| PREDICTED: uncharacterized protein LOC108155773 [Drosophila
>gi|1037067424|ref|XP_017142335.1| PREDICTED: BBSome-interacting protein 1 [Drosophila miranda]
>gi|1037067376|ref|XP_017142332.1| PREDICTED: LMBR1 domain-containing protein 2 homolog [Drosop
>gi|1037067392|ref|XP_017142333.1| PREDICTED: tubulin beta-2 chain [Drosophila miranda]
>gi|1037067408|ref|XP_017142334.1| PREDICTED: two pore potassium channel protein sup-9 [Drosoph
>gi|1037067360|ref|XP_017142331.1| PREDICTED: LOW QUALITY PROTEIN: alpha-mannosidase 2 [Drosoph
>gi|1037073064|ref|XP_017142726.1| PREDICTED: RNA-binding protein Nova-1 isoform X11 [Drosophil
>gi|1037073016|ref|XP_017142723.1| PREDICTED: RNA-binding protein Nova-1 isoform X8 [Drosophila
>gi|1037073048|ref|XP_017142725.1| PREDICTED: RNA-binding protein Nova-1 isoform X10 [Drosophil
>gi|1037072847|ref|XP_017142710.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
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>gi|1037072863|ref|XP_017142711.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
>gi|1037072879|ref|XP_017142712.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
>gi|1037072895|ref|XP_017142713.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
>gi|1037072911|ref|XP_017142715.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
>gi|1037072927|ref|XP_017142716.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
>gi|1037072943|ref|XP_017142717.1| PREDICTED: RNA-binding protein Nova-1 isoform X3 [Drosophila
>gi|1037073002|ref|XP_017142722.1| PREDICTED: RNA-binding protein Nova-1 isoform X7 [Drosophila
>gi|1037073032|ref|XP_017142724.1| PREDICTED: RNA-binding protein Nova-1 isoform X9 [Drosophila
>gi|1037072959|ref|XP_017142718.1| PREDICTED: RNA-binding protein Nova-1 isoform X4 [Drosophila
>gi|1037072987|ref|XP_017142721.1| PREDICTED: RNA-binding protein Nova-1 isoform X6 [Drosophila
>gi|1037072973|ref|XP_017142719.1| PREDICTED: RNA-binding protein Nova-1 isoform X5 [Drosophila
>gi|1037072785|ref|XP_017142706.1| PREDICTED: RNA-binding protein Nova-1 isoform X1 [Drosophila
>gi|1037072799|ref|XP_017142707.1| PREDICTED: RNA-binding protein Nova-1 isoform X1 [Drosophila
>gi|1037072815|ref|XP_017142708.1| PREDICTED: RNA-binding protein Nova-1 isoform X1 [Drosophila
>gi|1037072831|ref|XP_017142709.1| PREDICTED: RNA-binding protein Nova-1 isoform X2 [Drosophila
>gi|1037105742|ref|XP_017145400.1| PREDICTED: probable maleylacetoacetate isomerase 1 isoform X
>gi|1037105750|ref|XP_017145401.1| PREDICTED: probable maleylacetoacetate isomerase 1 isoform X
>gi|1037104255|ref|XP_017145188.1| PREDICTED: probable maleylacetoacetate isomerase 2 isoform X
>gi|1037104265|ref|XP_017145189.1| PREDICTED: probable maleylacetoacetate isomerase 2 isoform X
>gi|1037088036|ref|XP_017143782.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 1 [
>gi|1037099650|ref|XP_017144721.1| PREDICTED: uncharacterized protein LOC108157254 [Drosophila
>gi|1037079830|ref|XP_017143190.1| PREDICTED: zinc finger and SCAN domain-containing protein 12
>gi|1037079818|ref|XP_017143189.1| PREDICTED: arf-GAP with coiled-coil, ANK repeat and PH doma
>gi|1037079808|ref|XP_017143188.1| PREDICTED: arf-GAP with coiled-coil, ANK repeat and PH doma
>gi|1037090239|ref|XP_017143957.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1037090737|ref|XP_017143997.1| PREDICTED: probable WRKY transcription factor protein 1 iso
>gi|1037090751|ref|XP_017143998.1| PREDICTED: probable WRKY transcription factor protein 1 iso
>gi|1037090792|ref|XP_017144001.1| PREDICTED: aminoacylase-1-like [Drosophila miranda]
>gi|1037090806|ref|XP_017144002.1| PREDICTED: aminoacylase-1-like [Drosophila miranda]
>gi|1037090765|ref|XP_017143999.1| PREDICTED: aminoacylase-1-like [Drosophila miranda]
>gi|1037090778|ref|XP_017144000.1| PREDICTED: aminoacylase-1-like [Drosophila miranda]
>gi|1037025429|ref|XP_017139612.1| PREDICTED: vacuole membrane protein 1-like [Drosophila mira
>gi|1037096147|ref|XP_017144398.1| PREDICTED: uncharacterized protein LOC108157074 [Drosophila
>gi|1037096158|ref|XP_017144399.1| PREDICTED: ubiquitin-like domain-containing CTD phosphatase
>gi|1037084201|ref|XP_017143524.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 13
>gi|1037084214|ref|XP_017143525.1| PREDICTED: glutathione-specific gamma-glutamylcyclotransfera
>gi|1037084227|ref|XP_017143526.1| PREDICTED: glutathione-specific gamma-glutamylcyclotransfera
>gi|1037084239|ref|XP_017143527.1| PREDICTED: glutathione-specific gamma-glutamylcyclotransfera
>gi|1037083994|ref|XP_017143508.1| PREDICTED: myogenic-determination protein [Drosophila miran
>gi|1037084006|ref|XP_017143509.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1037084020|ref|XP_017143510.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1037083952|ref|XP_017143504.1| PREDICTED: glucosylceramidase isoform X1 [Drosophila miranda
>gi|1037083966|ref|XP_017143505.1| PREDICTED: glucosylceramidase isoform X2 [Drosophila miranda
>gi|1037083980|ref|XP_017143506.1| PREDICTED: glucosylceramidase [Drosophila miranda]
>gi|1037084060|ref|XP_017143513.1| PREDICTED: outer dense fiber protein 3-B [Drosophila miranda
>gi|1037083938|ref|XP_017143503.1| PREDICTED: synaptic vesicular amine transporter [Drosophila
>gi|1037084048|ref|XP_017143512.1| PREDICTED: histone chaperone asf1-like [Drosophila miranda]
>gi|1037084034|ref|XP_017143511.1| PREDICTED: outer dense fiber protein 3-B [Drosophila miranda
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>gi|1037101619|ref|XP_017144891.1| PREDICTED: uncharacterized protein LOC108157374 [Drosophila
>gi|1037063180|ref|XP_017142059.1| PREDICTED: (E3-independent) E2 ubiquitin-conjugating enzyme
>gi|1037063225|ref|XP_017142062.1| PREDICTED: serine protease easter-like isoform X2 [Drosophil
>gi|1037063210|ref|XP_017142061.1| PREDICTED: serine protease easter-like isoform X1 [Drosophil
>gi|1037063239|ref|XP_017142064.1| PREDICTED: serine protease easter-like [Drosophila miranda]
>gi|1037063194|ref|XP_017142060.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037098164|ref|XP_017144586.1| PREDICTED: GILT-like protein F37H8.5 [Drosophila miranda]
>gi|1037098154|ref|XP_017144585.1| PREDICTED: gamma-interferon-inducible lysosomal thiol reduct
>gi|1037025449|ref|XP_017139613.1| PREDICTED: G2/mitotic-specific cyclin-B-like [Drosophila mi
>gi|1037088444|ref|XP_017143813.1| PREDICTED: extracellular serine/threonine protein CG31145 is
>gi|1037088458|ref|XP_017143814.1| PREDICTED: extracellular serine/threonine protein CG31145 is
>gi|1037104863|ref|XP_017145275.1| PREDICTED: probable low-specificity L-threonine aldolase 2
>gi|1037105140|ref|XP_017145314.1| PREDICTED: nose resistant to fluoxetine protein 6-like isofc
>gi|1037105148|ref|XP_017145315.1| PREDICTED: nose resistant to fluoxetine protein 6-like isofc
>gi|1037105130|ref|XP_017145313.1| PREDICTED: nose resistant to fluoxetine protein 6-like [Dros
>gi|1037037362|ref|XP_017140322.1| PREDICTED: LOW QUALITY PROTEIN: nose resistant to fluoxetine
>gi|1037106007|ref|XP_017145437.1| PREDICTED: uncharacterized protein LOC108157766 [Drosophila
>gi|1037106410|ref|XP_017145492.1| PREDICTED: uncharacterized protein LOC108157808 [Drosophila
>gi|1037106417|ref|XP_017145493.1| PREDICTED: uncharacterized protein LOC108157808 [Drosophila
>gi|1037025465|ref|XP_017139614.1| PREDICTED: uncharacterized protein LOC108154033 [Drosophila
>gi|1037106664|ref|XP_017145528.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1037098800|ref|XP_017144649.1| PREDICTED: esterase B1 isoform X1 [Drosophila miranda]
>gi|1037098811|ref|XP_017144650.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1037098823|ref|XP_017144651.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1037098833|ref|XP_017144652.1| PREDICTED: esterase B1 isoform X3 [Drosophila miranda]
>gi|1037025481|ref|XP_017139616.1| PREDICTED: serine/threonine-protein kinase zyg-8 [Drosophila
>gi|1037098846|ref|XP_017144653.1| PREDICTED: LOW QUALITY PROTEIN: G protein-activated inward r
>gi|1037036887|ref|XP_017140294.1| PREDICTED: uncharacterized RING finger protein T02C1.1 [Dros
>gi|1037040331|ref|XP_017140511.1| PREDICTED: uncharacterized protein LOC108154681 [Drosophila
>gi|1037025498|ref|XP_017139617.1| PREDICTED: uncharacterized protein LOC108154035 [Drosophila
>gi|1037025511|ref|XP_017139618.1| PREDICTED: chymotrypsin-1-like [Drosophila miranda]
>gi|1037099892|ref|XP_017144743.1| PREDICTED: putative uncharacterized protein DDB_G0277255 [D
>gi|1037025527|ref|XP_017139619.1| PREDICTED: uncharacterized protein LOC108154037 [Drosophila
>gi|1037106243|ref|XP_017145470.1| PREDICTED: uncharacterized protein LOC108157789 [Drosophila
>gi|1037106251|ref|XP_017145471.1| PREDICTED: uncharacterized protein LOC108157791 [Drosophila
>gi|1037025544|ref|XP_017139620.1| PREDICTED: U-actitoxin-Avd3f [Drosophila miranda]
>gi|1037103094|ref|XP_017145046.1| PREDICTED: ankyrin repeat and SAM domain-containing protein
>gi|1037103104|ref|XP_017145047.1| PREDICTED: ankyrin repeat and SAM domain-containing protein
>gi|1037037540|ref|XP_017140334.1| PREDICTED: ATP-binding cassette sub-family A member 17-like
>gi|1037025562|ref|XP_017139621.1| PREDICTED: ABC transporter ced-7-like [Drosophila miranda]
>gi|1037095200|ref|XP_017144320.1| PREDICTED: LOW QUALITY PROTEIN: G2/mitotic-specific cyclin-E
>gi|1037038187|ref|XP_017140370.1| PREDICTED: kunitz-type serine protease inhibitor 2-like [Dr
>gi|1037105636|ref|XP_017145385.1| PREDICTED: uncharacterized protein LOC108157726 isoform X1
>gi|1037105642|ref|XP_017145386.1| PREDICTED: uncharacterized protein LOC108157726 isoform X2
>gi|1037025579|ref|XP_017139623.1| PREDICTED: LOW QUALITY PROTEIN: ATP-binding cassette sub-fam
>gi|1037101057|ref|XP_017144845.1| PREDICTED: fatty-acid amide hydrolase 2 [Drosophila miranda]
>gi|1037084564|ref|XP_017143553.1| PREDICTED: transcription factor A, mitochondrial isoform X2
>gi|1037084550|ref|XP_017143552.1| PREDICTED: transcription factor A, mitochondrial isoform X1
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>gi|1037084536|ref|XP_017143550.1| PREDICTED: esterase GA18864 [Drosophila miranda]
>gi|1037084578|ref|XP_017143554.1| PREDICTED: transcription elongation factor B polypeptide 2
>gi|1037101365|ref|XP_017144871.1| PREDICTED: signal recognition particle 14 kDa protein [Droso
>gi|1037101331|ref|XP_017144870.1| PREDICTED: uncharacterized protein LOC108157358 [Drosophila
>gi|1037092588|ref|XP_017144131.1| PREDICTED: uncharacterized protein LOC108156900 [Drosophila
>gi|1037092601|ref|XP_017144132.1| PREDICTED: uncharacterized protein LOC108156900 [Drosophila
>gi|1037096709|ref|XP_017144448.1| PREDICTED: septin-2 [Drosophila miranda]
>gi|1037044352|ref|XP_017140776.1| PREDICTED: signal recognition particle subunit SRP72 [Drosop
>gi|1037044396|ref|XP_017140779.1| PREDICTED: tRNA pseudouridine synthase A, mitochondrial iso
>gi|1037044408|ref|XP_017140780.1| PREDICTED: tRNA pseudouridine synthase A, mitochondrial iso
>gi|1037044423|ref|XP_017140782.1| PREDICTED: tRNA pseudouridine synthase A, mitochondrial iso
>gi|1037044239|ref|XP_017140769.1| PREDICTED: transcription intermediary factor 1-alpha isoform
>gi|1037044255|ref|XP_017140770.1| PREDICTED: transcription intermediary factor 1-alpha isoform
>gi|1037044286|ref|XP_017140772.1| PREDICTED: transcription intermediary factor 1-alpha isoform
>gi|1037044302|ref|XP_017140773.1| PREDICTED: transcription intermediary factor 1-alpha isoform
>gi|1037044271|ref|XP_017140771.1| PREDICTED: transcription intermediary factor 1-alpha isoform
>gi|1037044318|ref|XP_017140774.1| PREDICTED: transcription intermediary factor 1-alpha isoform
>gi|1037044336|ref|XP_017140775.1| PREDICTED: meckelin [Drosophila miranda]
>gi|1037044437|ref|XP_017140783.1| PREDICTED: lipase 3-like [Drosophila miranda]
>gi|1037044366|ref|XP_017140777.1| PREDICTED: CCR4-NOT transcription complex subunit 10 [Drosop
>gi|1037044453|ref|XP_017140784.1| PREDICTED: sorting nexin-12 [Drosophila miranda]
>gi|1037044380|ref|XP_017140778.1| PREDICTED: V-type proton ATPase subunit B [Drosophila mirand
>gi|1037102915|ref|XP_017145025.1| PREDICTED: octopamine receptor beta-2R isoform X1 [Drosophil
>gi|1037102925|ref|XP_017145026.1| PREDICTED: octopamine receptor beta-2R isoform X2 [Drosophil
>gi|1037086201|ref|XP_017143672.1| PREDICTED: protein hugin [Drosophila miranda]
>gi|1037086142|ref|XP_017143667.1| PREDICTED: octopamine receptor beta-3R-like [Drosophila mira
>gi|1037086170|ref|XP_017143670.1| PREDICTED: uncharacterized protein LOC108156624 isoform X1
>gi|1037086183|ref|XP_017143671.1| PREDICTED: uncharacterized protein LOC108156624 isoform X2
>gi|1037086156|ref|XP_017143668.1| PREDICTED: glutamyl-tRNA(Gln) amidotransferase subunit A, m
>gi|1037086215|ref|XP_017143673.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomple
>gi|1037025596|ref|XP_017139624.1| PREDICTED: pyrroline-5-carboxylate reductase 1, mitochondri
>gi|1037025610|ref|XP_017139625.1| PREDICTED: pre-mRNA-splicing factor 18 [Drosophila miranda]
>gi|1037076980|ref|XP_017142993.1| PREDICTED: WD repeat-containing protein 24 [Drosophila mira
>gi|1037077045|ref|XP_017142998.1| PREDICTED: probable glucosamine 6-phosphate N-acetyltransfe
>gi|1037077027|ref|XP_017142997.1| PREDICTED: probable glucosamine 6-phosphate N-acetyltransfe
>gi|1037077012|ref|XP_017142995.1| PREDICTED: complex I intermediate-associated protein 30, mi
>gi|1037076996|ref|XP_017142994.1| PREDICTED: dehydrogenase/reductase SDR family protein 7-lik
>gi|1037075340|ref|XP_017142884.1| PREDICTED: uncharacterized protein LOC108156102 [Drosophila
>gi|1037075358|ref|XP_017142885.1| PREDICTED: uncharacterized protein LOC108156102 [Drosophila
>gi|1037075467|ref|XP_017142893.1| PREDICTED: uncharacterized protein LOC108156108 [Drosophila
>gi|1037075404|ref|XP_017142888.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1037075421|ref|XP_017142889.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1037075437|ref|XP_017142890.1| PREDICTED: N-alpha-acetyltransferase 40 [Drosophila miranda]
>gi|1037075452|ref|XP_017142891.1| PREDICTED: pterin-4-alpha-carbinolamine dehydratase [Drosoph
>gi|1037075483|ref|XP_017142894.1| PREDICTED: uncharacterized protein LOC108156109 [Drosophila
>gi|1037075372|ref|XP_017142886.1| PREDICTED: proline-rich protein 2 isoform X1 [Drosophila mi
>gi|1037075388|ref|XP_017142887.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1037104995|ref|XP_017145293.1| PREDICTED: general odorant-binding protein 99b [Drosophila
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>gi|1037105001|ref|XP_017145294.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037104973|ref|XP_017145289.1| PREDICTED: doublesex- and mab-3-related transcription factor
>gi|1037098375|ref|XP_017144609.1| PREDICTED: general odorant-binding protein 99a [Drosophila r
>gi|1037098351|ref|XP_017144607.1| PREDICTED: head-specific guanylate cyclase [Drosophila miran
>gi|1037098363|ref|XP_017144608.1| PREDICTED: lysoplasmalogenase-like protein TMEM86A [Drosophi
>gi|1037074751|ref|XP_017142842.1| PREDICTED: mitotic checkpoint protein BUB3 [Drosophila miran
>gi|1037074721|ref|XP_017142839.1| PREDICTED: pickpocket protein 19 isoform X3 [Drosophila mira
>gi|1037074707|ref|XP_017142838.1| PREDICTED: pickpocket protein 19 isoform X2 [Drosophila mira
>gi|1037074691|ref|XP_017142837.1| PREDICTED: pickpocket protein 19 isoform X1 [Drosophila mira
>gi|1037074767|ref|XP_017142843.1| PREDICTED: general odorant-binding protein 99a [Drosophila r
>gi|1037025625|ref|XP_017139626.1| PREDICTED: pickpocket protein 28, partial [Drosophila mirano
>gi|1037074675|ref|XP_017142836.1| PREDICTED: pickpocket protein 19 [Drosophila miranda]
>gi|1037074733|ref|XP_017142841.1| PREDICTED: pickpocket protein 19 [Drosophila miranda]
>gi|1037025643|ref|XP_017139627.1| PREDICTED: pickpocket protein 19-like [Drosophila miranda]
>gi|1037101124|ref|XP_017144852.1| PREDICTED: LOW QUALITY PROTEIN: dopamine receptor 2 [Drosoph
>gi|1037098217|ref|XP_017144593.1| PREDICTED: mitochondrial 2-oxoglutarate/malate carrier prote
>gi|1037080608|ref|XP_017143249.1| PREDICTED: LOW QUALITY PROTEIN: dynein assembly factor with
>gi|1037080594|ref|XP_017143248.1| PREDICTED: protein phosphatase 1B [Drosophila miranda]
>gi|1037099970|ref|XP_017144752.1| PREDICTED: LOW QUALITY PROTEIN: pre-mRNA-splicing factor 18-
>gi|1037093612|ref|XP_017144199.1| PREDICTED: protein TsetseEP [Drosophila miranda]
>gi|1037037047|ref|XP_017140303.1| PREDICTED: uncharacterized protein LOC108154514 [Drosophila
>gi|1037035429|ref|XP_017140213.1| PREDICTED: uncharacterized protein LOC108154452 [Drosophila
>gi|1037096731|ref|XP_017144451.1| PREDICTED: muscle segmentation homeobox [Drosophila miranda]
>gi|1037034577|ref|XP_017140168.1| PREDICTED: NADH-quinone oxidoreductase subunit B 2 [Drosophi
>gi|1037034595|ref|XP_017140169.1| PREDICTED: NADH-quinone oxidoreductase subunit B 2 [Drosophi
>gi|1037081277|ref|XP_017143300.1| PREDICTED: protein TRC8 homolog isoform X1 [Drosophila miran
>gi|1037081293|ref|XP_017143301.1| PREDICTED: protein TRC8 homolog isoform X1 [Drosophila miran
>gi|1037081305|ref|XP_017143302.1| PREDICTED: protein TRC8 homolog isoform X1 [Drosophila miran
>gi|1037081319|ref|XP_017143304.1| PREDICTED: protein TRC8 homolog isoform X2 [Drosophila miran
>gi|1037081332|ref|XP_017143305.1| PREDICTED: protein TRC8 homolog isoform X3 [Drosophila miran
>gi|1037081345|ref|XP_017143306.1| PREDICTED: protein TRC8 homolog isoform X4 [Drosophila miran
>gi|1037081389|ref|XP_017143309.1| PREDICTED: F-box/LRR-repeat protein 7 isoform X2 [Drosophila
>gi|1037081403|ref|XP_017143310.1| PREDICTED: F-box/LRR-repeat protein 7 isoform X2 [Drosophila
>gi|1037081359|ref|XP_017143307.1| PREDICTED: uncharacterized F-box/LRR-repeat protein C02F5.7
>gi|1037081373|ref|XP_017143308.1| PREDICTED: uncharacterized F-box/LRR-repeat protein C02F5.7
>gi|1037098237|ref|XP_017144595.1| PREDICTED: uncharacterized protein LOC108157183 [Drosophila
>gi|1037098247|ref|XP_017144596.1| PREDICTED: uncharacterized protein LOC108157183 [Drosophila
>gi|1037098259|ref|XP_017144598.1| PREDICTED: uncharacterized protein LOC108157183 [Drosophila
>gi|1037098269|ref|XP_017144599.1| PREDICTED: uncharacterized protein LOC108157183 [Drosophila
>gi|1037098281|ref|XP_017144600.1| PREDICTED: uncharacterized protein LOC108157183 [Drosophila
>gi|1037067073|ref|XP_017142311.1| PREDICTED: signal peptidase complex subunit 1 [Drosophila m
>gi|1037067053|ref|XP_017142310.1| PREDICTED: DTW domain-containing protein 1 [Drosophila miran
>gi|1037067037|ref|XP_017142309.1| PREDICTED: uncharacterized protein C2orf42 homolog [Drosophi
>gi|1037066975|ref|XP_017142304.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X4 [Dros
>gi|1037066945|ref|XP_017142302.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X2 [Dros
>gi|1037066931|ref|XP_017142301.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X1 [Dros
>gi|1037066991|ref|XP_017142305.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X5 [Dros
>gi|1037066961|ref|XP_017142303.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X3 [Dros
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>gi|1037067021|ref|XP_017142308.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X7 [Dros
>gi|1037067005|ref|XP_017142307.1| PREDICTED: tyrosine-protein phosphatase 99A isoform X6 [Dros
>gi|1037025659|ref|XP_017139628.1| PREDICTED: protein Diedel-like [Drosophila miranda]
>gi|1037034778|ref|XP_017140179.1| PREDICTED: protein Diedel-like [Drosophila miranda]
>gi|1037089830|ref|XP_017143923.1| PREDICTED: calnexin [Drosophila miranda]
>gi|1037025677|ref|XP_017139629.1| PREDICTED: uncharacterized protein LOC108154049 [Drosophila
>gi|1037025695|ref|XP_017139630.1| PREDICTED: uncharacterized protein LOC108154050 [Drosophila
>gi|1037025712|ref|XP_017139631.1| PREDICTED: uncharacterized protein LOC108154051 [Drosophila
>gi|1037025728|ref|XP_017139633.1| PREDICTED: uncharacterized protein LOC108154052 [Drosophila
>gi|1037025745|ref|XP_017139634.1| PREDICTED: uncharacterized protein LOC108154053 [Drosophila
>gi|1037091954|ref|XP_017144087.1| PREDICTED: M-phase inducer phosphatase [Drosophila miranda]
>gi|1037025761|ref|XP_017139635.1| PREDICTED: uncharacterized protein LOC108154054 [Drosophila
>gi|1037100326|ref|XP_017144773.1| PREDICTED: aminopeptidase N-like [Drosophila miranda]
>gi|1037054463|ref|XP_017141464.1| PREDICTED: aminopeptidase N-like [Drosophila miranda]
>gi|1037054491|ref|XP_017141465.1| PREDICTED: aminopeptidase N-like [Drosophila miranda]
>gi|1037054445|ref|XP_017141463.1| PREDICTED: uncharacterized protein LOC108155268 isoform X3
>gi|1037054431|ref|XP_017141462.1| PREDICTED: uncharacterized protein LOC108155268 isoform X2
>gi|1037054398|ref|XP_017141460.1| PREDICTED: uncharacterized protein LOC108155268 isoform X1
>gi|1037054415|ref|XP_017141461.1| PREDICTED: uncharacterized protein LOC108155268 isoform X1
>gi|1037054523|ref|XP_017141467.1| PREDICTED: elongation factor 1-gamma [Drosophila miranda]
>gi|1037054555|ref|XP_017141470.1| PREDICTED: CDGSH iron-sulfur domain-containing protein 2 hom
>gi|1037054507|ref|XP_017141466.1| PREDICTED: bromodomain-containing protein 8 [Drosophila mira
>gi|1037054539|ref|XP_017141469.1| PREDICTED: uncharacterized protein LOC108155273 [Drosophila
>gi|1037104921|ref|XP_017145281.1| PREDICTED: cytochrome c1, heme protein, mitochondrial [Drosc
>gi|1037090099|ref|XP_017143945.1| PREDICTED: guanosine-3',5'-bis(diphosphate) 3'-pyrophosphohy
>gi|1037090085|ref|XP_017143943.1| PREDICTED: probable phosphoserine aminotransferase [Drosophi
>gi|1037090113|ref|XP_017143946.1| PREDICTED: putative defense protein 3 [Drosophila miranda]
>gi|1037090057|ref|XP_017143941.1| PREDICTED: uncharacterized protein LOC108156780 isoform X2
>gi|1037090071|ref|XP_017143942.1| PREDICTED: uncharacterized protein LOC108156780 isoform X3
>gi|1037090043|ref|XP_017143940.1| PREDICTED: uncharacterized protein LOC108156780 isoform X1
>gi|1037090029|ref|XP_017143939.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037081832|ref|XP_017143346.1| PREDICTED: multidrug resistance-associated protein 4 [Drosop
>gi|1037042660|ref|XP_017140667.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1037042616|ref|XP_017140665.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1037042674|ref|XP_017140668.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1037042739|ref|XP_017140672.1| PREDICTED: alkaline phosphatase [Drosophila miranda]
>gi|1037042690|ref|XP_017140669.1| PREDICTED: putative tyrosine-protein kinase Wsck [Drosophila
>gi|1037042755|ref|XP_017140673.1| PREDICTED: atlastin [Drosophila miranda]
>gi|1037042796|ref|XP_017140676.1| PREDICTED: metallothionein-2 [Drosophila miranda]
>gi|1037042600|ref|XP_017140664.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 34 isoform
>gi|1037042584|ref|XP_017140663.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 34 isoform
>gi|1037042569|ref|XP_017140662.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 34 isoform
>gi|1037042705|ref|XP_017140670.1| PREDICTED: set1/Ash2 histone methyltransferase complex subur
>gi|1037042719|ref|XP_017140671.1| PREDICTED: set1/Ash2 histone methyltransferase complex subur
>gi|1037042780|ref|XP_017140675.1| PREDICTED: myb-like protein P [Drosophila miranda]
>gi|1037081969|ref|XP_017143357.1| PREDICTED: CLK4-associating serine/arginine rich protein isc
>gi|1037081983|ref|XP_017143358.1| PREDICTED: CLK4-associating serine/arginine rich protein isc
>gi|1037082034|ref|XP_017143360.1| PREDICTED: bolA-like protein DDB_G0274169 [Drosophila mirand
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>gi|1037082022|ref|XP_017143359.1| PREDICTED: protoporphyrinogen oxidase [Drosophila miranda]
>gi|1037086698|ref|XP_017143709.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086299|ref|XP_017143679.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086399|ref|XP_017143686.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086515|ref|XP_017143694.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086557|ref|XP_017143697.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086656|ref|XP_017143706.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086586|ref|XP_017143699.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086271|ref|XP_017143677.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086447|ref|XP_017143689.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086598|ref|XP_017143702.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086355|ref|XP_017143683.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086503|ref|XP_017143693.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086544|ref|XP_017143696.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086626|ref|XP_017143704.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086460|ref|XP_017143690.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086530|ref|XP_017143695.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086612|ref|XP_017143703.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086229|ref|XP_017143674.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086285|ref|XP_017143678.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086431|ref|XP_017143688.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086313|ref|XP_017143680.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086489|ref|XP_017143692.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086572|ref|XP_017143698.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086726|ref|XP_017143711.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086712|ref|XP_017143710.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086738|ref|XP_017143713.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086341|ref|XP_017143682.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086473|ref|XP_017143691.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086684|ref|XP_017143708.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086257|ref|XP_017143676.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086327|ref|XP_017143681.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086670|ref|XP_017143707.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086385|ref|XP_017143685.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086642|ref|XP_017143705.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086243|ref|XP_017143675.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086417|ref|XP_017143687.1| PREDICTED: calcium-activated potassium channel slowpoke iso
>gi|1037086751|ref|XP_017143714.1| PREDICTED: short coiled-coil protein B-like [Drosophila mira
>gi|1037033851|ref|XP_017140122.1| PREDICTED: putative uncharacterized protein DDB_G0271606 [D
>gi|1037098757|ref|XP_017144644.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu
>gi|1037098745|ref|XP_017144643.1| PREDICTED: methionine aminopeptidase 1 [Drosophila miranda]
>gi|1037066641|ref|XP_017142281.1| PREDICTED: dorsal-ventral patterning protein tolloid isofor
>gi|1037066655|ref|XP_017142282.1| PREDICTED: dorsal-ventral patterning protein tolloid isofor
>gi|1037066668|ref|XP_017142283.1| PREDICTED: dorsal-ventral patterning protein tolloid isofor
>gi|1037066685|ref|XP_017142284.1| PREDICTED: dorsal-ventral patterning protein tolloid [Drosop
>gi|1037071374|ref|XP_017142609.1| PREDICTED: protein abnormal spindle [Drosophila miranda]
>gi|1037071405|ref|XP_017142611.1| PREDICTED: allatostatin-A isoform X2 [Drosophila miranda]
>gi|1037071390|ref|XP_017142610.1| PREDICTED: allatostatin-A isoform X1 [Drosophila miranda]
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>gi|1037103740|ref|XP_017145121.1| PREDICTED: uncharacterized protein LOC108157529 [Drosophila
>gi|1037094566|ref|XP_017144271.1| PREDICTED: uncharacterized protein LOC108156990 [Drosophila
>gi|1037094578|ref|XP_017144272.1| PREDICTED: solute carrier family 25 member 44 [Drosophila mi
>gi|1037094592|ref|XP_017144273.1| PREDICTED: solute carrier family 25 member 44 [Drosophila mi
>gi|1037094606|ref|XP_017144274.1| PREDICTED: solute carrier family 25 member 44 [Drosophila mi
>gi|1037042275|ref|XP_017140642.1| PREDICTED: uncharacterized protein LOC108154770 [Drosophila
>gi|1037025779|ref|XP_017139636.1| PREDICTED: V-type proton ATPase subunit C [Drosophila mirand
>gi|1037042308|ref|XP_017140645.1| PREDICTED: uncharacterized protein LOC108154772 [Drosophila
>gi|1037042138|ref|XP_017140634.1| PREDICTED: protein polybromo-1 [Drosophila miranda]
>gi|1037042259|ref|XP_017140641.1| PREDICTED: protein LMBR1L isoform X2 [Drosophila miranda]
>gi|1037042228|ref|XP_017140640.1| PREDICTED: protein LMBR1L isoform X1 [Drosophila miranda]
>gi|1037042324|ref|XP_017140646.1| PREDICTED: tetratricopeptide repeat protein 12 isoform X1 [D
>gi|1037042342|ref|XP_017140647.1| PREDICTED: tetratricopeptide repeat protein 12 isoform X2 [D
>gi|1037042374|ref|XP_017140649.1| PREDICTED: complex III assembly factor LYRM7 [Drosophila mi
>gi|1037042214|ref|XP_017140639.1| PREDICTED: peptidyl-prolyl cis-trans isomerase-like 4 [Droso
>gi|1037042152|ref|XP_017140635.1| PREDICTED: SAFB-like transcription modulator isoform X1 [Dro
>gi|1037042168|ref|XP_017140636.1| PREDICTED: SAFB-like transcription modulator isoform X2 [Dro
>gi|1037042390|ref|XP_017140650.1| PREDICTED: serine/threonine-protein kinase Nek8 [Drosophila
>gi|1037042358|ref|XP_017140648.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1037042290|ref|XP_017140644.1| PREDICTED: macrophage erythroblast attacher [Drosophila mira
>gi|1037042199|ref|XP_017140638.1| PREDICTED: endothelin-converting enzyme 2 [Drosophila mirand
>gi|1037042184|ref|XP_017140637.1| PREDICTED: phosphate-regulating neutral endopeptidase [Droso
>gi|1037088597|ref|XP_017143825.1| PREDICTED: 40S ribosomal protein S3 [Drosophila miranda]
>gi|1037088583|ref|XP_017143824.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1037106394|ref|XP_017145490.1| PREDICTED: ATP-binding cassette sub-family A member 3-like
>gi|1037106402|ref|XP_017145491.1| PREDICTED: ATP-binding cassette sub-family A member 3-like
>gi|1037073189|ref|XP_017142735.1| PREDICTED: uncharacterized protein LOC108156017 [Drosophila
>gi|1037073205|ref|XP_017142736.1| PREDICTED: uncharacterized protein LOC108156017 [Drosophila
>gi|1037025797|ref|XP_017139637.1| PREDICTED: trypsin alpha-3 [Drosophila miranda]
>gi|1037073080|ref|XP_017142727.1| PREDICTED: uncharacterized protein LOC108156014 [Drosophila
>gi|1037073094|ref|XP_017142728.1| PREDICTED: uncharacterized protein LOC108156015 [Drosophila
>gi|1037073221|ref|XP_017142737.1| PREDICTED: triosephosphate isomerase isoform X1 [Drosophila
>gi|1037073237|ref|XP_017142738.1| PREDICTED: triosephosphate isomerase isoform X2 [Drosophila
>gi|1037073174|ref|XP_017142734.1| PREDICTED: uncharacterized protein LOC108156016 isoform X4
>gi|1037073142|ref|XP_017142732.1| PREDICTED: G-protein coupled receptor moody isoform X2 [Dros
>gi|1037073110|ref|XP_017142730.1| PREDICTED: D(1C) dopamine receptor isoform X1 [Drosophila mi
>gi|1037073126|ref|XP_017142731.1| PREDICTED: D(1C) dopamine receptor isoform X1 [Drosophila mi
>gi|1037073158|ref|XP_017142733.1| PREDICTED: D(1C) dopamine receptor isoform X3 [Drosophila mi
>gi|1037105406|ref|XP_017145353.1| PREDICTED: uncharacterized protein LOC108157694 [Drosophila
>gi|1037105398|ref|XP_017145352.1| PREDICTED: uncharacterized protein LOC108157693 [Drosophila
>gi|1037100626|ref|XP_017144800.1| PREDICTED: uncharacterized protein LOC108157306 isoform X2
>gi|1037100616|ref|XP_017144799.1| PREDICTED: uncharacterized protein LOC108157306 isoform X1
>gi|1037100606|ref|XP_017144798.1| PREDICTED: uncharacterized protein LOC108157305 [Drosophila
>gi|1037102323|ref|XP_017144965.1| PREDICTED: myosin regulatory light chain 2 [Drosophila miran
>gi|1037089400|ref|XP_017143888.1| PREDICTED: proline synthase co-transcribed bacterial homolog
>gi|1037089386|ref|XP_017143887.1| PREDICTED: cell cycle checkpoint protein RAD17 [Drosophila m
>gi|1037025813|ref|XP_017139638.1| PREDICTED: serine/arginine repetitive matrix protein 1 [Dros
>gi|1037103464|ref|XP_017145085.1| PREDICTED: trafficking protein particle complex subunit 1 [
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>gi|1037070129|ref|XP_017142525.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1037070145|ref|XP_017142526.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1037070099|ref|XP_017142523.1| PREDICTED: stearoyl-CoA desaturase 5 [Drosophila miranda]
>gi|1037070115|ref|XP_017142524.1| PREDICTED: stearoyl-CoA desaturase 5 [Drosophila miranda]
>gi|1037070160|ref|XP_017142527.1| PREDICTED: 40S ribosomal protein S7 [Drosophila miranda]
>gi|1037070188|ref|XP_017142529.1| PREDICTED: cecropin-2 [Drosophila miranda]
>gi|1037070174|ref|XP_017142528.1| PREDICTED: cecropin-2 [Drosophila miranda]
>gi|1037070204|ref|XP_017142531.1| PREDICTED: cecropin-A2-like [Drosophila miranda]
>gi|1037032188|ref|XP_017140021.1| PREDICTED: cecropin-C-like [Drosophila miranda]
>gi|1037032171|ref|XP_017140020.1| PREDICTED: cecropin-C-like [Drosophila miranda]
>gi|1037034678|ref|XP_017140175.1| PREDICTED: cecropin-C-like [Drosophila miranda]
>gi|1037101649|ref|XP_017144894.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1037101639|ref|XP_017144893.1| PREDICTED: serine protease easter isoform X2 [Drosophila mir
>gi|1037101629|ref|XP_017144892.1| PREDICTED: serine protease easter isoform X1 [Drosophila mir
>gi|1037101671|ref|XP_017144897.1| PREDICTED: uncharacterized protein LOC108157378 [Drosophila
>gi|1037101661|ref|XP_017144896.1| PREDICTED: circumsporozoite protein [Drosophila miranda]
>gi|1037101683|ref|XP_017144898.1| PREDICTED: uncharacterized protein LOC108157379 [Drosophila
>gi|1037079044|ref|XP_017143142.1| PREDICTED: LOW QUALITY PROTEIN: headcase protein [Drosophila
>gi|1037082638|ref|XP_017143404.1| PREDICTED: ferritin subunit isoform X2 [Drosophila miranda]
>gi|1037082624|ref|XP_017143403.1| PREDICTED: ferritin subunit isoform X1 [Drosophila miranda]
>gi|1037082651|ref|XP_017143405.1| PREDICTED: ferritin heavy chain [Drosophila miranda]
>gi|1037099791|ref|XP_017144733.1| PREDICTED: uncharacterized protein LOC108157263 [Drosophila
>gi|1037099746|ref|XP_017144730.1| PREDICTED: thyroid transcription factor 1-associated protei
>gi|1037099767|ref|XP_017144731.1| PREDICTED: thyroid transcription factor 1-associated protei
>gi|1037099779|ref|XP_017144732.1| PREDICTED: thyroid transcription factor 1-associated protei
>gi|1037099696|ref|XP_017144725.1| PREDICTED: 28S ribosomal protein S18c, mitochondrial [Drosop
>gi|1037099674|ref|XP_017144723.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037099684|ref|XP_017144724.1| PREDICTED: telomere-binding protein cav-like [Drosophila mi
>gi|1037094444|ref|XP_017144261.1| PREDICTED: RING finger protein 37 [Drosophila miranda]
>gi|1037094458|ref|XP_017144262.1| PREDICTED: RING finger protein 37 [Drosophila miranda]
>gi|1037104316|ref|XP_017145197.1| PREDICTED: sphingomyelin phosphodiesterase [Drosophila mira
>gi|1037103937|ref|XP_017145147.1| PREDICTED: sphingomyelin phosphodiesterase [Drosophila mira
>gi|1037103945|ref|XP_017145148.1| PREDICTED: corticotropin-releasing factor-binding protein [
>gi|1037025831|ref|XP_017139639.1| PREDICTED: uncharacterized protein LOC108154059 [Drosophila
>gi|1037087842|ref|XP_017143767.1| PREDICTED: calcium-binding mitochondrial carrier protein Ara
>gi|1037087829|ref|XP_017143766.1| PREDICTED: calcium-binding mitochondrial carrier protein Ara
>gi|1037087856|ref|XP_017143768.1| PREDICTED: STAM-binding protein-like A [Drosophila miranda]
>gi|1037101281|ref|XP_017144864.1| PREDICTED: neutral ceramidase [Drosophila miranda]
>gi|1037101291|ref|XP_017144865.1| PREDICTED: neutral ceramidase [Drosophila miranda]
>gi|1037101302|ref|XP_017144866.1| PREDICTED: neutral ceramidase [Drosophila miranda]
>gi|1037069313|ref|XP_017142468.1| PREDICTED: uncharacterized protein LOC108155861 [Drosophila
>gi|1037069329|ref|XP_017142469.1| PREDICTED: uncharacterized protein LOC108155861 [Drosophila
>gi|1037069111|ref|XP_017142453.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1037069095|ref|XP_017142452.1| PREDICTED: putative uncharacterized protein DDB_G0271606 [Dr
>gi|1037069205|ref|XP_017142459.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1037069079|ref|XP_017142451.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1037069175|ref|XP_017142457.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 isoform X1
>gi|1037069189|ref|XP_017142458.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 isoform X2
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>gi|1037069219|ref|XP_017142460.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1037069143|ref|XP_017142455.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1037069361|ref|XP_017142471.1| PREDICTED: uncharacterized protein LOC108155863 [Drosophila
>gi|1037069345|ref|XP_017142470.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1037069159|ref|XP_017142456.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1037069235|ref|XP_017142461.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isoform
>gi|1037069267|ref|XP_017142465.1| PREDICTED: uncharacterized protein LOC108155860 isoform X3
>gi|1037069251|ref|XP_017142464.1| PREDICTED: uncharacterized protein LOC108155860 isoform X2
>gi|1037069283|ref|XP_017142466.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like isoform
>gi|1037069299|ref|XP_017142467.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like isoform
>gi|1037069127|ref|XP_017142454.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1037096767|ref|XP_017144454.1| PREDICTED: J domain-containing protein isoform X2 [Drosophil
>gi|1037096755|ref|XP_017144453.1| PREDICTED: J domain-containing protein isoform X1 [Drosophil
>gi|1037096779|ref|XP_017144455.1| PREDICTED: J domain-containing protein isoform X3 [Drosophil
>gi|1037067579|ref|XP_017142346.1| PREDICTED: tropomodulin isoform X7 [Drosophila miranda]
>gi|1037067615|ref|XP_017142348.1| PREDICTED: tropomodulin isoform X9 [Drosophila miranda]
>gi|1037067499|ref|XP_017142341.1| PREDICTED: tropomodulin isoform X2 [Drosophila miranda]
>gi|1037067547|ref|XP_017142344.1| PREDICTED: tropomodulin isoform X5 [Drosophila miranda]
>gi|1037067456|ref|XP_017142337.1| PREDICTED: tropomodulin isoform X1 [Drosophila miranda]
>gi|1037067469|ref|XP_017142338.1| PREDICTED: tropomodulin isoform X1 [Drosophila miranda]
>gi|1037067485|ref|XP_017142340.1| PREDICTED: tropomodulin isoform X1 [Drosophila miranda]
>gi|1037067563|ref|XP_017142345.1| PREDICTED: tropomodulin isoform X6 [Drosophila miranda]
>gi|1037067515|ref|XP_017142342.1| PREDICTED: tropomodulin isoform X3 [Drosophila miranda]
>gi|1037067645|ref|XP_017142350.1| PREDICTED: uncharacterized protein LOC108155791 [Drosophila
>gi|1037067599|ref|XP_017142347.1| PREDICTED: tropomodulin isoform X8 [Drosophila miranda]
>gi|1037067631|ref|XP_017142349.1| PREDICTED: tropomodulin isoform X10 [Drosophila miranda]
>gi|1037067529|ref|XP_017142343.1| PREDICTED: tropomodulin isoform X4 [Drosophila miranda]
>gi|1037067440|ref|XP_017142336.1| PREDICTED: sodium-independent sulfate anion transporter [Dr
>gi|1037072036|ref|XP_017142653.1| PREDICTED: uncharacterized protein LOC108155975 [Drosophila
>gi|1037071974|ref|XP_017142649.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037071990|ref|XP_017142650.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037072022|ref|XP_017142652.1| PREDICTED: uncharacterized protein LOC108155974 isoform X3
>gi|1037072006|ref|XP_017142651.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1037071957|ref|XP_017142648.1| PREDICTED: WD repeat-containing protein 44 [Drosophila mira
>gi|1037084860|ref|XP_017143574.1| PREDICTED: zinc transporter ZIP13 homolog [Drosophila miran
>gi|1037084874|ref|XP_017143577.1| PREDICTED: zinc transporter ZIP13 homolog [Drosophila miran
>gi|1037084888|ref|XP_017143578.1| PREDICTED: zinc transporter ZIP13 homolog [Drosophila miran
>gi|1037084902|ref|XP_017143579.1| PREDICTED: zinc transporter ZIP13 homolog [Drosophila miran
>gi|1037084916|ref|XP_017143580.1| PREDICTED: 40S ribosomal protein S8 [Drosophila miranda]
>gi|1037096096|ref|XP_017144394.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037099031|ref|XP_017144667.1| PREDICTED: tudor domain-containing protein 5-like [Drosophil
>gi|1037073824|ref|XP_017142779.1| PREDICTED: KAT8 regulatory NSL complex subunit 2 [Drosophila
>gi|1037073794|ref|XP_017142777.1| PREDICTED: protocadherin-15 [Drosophila miranda]
>gi|1037073809|ref|XP_017142778.1| PREDICTED: protocadherin-15 [Drosophila miranda]
>gi|1037045604|ref|XP_017140863.1| PREDICTED: protein similar [Drosophila miranda]
>gi|1037045667|ref|XP_017140866.1| PREDICTED: mitochondrial intermembrane space import and asse
>gi|1037045716|ref|XP_017140869.1| PREDICTED: uncharacterized protein LOC108154921 [Drosophila
>gi|1037045685|ref|XP_017140867.1| PREDICTED: uncharacterized protein LOC108154919 [Drosophila
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>gi|1037045701|ref|XP_017140868.1| PREDICTED: uncharacterized protein LOC108154919 [Drosophila
>gi|1037045651|ref|XP_017140865.1| PREDICTED: DNA repair protein Rad51 homolog [Drosophila mira
>gi|1037045620|ref|XP_017140864.1| PREDICTED: PC4 and SFRS1-interacting protein [Drosophila mi
>gi|1037081939|ref|XP_017143354.1| PREDICTED: dual specificity protein phosphatase 18 [Drosoph:
>gi|1037081928|ref|XP_017143353.1| PREDICTED: solute carrier family 25 member 51 [Drosophila mi
>gi|1037081955|ref|XP_017143355.1| PREDICTED: 60S ribosomal protein L32 [Drosophila miranda]
>gi|1037081913|ref|XP_017143352.1| PREDICTED: serendipity locus protein delta [Drosophila miran
>gi|1037080846|ref|XP_017143268.1| PREDICTED: serendipity locus protein alpha [Drosophila miran
>gi|1037080860|ref|XP_017143269.1| PREDICTED: serendipity locus protein beta [Drosophila miranc
>gi|1037080872|ref|XP_017143270.1| PREDICTED: sex-regulated protein janus-A [Drosophila miranda
>gi|1037080886|ref|XP_017143271.1| PREDICTED: sex-regulated protein janus-B [Drosophila miranda
>gi|1037097205|ref|XP_017144494.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1037081081|ref|XP_017143286.1| PREDICTED: axin isoform X2 [Drosophila miranda]
>gi|1037081051|ref|XP_017143284.1| PREDICTED: axin isoform X1 [Drosophila miranda]
>gi|1037081067|ref|XP_017143285.1| PREDICTED: axin isoform X1 [Drosophila miranda]
>gi|1037081094|ref|XP_017143287.1| PREDICTED: axin isoform X3 [Drosophila miranda]
>gi|1037057432|ref|XP_017141661.1| PREDICTED: alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylgl
>gi|1037057458|ref|XP_017141663.1| PREDICTED: alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylgl
>gi|1037057489|ref|XP_017141665.1| PREDICTED: uncharacterized protein LOC108155391 isoform X5
>gi|1037057473|ref|XP_017141664.1| PREDICTED: alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylgl
>gi|1037057446|ref|XP_017141662.1| PREDICTED: alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylgl
>gi|1037057539|ref|XP_017141668.1| PREDICTED: kinesin-like protein subito [Drosophila miranda]
>gi|1037057555|ref|XP_017141669.1| PREDICTED: 4-hydroxybutyrate coenzyme A transferase [Drosoph
>gi|1037057571|ref|XP_017141671.1| PREDICTED: nucleoplasmin-like protein [Drosophila miranda]
>gi|1037057521|ref|XP_017141667.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1037057505|ref|XP_017141666.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1037055199|ref|XP_017141516.1| PREDICTED: nucleoplasmin-like protein [Drosophila miranda]
>gi|1037055060|ref|XP_017141506.1| PREDICTED: ADAM 17-like protease isoform X1 [Drosophila mira
>gi|1037055075|ref|XP_017141507.1| PREDICTED: ADAM 17-like protease isoform X2 [Drosophila mira
>gi|1037055090|ref|XP_017141508.1| PREDICTED: ADAM 17-like protease isoform X3 [Drosophila mira
>gi|1037055154|ref|XP_017141513.1| PREDICTED: uncharacterized protein LOC108155307 isoform X1
>gi|1037055169|ref|XP_017141514.1| PREDICTED: uncharacterized protein LOC108155307 isoform X2
>gi|1037055138|ref|XP_017141512.1| PREDICTED: uncharacterized protein LOC108155306 [Drosophila
>gi|1037055184|ref|XP_017141515.1| PREDICTED: coiled-coil domain-containing protein 12 [Drosoph
>gi|1037055107|ref|XP_017141509.1| PREDICTED: spindle assembly abnormal protein 6 homolog [Dros
>gi|1037025849|ref|XP_017139640.1| PREDICTED: glycine-rich RNA-binding protein 3, mitochondrial
>gi|1037055122|ref|XP_017141511.1| PREDICTED: prostatic acid phosphatase [Drosophila miranda]
>gi|1037055045|ref|XP_017141505.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1037103259|ref|XP_017145064.1| PREDICTED: tachykinin-like peptides receptor 99D isoform X1
>gi|1037103289|ref|XP_017145067.1| PREDICTED: tachykinin-like peptides receptor 99D isoform X4
>gi|1037103297|ref|XP_017145068.1| PREDICTED: tachykinin-like peptides receptor 99D isoform X5
>gi|1037103269|ref|XP_017145065.1| PREDICTED: tachykinin-like peptides receptor 99D isoform X2
>gi|1037103279|ref|XP_017145066.1| PREDICTED: tachykinin-like peptides receptor 99D isoform X3
>gi|1037103309|ref|XP_017145069.1| PREDICTED: tachykinin-like peptides receptor 99D isoform X6
>gi|1037047021|ref|XP_017140957.1| PREDICTED: vacuolar protein sorting-associated protein 13B
>gi|1037046969|ref|XP_017140953.1| PREDICTED: vacuolar protein sorting-associated protein 13B
>gi|1037046985|ref|XP_017140955.1| PREDICTED: vacuolar protein sorting-associated protein 13B
>gi|1037047003|ref|XP_017140956.1| PREDICTED: vacuolar protein sorting-associated protein 13B
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>gi|1037047037|ref|XP_017140958.1| PREDICTED: vacuolar protein sorting-associated protein 13B
>gi|1037047055|ref|XP_017140959.1| PREDICTED: translation initiation factor eIF-2B subunit alp
>gi|1037047100|ref|XP_017140962.1| PREDICTED: uncharacterized protein LOC108154960 [Drosophila
>gi|1037047116|ref|XP_017140965.1| PREDICTED: uncharacterized protein LOC108154961 [Drosophila
>gi|1037047084|ref|XP_017140961.1| PREDICTED: uncharacterized protein LOC108154959 [Drosophila
>gi|1037047069|ref|XP_017140960.1| PREDICTED: gametocyte-specific factor 1 homolog [Drosophila
>gi|1037084088|ref|XP_017143515.1| PREDICTED: conserved oligomeric Golgi complex subunit 7 [Dro
>gi|1037025867|ref|XP_017139641.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1037084102|ref|XP_017143516.1| PREDICTED: uncharacterized protein LOC108156523 [Drosophila
>gi|1037032280|ref|XP_017140026.1| PREDICTED: serine protease 3 [Drosophila miranda]
>gi|1037103480|ref|XP_017145087.1| PREDICTED: serine proteases 1/2 isoform X2 [Drosophila miran
>gi|1037103472|ref|XP_017145086.1| PREDICTED: serine proteases 1/2 isoform X1 [Drosophila miran
>gi|1037103488|ref|XP_017145088.1| PREDICTED: LOW QUALITY PROTEIN: serine proteases 1/2-like [I
>gi|1037103033|ref|XP_017145038.1| PREDICTED: transient receptor potential protein [Drosophila
>gi|1037103043|ref|XP_017145039.1| PREDICTED: cardio acceleratory peptide 2b [Drosophila mirand
>gi|1037104666|ref|XP_017145245.1| PREDICTED: uncharacterized protein LOC108157620 [Drosophila
>gi|1037037101|ref|XP_017140307.1| PREDICTED: endocuticle structural glycoprotein SgAbd-9 [Dros
>gi|1037064522|ref|XP_017142152.1| PREDICTED: sodium-independent sulfate anion transporter [Dro
>gi|1037064691|ref|XP_017142164.1| PREDICTED: phosphoribosyl pyrophosphate synthase-associated
>gi|1037064674|ref|XP_017142163.1| PREDICTED: phosphoribosyl pyrophosphate synthase-associated
>gi|1037064707|ref|XP_017142165.1| PREDICTED: phosphoribosyl pyrophosphate synthase-associated
>gi|1037064723|ref|XP_017142166.1| PREDICTED: phosphoribosyl pyrophosphate synthase-associated
>gi|1037064935|ref|XP_017142170.1| PREDICTED: cytosolic carboxypeptidase 6 [Drosophila miranda]
>gi|1037064627|ref|XP_017142160.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1037064643|ref|XP_017142161.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 isoform X1
>gi|1037064658|ref|XP_017142162.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 isoform X2
>gi|1037064552|ref|XP_017142154.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Drosop
>gi|1037064611|ref|XP_017142158.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Drosop
>gi|1037064596|ref|XP_017142157.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isoform
>gi|1037064568|ref|XP_017142155.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isoform
>gi|1037064581|ref|XP_017142156.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isoform
>gi|1037064536|ref|XP_017142153.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Drosop
>gi|1037064739|ref|XP_017142167.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prote
>gi|1037025883|ref|XP_017139643.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1037064753|ref|XP_017142168.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1037064767|ref|XP_017142169.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Drosop
>gi|1037106516|ref|XP_017145507.1| PREDICTED: uncharacterized protein LOC108157819 [Drosophila
>gi|1037105910|ref|XP_017145423.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Drosop
>gi|1037106786|ref|XP_017145545.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1037106792|ref|XP_017145546.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1037025904|ref|XP_017139644.1| PREDICTED: chromobox protein homolog 1-like [Drosophila mira
>gi|1037032706|ref|XP_017140054.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1037035788|ref|XP_017140229.1| PREDICTED: uncharacterized protein LOC108154469 [Drosophila
>gi|1037105079|ref|XP_017145305.1| PREDICTED: uncharacterized protein LOC108157659 [Drosophila
>gi|1037105073|ref|XP_017145304.1| PREDICTED: protein kinase DC1 [Drosophila miranda]
>gi|1037105049|ref|XP_017145301.1| PREDICTED: protein kinase DC1 isoform X4 [Drosophila miranda
>gi|1037105041|ref|XP_017145299.1| PREDICTED: protein kinase DC1 isoform X3 [Drosophila miranda
>gi|1037105031|ref|XP_017145298.1| PREDICTED: protein kinase DC1 isoform X2 [Drosophila miranda
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>gi|1037105023|ref|XP_017145297.1| PREDICTED: protein kinase DC1 isoform X1 [Drosophila miranda
>gi|1037105065|ref|XP_017145303.1| PREDICTED: protein kinase DC1 isoform X6 [Drosophila miranda
>gi|1037105057|ref|XP_017145302.1| PREDICTED: protein kinase DC1 isoform X5 [Drosophila miranda
>gi|1037073721|ref|XP_017142772.1| PREDICTED: uncharacterized protein LOC108156030 isoform X2
>gi|1037073705|ref|XP_017142771.1| PREDICTED: uncharacterized protein LOC108156030 isoform X1
>gi|1037073689|ref|XP_017142770.1| PREDICTED: protein no-on-transient A [Drosophila miranda]
>gi|1037073735|ref|XP_017142773.1| PREDICTED: uncharacterized protein LOC108156032 [Drosophila
>gi|1037073751|ref|XP_017142774.1| PREDICTED: MD-2-related lipid-recognition protein-like [Dros
>gi|1037073673|ref|XP_017142769.1| PREDICTED: zinc finger protein 1 [Drosophila miranda]
>gi|1037078140|ref|XP_017143080.1| PREDICTED: serine/threonine-protein kinase Warts [Drosophila
>gi|1037105316|ref|XP_017145339.1| PREDICTED: uncharacterized protein LOC108157684 [Drosophila
>gi|1037103367|ref|XP_017145076.1| PREDICTED: protein deglycase DJ-1 [Drosophila miranda]
>gi|1037079784|ref|XP_017143185.1| PREDICTED: SH3 domain-containing kinase-binding protein 1 [D
>gi|1037105765|ref|XP_017145404.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1037105771|ref|XP_017145405.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1037102866|ref|XP_017145019.1| PREDICTED: protein tailless [Drosophila miranda]
>gi|1037059971|ref|XP_017141840.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1037059878|ref|XP_017141834.1| PREDICTED: structural maintenance of chromosomes protein 5
>gi|1037059955|ref|XP_017141839.1| PREDICTED: uncharacterized protein LOC108155510 [Drosophila
>gi|1037059925|ref|XP_017141837.1| PREDICTED: uncharacterized protein LOC108155508 [Drosophila
>gi|1037059909|ref|XP_017141836.1| PREDICTED: LOW QUALITY PROTEIN: putative uncharacterized pr
>gi|1037059939|ref|XP_017141838.1| PREDICTED: uncharacterized protein LOC108155509 [Drosophila
>gi|1037059894|ref|XP_017141835.1| PREDICTED: uncharacterized protein LOC108155506 [Drosophila
>gi|1037031579|ref|XP_017139989.1| PREDICTED: cytochrome P450 4c3 [Drosophila miranda]
>gi|1037025924|ref|XP_017139645.1| PREDICTED: uncharacterized protein LOC108154065 [Drosophila
>gi|1037082102|ref|XP_017143366.1| PREDICTED: pituitary homeobox homolog Ptx1 isoform X1 [Droso
>gi|1037082116|ref|XP_017143367.1| PREDICTED: pituitary homeobox homolog Ptx1 isoform X2 [Droso
>gi|1037082213|ref|XP_017143375.1| PREDICTED: uncharacterized protein LOC108156431 [Drosophila
>gi|1037082227|ref|XP_017143376.1| PREDICTED: uncharacterized protein LOC108156432 [Drosophila
>gi|1037082185|ref|XP_017143373.1| PREDICTED: uncharacterized protein LOC108156430 [Drosophila
>gi|1037082199|ref|XP_017143374.1| PREDICTED: uncharacterized protein LOC108156430 [Drosophila
>gi|1037082048|ref|XP_017143361.1| PREDICTED: uncharacterized protein LOC108156423 [Drosophila
>gi|1037082144|ref|XP_017143369.1| PREDICTED: uncharacterized protein LOC108156427 [Drosophila
>gi|1037082130|ref|XP_017143368.1| PREDICTED: 5-hydroxytryptamine receptor 1 [Drosophila miranc
>gi|1037082158|ref|XP_017143370.1| PREDICTED: coiled-coil-helix-coiled-coil-helix domain-contai
>gi|1037082172|ref|XP_017143371.1| PREDICTED: uncharacterized protein LOC108156429 [Drosophila
>gi|1037025942|ref|XP_017139646.1| PREDICTED: uncharacterized protein LOC108154066 [Drosophila
>gi|1037025959|ref|XP_017139647.1| PREDICTED: uncharacterized protein LOC108154067 [Drosophila
>gi|1037082062|ref|XP_017143362.1| PREDICTED: serine/threonine-protein phosphatase 2B catalytic
>gi|1037082075|ref|XP_017143363.1| PREDICTED: serine/threonine-protein phosphatase 2B catalytic
>gi|1037082088|ref|XP_017143365.1| PREDICTED: serine/threonine-protein phosphatase 2B catalytic
>gi|1037022574|ref|XP_017139430.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037097109|ref|XP_017144485.1| PREDICTED: alkaline phosphatase 4 [Drosophila miranda]
>gi|1037097121|ref|XP_017144486.1| PREDICTED: MICOS complex subunit MIC19 [Drosophila miranda]
>gi|1037097133|ref|XP_017144487.1| PREDICTED: MICOS complex subunit MIC19 [Drosophila miranda]
>gi|1037080899|ref|XP_017143272.1| PREDICTED: discs overgrown protein kinase [Drosophila mirand
>gi|1037080913|ref|XP_017143273.1| PREDICTED: discs overgrown protein kinase [Drosophila mirand
>gi|1037080927|ref|XP_017143274.1| PREDICTED: discs overgrown protein kinase [Drosophila mirand
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>gi|1037080941|ref|XP_017143275.1| PREDICTED: discs overgrown protein kinase [Drosophila miran
>gi|1037080954|ref|XP_017143276.1| PREDICTED: discs overgrown protein kinase [Drosophila miran
>gi|1037105177|ref|XP_017145320.1| PREDICTED: transcription factor SOX-10 [Drosophila miranda]
>gi|1037105187|ref|XP_017145321.1| PREDICTED: uncharacterized protein LOC108157670 [Drosophila
>gi|1037103013|ref|XP_017145035.1| PREDICTED: serpin B3-like isoform X2 [Drosophila miranda]
>gi|1037103003|ref|XP_017145034.1| PREDICTED: leukocyte elastase inhibitor-like isoform X1 [Dro
>gi|1037025975|ref|XP_017139648.1| PREDICTED: uncharacterized protein LOC108154068 [Drosophila
>gi|1037034072|ref|XP_017140135.1| PREDICTED: uncharacterized protein LOC108154382 [Drosophila
>gi|1037037328|ref|XP_017140320.1| PREDICTED: proton-coupled folate transporter [Drosophila mi
>gi|1037105195|ref|XP_017145322.1| PREDICTED: adhesion G-protein coupled receptor G2 [Drosophil
>gi|1037037169|ref|XP_017140311.1| PREDICTED: uncharacterized protein LOC108154521 [Drosophila
>gi|1037105757|ref|XP_017145402.1| PREDICTED: adhesion G-protein coupled receptor G2 [Drosophil
>gi|1037039171|ref|XP_017140431.1| PREDICTED: guanylate cyclase soluble subunit beta-1 [Drosop
>gi|1037039263|ref|XP_017140435.1| PREDICTED: uncharacterized protein LOC108154624 [Drosophila
>gi|1037039039|ref|XP_017140424.1| PREDICTED: protein mesh isoform X3 [Drosophila miranda]
>gi|1037039025|ref|XP_017140423.1| PREDICTED: protein mesh isoform X2 [Drosophila miranda]
>gi|1037039007|ref|XP_017140422.1| PREDICTED: protein mesh isoform X1 [Drosophila miranda]
>gi|1037039280|ref|XP_017140437.1| PREDICTED: protein bottleneck [Drosophila miranda]
>gi|1037039105|ref|XP_017140428.1| PREDICTED: probable 2-oxoglutarate dehydrogenase E1 componen
>gi|1037039057|ref|XP_017140425.1| PREDICTED: chaoptin isoform X1 [Drosophila miranda]
>gi|1037039075|ref|XP_017140426.1| PREDICTED: chaoptin isoform X1 [Drosophila miranda]
>gi|1037039093|ref|XP_017140427.1| PREDICTED: chaoptin isoform X2 [Drosophila miranda]
>gi|1037039246|ref|XP_017140434.1| PREDICTED: protein zwilch [Drosophila miranda]
>gi|1037039298|ref|XP_017140438.1| PREDICTED: probable rRNA-processing protein EBP2 homolog [D
>gi|1037039312|ref|XP_017140439.1| PREDICTED: G patch domain-containing protein 4 [Drosophila
>gi|1037039328|ref|XP_017140440.1| PREDICTED: ATP synthase lipid-binding protein, mitochondrial
>gi|1037039342|ref|XP_017140441.1| PREDICTED: ATP synthase lipid-binding protein, mitochondrial
>gi|1037039230|ref|XP_017140433.1| PREDICTED: uncharacterized membrane protein DDB_G0293934 is
>gi|1037039215|ref|XP_017140432.1| PREDICTED: alpha-protein kinase 1 isoform X1 [Drosophila mi
>gi|1037039115|ref|XP_017140429.1| PREDICTED: vacuolar protein sorting-associated protein 16 h
>gi|1037054891|ref|XP_017141493.1| PREDICTED: tryptophan--tRNA ligase, cytoplasmic [Drosophila
>gi|1037054951|ref|XP_017141497.1| PREDICTED: uncharacterized protein LOC108155293 [Drosophila
>gi|1037054905|ref|XP_017141494.1| PREDICTED: renin receptor [Drosophila miranda]
>gi|1037054959|ref|XP_017141498.1| PREDICTED: uncharacterized protein LOC108155294 [Drosophila
>gi|1037054973|ref|XP_017141499.1| PREDICTED: uncharacterized protein LOC108155295 [Drosophila
>gi|1037054933|ref|XP_017141496.1| PREDICTED: mitochondrial RNA-splicing protein MRS4 [Drosoph
>gi|1037054984|ref|XP_017141501.1| PREDICTED: uncharacterized protein LOC108155296, partial [D
>gi|1037054875|ref|XP_017141492.1| PREDICTED: protein maelstrom 2 [Drosophila miranda]
>gi|1037054919|ref|XP_017141495.1| PREDICTED: estrogen sulfotransferase [Drosophila miranda]
>gi|1037056502|ref|XP_017141598.1| PREDICTED: uncharacterized protein LOC108155350 [Drosophila
>gi|1037056462|ref|XP_017141596.1| PREDICTED: phosphoinositide 3-kinase regulatory subunit 4 [
>gi|1037056486|ref|XP_017141597.1| PREDICTED: G2/mitotic-specific cyclin-B3 [Drosophila mirand
>gi|1037056598|ref|XP_017141604.1| PREDICTED: splicing factor 3A subunit 2 [Drosophila miranda]
>gi|1037056614|ref|XP_017141605.1| PREDICTED: metacaspase-1 [Drosophila miranda]
>gi|1037056520|ref|XP_017141599.1| PREDICTED: protein pygopus [Drosophila miranda]
>gi|1037056536|ref|XP_017141600.1| PREDICTED: protein pygopus [Drosophila miranda]
>gi|1037056552|ref|XP_017141601.1| PREDICTED: protein pygopus [Drosophila miranda]
>gi|1037056568|ref|XP_017141602.1| PREDICTED: protein pygopus [Drosophila miranda]
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>gi|1037056582|ref|XP_017141603.1| PREDICTED: protein pygopus [Drosophila miranda]
>gi|1037056630|ref|XP_017141606.1| PREDICTED: keratin-associated protein 19-2 [Drosophila mira
>gi|1037037860|ref|XP_017140350.1| PREDICTED: uncharacterized protein LOC108154560 [Drosophila
>gi|1037052344|ref|XP_017141318.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1037052438|ref|XP_017141324.1| PREDICTED: carbonic anhydrase 2 [Drosophila miranda]
>gi|1037052422|ref|XP_017141323.1| PREDICTED: histone deacetylase 11 [Drosophila miranda]
>gi|1037052313|ref|XP_017141316.1| PREDICTED: nicastrin [Drosophila miranda]
>gi|1037052454|ref|XP_017141326.1| PREDICTED: 60S ribosome subunit biogenesis protein NIP7 homo
>gi|1037052376|ref|XP_017141320.1| PREDICTED: mitochondrial coenzyme A transporter SLC25A42 [Dr
>gi|1037052390|ref|XP_017141321.1| PREDICTED: mitochondrial coenzyme A transporter SLC25A42 [Dr
>gi|1037052297|ref|XP_017141315.1| PREDICTED: LOW QUALITY PROTEIN: lon protease homolog, mitoch
>gi|1037052406|ref|XP_017141322.1| PREDICTED: uncharacterized protein LOC108155167 [Drosophila
>gi|1037052470|ref|XP_017141327.1| PREDICTED: uncharacterized protein LOC108155171 [Drosophila
>gi|1037052360|ref|XP_017141319.1| PREDICTED: succinate--hydroxymethylglutarate CoA-transferase
>gi|1037025993|ref|XP_017139650.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-3-like [Drosop
>gi|1037026010|ref|XP_017139651.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like, partia
>gi|1037021943|ref|XP_017150787.1| PREDICTED: troponin C [Drosophila miranda]
>gi|1037021961|ref|XP_017150788.1| PREDICTED: troponin C [Drosophila miranda]
>gi|1036993792|ref|XP_017149069.1| PREDICTED: troponin C isoform X1 [Drosophila miranda]
>gi|1036993806|ref|XP_017149070.1| PREDICTED: troponin C isoform X2 [Drosophila miranda]
>gi|1036993780|ref|XP_017149067.1| PREDICTED: laccase-5 [Drosophila miranda]
>gi|1036936108|ref|XP_017145572.1| PREDICTED: laccase-21-like, partial [Drosophila miranda]
>gi|1036936126|ref|XP_017145573.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037002313|ref|XP_017149591.1| PREDICTED: ataxin-8-like, partial [Drosophila miranda]
>gi|1036936142|ref|XP_017145574.1| PREDICTED: BTB/POZ domain-containing adapter for CUL3-mediat
>gi|1037003278|ref|XP_017149645.1| PREDICTED: probable protein phosphatase CG10417 [Drosophila
>gi|1036936160|ref|XP_017145575.1| PREDICTED: uncharacterized protein LOC108157943 [Drosophila
>gi|1036997515|ref|XP_017149299.1| PREDICTED: uncharacterized protein LOC108160063 isoform X1
>gi|1036997531|ref|XP_017149300.1| PREDICTED: uncharacterized protein LOC108160063 isoform X2
>gi|1036997497|ref|XP_017149298.1| PREDICTED: presequence protease, mitochondrial [Drosophila m
>gi|1037014063|ref|XP_017150296.1| PREDICTED: ARL14 effector protein [Drosophila miranda]
>gi|1037014019|ref|XP_017150293.1| PREDICTED: uncharacterized protein LOC108160663 isoform X1
>gi|1037014048|ref|XP_017150295.1| PREDICTED: uncharacterized protein LOC108160663 isoform X3
>gi|1037014036|ref|XP_017150294.1| PREDICTED: uncharacterized protein LOC108160663 isoform X2
>gi|1037007965|ref|XP_017149932.1| PREDICTED: zinc finger protein DPF3 [Drosophila miranda]
>gi|1037010764|ref|XP_017150093.1| PREDICTED: uncharacterized protein LOC108160540 [Drosophila
>gi|1037010798|ref|XP_017150095.1| PREDICTED: josephin-like protein isoform X2 [Drosophila mira
>gi|1037010782|ref|XP_017150094.1| PREDICTED: josephin-like protein isoform X1 [Drosophila mira
>gi|1037016781|ref|XP_017150456.1| PREDICTED: ribonuclease kappa [Drosophila miranda]
>gi|1036984049|ref|XP_017148478.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X
>gi|1036984031|ref|XP_017148477.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X
>gi|1036984102|ref|XP_017148481.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X5
>gi|1036984120|ref|XP_017148483.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X6
>gi|1036984141|ref|XP_017148484.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X6
>gi|1036984159|ref|XP_017148485.1| PREDICTED: rhoGEF domain-containing protein gxcJ isoform X7
>gi|1036984084|ref|XP_017148480.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X4
>gi|1036984068|ref|XP_017148479.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X3
>gi|1036984175|ref|XP_017148486.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X8
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>gi|1036935929|ref|XP_017145561.1| PREDICTED: uncharacterized protein LOC108157929 [Drosophila
>gi|1036984193|ref|XP_017148487.1| PREDICTED: guanine nucleotide exchange factor DBS isoform X9
>gi|1036997724|ref|XP_017149312.1| PREDICTED: craniofacial development protein 1 [Drosophila mi
>gi|1036997623|ref|XP_017149306.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036997635|ref|XP_017149307.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036997653|ref|XP_017149308.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036997671|ref|XP_017149309.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036997687|ref|XP_017149310.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036997707|ref|XP_017149311.1| PREDICTED: LOW QUALITY PROTEIN: cytosolic non-specific dipep
>gi|1037007656|ref|XP_017149913.1| PREDICTED: uncharacterized protein LOC108160422 [Drosophila
>gi|1037018816|ref|XP_017150590.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1037009879|ref|XP_017150044.1| PREDICTED: LOW QUALITY PROTEIN: cytosolic non-specific dipep
>gi|1036993260|ref|XP_017149036.1| PREDICTED: uncharacterized protein LOC108159914 [Drosophila
>gi|1037018026|ref|XP_017150536.1| PREDICTED: LOW QUALITY PROTEIN: BTB/POZ domain-containing ad
>gi|1036942154|ref|XP_017145951.1| PREDICTED: uncharacterized protein LOC108158230 [Drosophila
>gi|1037019099|ref|XP_017150610.1| PREDICTED: uncharacterized protein LOC108160858 [Drosophila
>gi|1037001889|ref|XP_017149564.1| PREDICTED: U4/U6.U5 small nuclear ribonucleoprotein 27 kDa p
>gi|1037001894|ref|XP_017149565.1| PREDICTED: U4/U6.U5 small nuclear ribonucleoprotein 27 kDa p
>gi|1037001872|ref|XP_017149563.1| PREDICTED: eukaryotic translation initiation factor 6 [Droso
>gi|1037009589|ref|XP_017150030.1| PREDICTED: thioredoxin-related transmembrane protein 1 isofo
>gi|1037009561|ref|XP_017150029.1| PREDICTED: thioredoxin-related transmembrane protein 1 isofo
>gi|1036979477|ref|XP_017148193.1| PREDICTED: synaptosomal-associated protein 29 isoform X2 [Dr
>gi|1036979494|ref|XP_017148194.1| PREDICTED: synaptosomal-associated protein 29 isoform X3 [Dr
>gi|1036979459|ref|XP_017148192.1| PREDICTED: synaptosomal-associated protein 29 isoform X1 [Dr
>gi|1036979443|ref|XP_017148191.1| PREDICTED: inactive peptidyl-prolyl cis-trans isomerase shut
>gi|1036979569|ref|XP_017148199.1| PREDICTED: inactive peptidyl-prolyl cis-trans isomerase shut
>gi|1036979512|ref|XP_017148195.1| PREDICTED: uncharacterized protein LOC108159433 isoform X1
>gi|1036979551|ref|XP_017148198.1| PREDICTED: dynein light chain roadblock-type 1-like isoform
>gi|1036979534|ref|XP_017148196.1| PREDICTED: dynein light chain roadblock-type 1-like isoform
>gi|1037015842|ref|XP_017150401.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1037015860|ref|XP_017150402.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1037015898|ref|XP_017150404.1| PREDICTED: uncharacterized protein LOC108160730 isoform X2
>gi|1037015880|ref|XP_017150403.1| PREDICTED: uncharacterized protein LOC108160730 isoform X1
>gi|1037015916|ref|XP_017150405.1| PREDICTED: uncharacterized protein LOC108160730 isoform X3
>gi|1037017974|ref|XP_017150532.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1037017990|ref|XP_017150533.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1036987099|ref|XP_017148667.1| PREDICTED: NTF2-related export protein [Drosophila miranda]
>gi|1036987027|ref|XP_017148662.1| PREDICTED: gastrulation defective protein 1 homolog [Drosoph
>gi|1036987015|ref|XP_017148661.1| PREDICTED: FACT complex subunit Ssrp1 [Drosophila miranda]
>gi|1036987081|ref|XP_017148665.1| PREDICTED: uncharacterized protein LOC108159682 [Drosophila
>gi|1036987045|ref|XP_017148663.1| PREDICTED: receptor expression-enhancing protein 2 [Drosophi
>gi|1036987063|ref|XP_017148664.1| PREDICTED: receptor expression-enhancing protein 2 [Drosophi
>gi|1037009759|ref|XP_017150038.1| PREDICTED: uncharacterized protein LOC108160508 isoform X2
>gi|1037009743|ref|XP_017150037.1| PREDICTED: uncharacterized protein LOC108160508 isoform X1
>gi|1037009809|ref|XP_017150039.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1037018556|ref|XP_017150572.1| PREDICTED: transmembrane protein 14 homolog [Drosophila mira
>gi|1036981253|ref|XP_017148307.1| PREDICTED: protein preli-like [Drosophila miranda]
>gi|1036981201|ref|XP_017148304.1| PREDICTED: chromatin-remodeling complex ATPase chain Iswi [D
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>gi|1036981221|ref|XP_017148305.1| PREDICTED: proton-coupled amino acid transporter 1 [Drosoph
>gi|1036981238|ref|XP_017148306.1| PREDICTED: proton-coupled amino acid transporter 1 [Drosoph
>gi|1037004350|ref|XP_017149707.1| PREDICTED: methylglutaconyl-CoA hydratase, mitochondrial [D
>gi|1037004313|ref|XP_017149706.1| PREDICTED: zinc transporter 9 [Drosophila miranda]
>gi|1037020057|ref|XP_017150670.1| PREDICTED: arylsulfatase B isoform X1 [Drosophila miranda]
>gi|1037020073|ref|XP_017150671.1| PREDICTED: arylsulfatase B isoform X2 [Drosophila miranda]
>gi|1037020091|ref|XP_017150672.1| PREDICTED: arylsulfatase B isoform X2 [Drosophila miranda]
>gi|1037020108|ref|XP_017150674.1| PREDICTED: uncharacterized protein LOC108160909 [Drosophila
>gi|1036982685|ref|XP_017148395.1| PREDICTED: cytochrome c oxidase assembly factor 4 homolog, 
>gi|1036982667|ref|XP_017148394.1| PREDICTED: peptidyl-tRNA hydrolase 2, mitochondrial [Drosop
>gi|1036982615|ref|XP_017148390.1| PREDICTED: GTPase Era, mitochondrial [Drosophila miranda]
>gi|1036982633|ref|XP_017148391.1| PREDICTED: small RNA 2'-O-methyltransferase [Drosophila mir
>gi|1036982649|ref|XP_017148393.1| PREDICTED: small RNA 2'-O-methyltransferase [Drosophila mir
>gi|1036982599|ref|XP_017148389.1| PREDICTED: uncharacterized protein LOC108159528 isoform X4 
>gi|1036982563|ref|XP_017148387.1| PREDICTED: uncharacterized protein LOC108159528 isoform X2 
>gi|1036982581|ref|XP_017148388.1| PREDICTED: uncharacterized protein LOC108159528 isoform X3 
>gi|1036982546|ref|XP_017148386.1| PREDICTED: uncharacterized protein LOC108159528 isoform X1 
>gi|1036976685|ref|XP_017148018.1| PREDICTED: dynein light chain 4, axonemal [Drosophila miran
>gi|1036976702|ref|XP_017148019.1| PREDICTED: dynein light chain 4, axonemal [Drosophila miran
>gi|1036976725|ref|XP_017148020.1| PREDICTED: dynein light chain 4, axonemal [Drosophila miran
>gi|1036976667|ref|XP_017148017.1| PREDICTED: pre-mRNA-processing-splicing factor 8 [Drosophil
>gi|1036992480|ref|XP_017148986.1| PREDICTED: uncharacterized protein LOC108159883 [Drosophila
>gi|1036992466|ref|XP_017148985.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036992433|ref|XP_017148982.1| PREDICTED: LOW QUALITY PROTEIN: ornithine decarboxylase ant
>gi|1036992451|ref|XP_017148983.1| PREDICTED: small nuclear ribonucleoprotein Sm D3 [Drosophil
>gi|1037007347|ref|XP_017149896.1| PREDICTED: endonuclease G, mitochondrial [Drosophila mirand
>gi|1037007333|ref|XP_017149895.1| PREDICTED: T-complex protein 1 subunit epsilon [Drosophila 
>gi|1036943809|ref|XP_017146053.1| PREDICTED: uncharacterized protein LOC108158324 [Drosophila
>gi|1037012915|ref|XP_017150226.1| PREDICTED: uncharacterized protein LOC108160622 isoform X1 
>gi|1037012933|ref|XP_017150227.1| PREDICTED: uncharacterized protein LOC108160622 isoform X1 
>gi|1037012951|ref|XP_017150228.1| PREDICTED: uncharacterized protein LOC108160622 isoform X1 
>gi|1037012971|ref|XP_017150230.1| PREDICTED: uncharacterized protein LOC108160622 isoform X2 
>gi|1037007126|ref|XP_017149882.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1037007108|ref|XP_017149881.1| PREDICTED: nuclear polyadenylated RNA-binding protein NAB2 
>gi|1037007090|ref|XP_017149880.1| PREDICTED: protein lunapark-B isoform X2 [Drosophila mirand
>gi|1037007071|ref|XP_017149879.1| PREDICTED: protein lunapark-B isoform X1 [Drosophila mirand
>gi|1036989131|ref|XP_017148787.1| PREDICTED: angiogenic factor with G patch and FHA domains 1
>gi|1036989147|ref|XP_017148788.1| PREDICTED: angiogenic factor with G patch and FHA domains 1
>gi|1036989165|ref|XP_017148789.1| PREDICTED: hexokinase type 2 [Drosophila miranda]
>gi|1036989112|ref|XP_017148786.1| PREDICTED: uncharacterized protein LOC108159752 isoform X2 
>gi|1036989094|ref|XP_017148785.1| PREDICTED: uncharacterized protein LOC108159752 isoform X1 
>gi|1036947419|ref|XP_017146268.1| PREDICTED: integrin alpha-PS3 isoform X1 [Drosophila mirand
>gi|1036947434|ref|XP_017146269.1| PREDICTED: integrin alpha-PS3 isoform X2 [Drosophila mirand
>gi|1036947207|ref|XP_017146255.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947225|ref|XP_017146256.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947245|ref|XP_017146257.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947170|ref|XP_017146253.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947189|ref|XP_017146254.1| PREDICTED: transient receptor potential cation channel trpm
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>gi|1036947277|ref|XP_017146259.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947261|ref|XP_017146258.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947152|ref|XP_017146251.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947135|ref|XP_017146250.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947119|ref|XP_017146249.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947312|ref|XP_017146261.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947294|ref|XP_017146260.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947348|ref|XP_017146264.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947332|ref|XP_017146262.1| PREDICTED: transient receptor potential cation channel trpm
>gi|1036947449|ref|XP_017146270.1| PREDICTED: NADH dehydrogenase [ubiquinone] flavoprotein 1, m
>gi|1036947467|ref|XP_017146271.1| PREDICTED: G patch domain and ankyrin repeat-containing prot
>gi|1036947483|ref|XP_017146272.1| PREDICTED: G patch domain and ankyrin repeat-containing prot
>gi|1036947365|ref|XP_017146265.1| PREDICTED: DNA repair protein complementing XP-C cells homol
>gi|1036947383|ref|XP_017146266.1| PREDICTED: DNA repair protein complementing XP-C cells homol
>gi|1036947401|ref|XP_017146267.1| PREDICTED: uncharacterized protein LOC108158474 [Drosophila
>gi|1036988250|ref|XP_017148733.1| PREDICTED: mitochondrial import receptor subunit TOM7 homolo
>gi|1036988200|ref|XP_017148730.1| PREDICTED: uncharacterized protein LOC108159720 [Drosophila
>gi|1036988210|ref|XP_017148731.1| PREDICTED: uncharacterized protein LOC108159720 [Drosophila
>gi|1036988226|ref|XP_017148732.1| PREDICTED: uncharacterized protein LOC108159722 [Drosophila
>gi|1036988169|ref|XP_017148729.1| PREDICTED: dymeclin [Drosophila miranda]
>gi|1036936178|ref|XP_017145577.1| PREDICTED: LOW QUALITY PROTEIN: odorant receptor 49a-like [D
>gi|1036943926|ref|XP_017146060.1| PREDICTED: kunitz-type serine protease inhibitor nigrescinin
>gi|1037015434|ref|XP_017150380.1| PREDICTED: uncharacterized protein LOC108160716 [Drosophila
>gi|1037021205|ref|XP_017150742.1| PREDICTED: uncharacterized protein LOC108160960 [Drosophila
>gi|1036936196|ref|XP_017145578.1| PREDICTED: gamma-butyrobetaine dioxygenase [Drosophila miran
>gi|1036944869|ref|XP_017146113.1| PREDICTED: tripartite motif-containing protein 75 isoform X
>gi|1036944905|ref|XP_017146115.1| PREDICTED: uncharacterized RING finger protein T02C1.1 isofo
>gi|1036944887|ref|XP_017146114.1| PREDICTED: ligand of Numb protein X 2 isoform X2 [Drosophila
>gi|1036942664|ref|XP_017145983.1| PREDICTED: dopamine N-acetyltransferase-like [Drosophila mir
>gi|1037019328|ref|XP_017150623.1| PREDICTED: uncharacterized protein LOC108160872 [Drosophila
>gi|1037019344|ref|XP_017150624.1| PREDICTED: uncharacterized protein LOC108160873 [Drosophila
>gi|1037019813|ref|XP_017150654.1| PREDICTED: calcitonin gene-related peptide type 1 receptor
>gi|1037019825|ref|XP_017150655.1| PREDICTED: calcitonin gene-related peptide type 1 receptor
>gi|1037019843|ref|XP_017150656.1| PREDICTED: calcitonin gene-related peptide type 1 receptor
>gi|1037020374|ref|XP_017150690.1| PREDICTED: uncharacterized protein LOC108160919 isoform X3
>gi|1037020345|ref|XP_017150688.1| PREDICTED: PDZ and LIM domain protein 7 isoform X1 [Drosophi
>gi|1037020357|ref|XP_017150689.1| PREDICTED: uncharacterized protein LOC108160919 isoform X2
>gi|1037019292|ref|XP_017150621.1| PREDICTED: transcriptional regulator ATRX [Drosophila mirand
>gi|1037015708|ref|XP_017150394.1| PREDICTED: uncharacterized protein LOC108160725 [Drosophila
>gi|1037015724|ref|XP_017150395.1| PREDICTED: muscle-specific protein 20 [Drosophila miranda]
>gi|1036996635|ref|XP_017149245.1| PREDICTED: glutathione S-transferase E14 isoform X2 [Drosoph
>gi|1036996617|ref|XP_017149244.1| PREDICTED: glutathione S-transferase E14 isoform X1 [Drosoph
>gi|1036996601|ref|XP_017149243.1| PREDICTED: RNA-binding protein 28 [Drosophila miranda]
>gi|1036996691|ref|XP_017149248.1| PREDICTED: uncharacterized protein LOC108160023 [Drosophila
>gi|1036996655|ref|XP_017149246.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036996673|ref|XP_017149247.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1037017257|ref|XP_017150487.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1037017241|ref|XP_017150486.1| PREDICTED: larval cuticle protein 4-like [Drosophila miranda
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>gi|1037017221|ref|XP_017150485.1| PREDICTED: isochorismatase domain-containing protein 1 [Dros
>gi|1037007450|ref|XP_017149901.1| PREDICTED: putative sodium-dependent multivitamin transporte
>gi|1036981630|ref|XP_017148331.1| PREDICTED: transcription factor Adf-1 isoform X1 [Drosophila
>gi|1036981646|ref|XP_017148332.1| PREDICTED: transcription factor Adf-1 isoform X2 [Drosophila
>gi|1036981675|ref|XP_017148334.1| PREDICTED: UPF0545 protein C22orf39 homolog [Drosophila mira
>gi|1036981657|ref|XP_017148333.1| PREDICTED: coenzyme Q-binding protein COQ10, mitochondrial
>gi|1036981592|ref|XP_017148329.1| PREDICTED: D-glucuronyl C5-epimerase [Drosophila miranda]
>gi|1036981611|ref|XP_017148330.1| PREDICTED: D-glucuronyl C5-epimerase [Drosophila miranda]
>gi|1036957094|ref|XP_017146857.1| PREDICTED: post-GPI attachment to proteins factor 3 [Drosop
>gi|1036957349|ref|XP_017146871.1| PREDICTED: peptidyl-prolyl cis-trans isomerase H [Drosophila
>gi|1036957056|ref|XP_017146855.1| PREDICTED: aromatic-L-amino-acid decarboxylase [Drosophila m
>gi|1036957112|ref|XP_017146858.1| PREDICTED: uncharacterized protein LOC108158757 isoform X1
>gi|1036957128|ref|XP_017146859.1| PREDICTED: uncharacterized protein LOC108158757 isoform X2
>gi|1036957023|ref|XP_017146853.1| PREDICTED: tyrosine decarboxylase isoform X1 [Drosophila mi
>gi|1036957039|ref|XP_017146854.1| PREDICTED: tyrosine decarboxylase isoform X2 [Drosophila mi
>gi|1036957237|ref|XP_017146863.1| PREDICTED: ribonuclease P protein subunit p25-like protein
>gi|1036957248|ref|XP_017146864.1| PREDICTED: ribonuclease P protein subunit p25-like protein
>gi|1036957260|ref|XP_017146865.1| PREDICTED: ribonuclease P protein subunit p25-like protein
>gi|1036957074|ref|XP_017146856.1| PREDICTED: FAD-dependent oxidoreductase domain-containing p
>gi|1036957180|ref|XP_017146860.1| PREDICTED: MOB kinase activator-like 4 [Drosophila miranda]
>gi|1036957194|ref|XP_017146862.1| PREDICTED: ras-related protein Rab-2A [Drosophila miranda]
>gi|1036957007|ref|XP_017146851.1| PREDICTED: monocarboxylate transporter 14 [Drosophila miran
>gi|1036957270|ref|XP_017146866.1| PREDICTED: uncharacterized protein LOC108158761 [Drosophila
>gi|1036957286|ref|XP_017146867.1| PREDICTED: uncharacterized protein LOC108158761 [Drosophila
>gi|1036957304|ref|XP_017146868.1| PREDICTED: uncharacterized protein LOC108158761 [Drosophila
>gi|1036957329|ref|XP_017146870.1| PREDICTED: uncharacterized protein LOC108158761 [Drosophila
>gi|1036942772|ref|XP_017145991.1| PREDICTED: zinc transporter ZIP1 [Drosophila miranda]
>gi|1036943755|ref|XP_017146050.1| PREDICTED: zinc transporter ZIP1 [Drosophila miranda]
>gi|1037009295|ref|XP_017150011.1| PREDICTED: uncharacterized protein LOC108160488 isoform X12
>gi|1037009228|ref|XP_017150007.1| PREDICTED: uncharacterized protein LOC108160488 isoform X8
>gi|1037009211|ref|XP_017150006.1| PREDICTED: uncharacterized protein LOC108160488 isoform X7
>gi|1037009178|ref|XP_017150004.1| PREDICTED: uncharacterized protein LOC108160488 isoform X5
>gi|1037009123|ref|XP_017150001.1| PREDICTED: uncharacterized protein LOC108160488 isoform X2
>gi|1037009194|ref|XP_017150005.1| PREDICTED: uncharacterized protein LOC108160488 isoform X6
>gi|1037009159|ref|XP_017150003.1| PREDICTED: uncharacterized protein LOC108160488 isoform X4
>gi|1037009134|ref|XP_017150002.1| PREDICTED: uncharacterized protein LOC108160488 isoform X3
>gi|1037009104|ref|XP_017150000.1| PREDICTED: uncharacterized protein LOC108160488 isoform X1
>gi|1037009313|ref|XP_017150012.1| PREDICTED: uncharacterized protein LOC108160488 isoform X13
>gi|1037009331|ref|XP_017150013.1| PREDICTED: uncharacterized protein LOC108160488 isoform X14
>gi|1037009262|ref|XP_017150009.1| PREDICTED: uncharacterized protein LOC108160488 isoform X10
>gi|1037009281|ref|XP_017150010.1| PREDICTED: uncharacterized protein LOC108160488 isoform X11
>gi|1037009245|ref|XP_017150008.1| PREDICTED: uncharacterized protein LOC108160488 isoform X9
>gi|1037000373|ref|XP_017149471.1| PREDICTED: putative homeodomain transcription factor [Drosop
>gi|1037000395|ref|XP_017149472.1| PREDICTED: putative homeodomain transcription factor [Drosop
>gi|1037012040|ref|XP_017150170.1| PREDICTED: probable methylcrotonoyl-CoA carboxylase beta cha
>gi|1036993868|ref|XP_017149074.1| PREDICTED: microtubule-associated protein RP/EB family membe
>gi|1036993830|ref|XP_017149071.1| PREDICTED: microtubule-associated protein RP/EB family membe
>gi|1036993842|ref|XP_017149072.1| PREDICTED: microtubule-associated protein RP/EB family membe
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>gi|1036993852|ref|XP_017149073.1| PREDICTED: microtubule-associated protein RP/EB family membe
>gi|1036993931|ref|XP_017149079.1| PREDICTED: CDGSH iron-sulfur domain-containing protein 3, m:
>gi|1036993901|ref|XP_017149077.1| PREDICTED: CDGSH iron-sulfur domain-containing protein 3, m:
>gi|1036993917|ref|XP_017149078.1| PREDICTED: CDGSH iron-sulfur domain-containing protein 3, m:
>gi|1036993883|ref|XP_017149075.1| PREDICTED: aldose reductase [Drosophila miranda]
>gi|1037006410|ref|XP_017149836.1| PREDICTED: rap guanine nucleotide exchange factor 4 isoform
>gi|1037006428|ref|XP_017149837.1| PREDICTED: rap guanine nucleotide exchange factor 4 isoform
>gi|1037006462|ref|XP_017149839.1| PREDICTED: rap guanine nucleotide exchange factor 4 isoform
>gi|1037006446|ref|XP_017149838.1| PREDICTED: rap guanine nucleotide exchange factor 4 isoform
>gi|1037006478|ref|XP_017149840.1| PREDICTED: rap guanine nucleotide exchange factor 4 isoform
>gi|1037006496|ref|XP_017149841.1| PREDICTED: cytochrome P450 6a2 [Drosophila miranda]
>gi|1036936212|ref|XP_017145579.1| PREDICTED: cytosol aminopeptidase-like [Drosophila miranda]
>gi|1037007796|ref|XP_017149921.1| PREDICTED: succinate dehydrogenase [ubiquinone] iron-sulfur
>gi|1037007813|ref|XP_017149922.1| PREDICTED: anaphase-promoting complex subunit 15A [Drosophil
>gi|1037007848|ref|XP_017149924.1| PREDICTED: ubiquitin-like protein 5 isoform X2 [Drosophila r
>gi|1037007831|ref|XP_017149923.1| PREDICTED: ubiquitin-like protein 5 isoform X1 [Drosophila r
>gi|1037007865|ref|XP_017149925.1| PREDICTED: uncharacterized protein LOC108160433 [Drosophila
>gi|1037000483|ref|XP_017149475.1| PREDICTED: probable RNA polymerase II nuclear localization p
>gi|1037000501|ref|XP_017149476.1| PREDICTED: uncharacterized protein LOC108160165 isoform X2
>gi|1037000519|ref|XP_017149477.1| PREDICTED: RNA-binding motif protein, X-linked 2-like [Dros
>gi|1036936228|ref|XP_017145580.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1036936246|ref|XP_017145581.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1036936264|ref|XP_017145582.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1037015580|ref|XP_017150385.1| PREDICTED: uncharacterized protein LOC108160720 [Drosophila
>gi|1036996220|ref|XP_017149219.1| PREDICTED: uncharacterized protein LOC108160004 isoform X1
>gi|1036996273|ref|XP_017149222.1| PREDICTED: uncharacterized protein LOC108160004 isoform X4
>gi|1036996238|ref|XP_017149220.1| PREDICTED: uncharacterized protein LOC108160004 isoform X2
>gi|1036996253|ref|XP_017149221.1| PREDICTED: uncharacterized protein LOC108160004 isoform X3
>gi|1037019151|ref|XP_017150613.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1037019167|ref|XP_017150614.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1036999876|ref|XP_017149440.1| PREDICTED: coronin-6 isoform X2 [Drosophila miranda]
>gi|1036999858|ref|XP_017149439.1| PREDICTED: coronin-6 isoform X1 [Drosophila miranda]
>gi|1036995984|ref|XP_017149203.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036995842|ref|XP_017149194.1| PREDICTED: alaserpin isoform X3 [Drosophila miranda]
>gi|1036995807|ref|XP_017149193.1| PREDICTED: alaserpin isoform X2 [Drosophila miranda]
>gi|1036995793|ref|XP_017149192.1| PREDICTED: alaserpin isoform X1 [Drosophila miranda]
>gi|1036995860|ref|XP_017149196.1| PREDICTED: alaserpin isoform X4 [Drosophila miranda]
>gi|1036995878|ref|XP_017149197.1| PREDICTED: alaserpin isoform X4 [Drosophila miranda]
>gi|1036995914|ref|XP_017149199.1| PREDICTED: serpin B3 isoform X1 [Drosophila miranda]
>gi|1036995928|ref|XP_017149200.1| PREDICTED: serpin B3 isoform X2 [Drosophila miranda]
>gi|1036995946|ref|XP_017149201.1| PREDICTED: serine protease inhibitor 3/4 isoform X3 [Drosopl
>gi|1036995896|ref|XP_017149198.1| PREDICTED: antichymotrypsin-2 [Drosophila miranda]
>gi|1036995966|ref|XP_017149202.1| PREDICTED: leukocyte elastase inhibitor [Drosophila miranda]
>gi|1037018201|ref|XP_017150548.1| PREDICTED: antichymotrypsin-2 [Drosophila miranda]
>gi|1037018186|ref|XP_017150546.1| PREDICTED: GTP-binding protein Di-Ras2 [Drosophila miranda]
>gi|1036979196|ref|XP_017148174.1| PREDICTED: putative deoxyribonuclease TATDN1 [Drosophila mi
>gi|1036979107|ref|XP_017148169.1| PREDICTED: MTSS1-like protein isoform X15 [Drosophila mirand
>gi|1036979144|ref|XP_017148171.1| PREDICTED: serine-rich adhesin for platelets isoform X17 [D1
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>gi|1036979091|ref|XP_017148168.1| PREDICTED: metastasis suppressor protein 1 isoform X14 [Dros
>gi|1036979176|ref|XP_017148173.1| PREDICTED: metastasis suppressor protein 1 isoform X19 [Dros
>gi|1036978982|ref|XP_017148161.1| PREDICTED: serine-rich adhesin for platelets isoform X8 [Dro
>gi|1036978874|ref|XP_017148155.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036979160|ref|XP_017148172.1| PREDICTED: uncharacterized protein LOC108159416 isoform X18
>gi|1036978910|ref|XP_017148157.1| PREDICTED: serine-rich adhesin for platelets isoform X4 [Dro
>gi|1036978892|ref|XP_017148156.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036979056|ref|XP_017148165.1| PREDICTED: AF4/FMR2 family member 4 isoform X12 [Drosophila
>gi|1036978944|ref|XP_017148159.1| PREDICTED: serine-rich adhesin for platelets isoform X6 [Dro
>gi|1036979074|ref|XP_017148166.1| PREDICTED: metastasis suppressor protein 1 isoform X13 [Dros
>gi|1036979019|ref|XP_017148163.1| PREDICTED: chromatin modification-related protein eaf-1 isor
>gi|1036979000|ref|XP_017148162.1| PREDICTED: serine-rich adhesin for platelets isoform X9 [Dro
>gi|1036978926|ref|XP_017148158.1| PREDICTED: serine-rich adhesin for platelets isoform X5 [Dro
>gi|1036978962|ref|XP_017148160.1| PREDICTED: serine-rich adhesin for platelets isoform X7 [Dro
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>gi|1036999039|ref|XP_017149395.1| PREDICTED: rap1 GTPase-GDP dissociation stimulator 1-B [Dros
>gi|1036999058|ref|XP_017149397.1| PREDICTED: dnaJ homolog subfamily B member 14 [Drosophila m
>gi|1036977682|ref|XP_017148081.1| PREDICTED: G protein pathway suppressor 2 isoform X1 [Droso
>gi|1036977712|ref|XP_017148083.1| PREDICTED: G protein pathway suppressor 2 isoform X3 [Droso
>gi|1036977694|ref|XP_017148082.1| PREDICTED: G protein pathway suppressor 2 isoform X2 [Droso
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>gi|1036977857|ref|XP_017148093.1| PREDICTED: 23 kDa integral membrane protein [Drosophila mira
>gi|1036977778|ref|XP_017148087.1| PREDICTED: 23 kDa integral membrane protein [Drosophila mira
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>gi|1036977814|ref|XP_017148089.1| PREDICTED: CD63 antigen [Drosophila miranda]
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>gi|1036977794|ref|XP_017148088.1| PREDICTED: 23 kDa integral membrane protein [Drosophila mira
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>gi|1037018112|ref|XP_017150541.1| PREDICTED: protein late bloomer [Drosophila miranda]
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>gi|1036941131|ref|XP_017145888.1| PREDICTED: 23 kDa integral membrane protein [Drosophila mira
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>gi|1036944024|ref|XP_017146066.1| PREDICTED: uncharacterized protein LOC108158337 [Drosophila
>gi|1037020580|ref|XP_017150702.1| PREDICTED: 23 kDa integral membrane protein [Drosophila mira
>gi|1037020595|ref|XP_017150703.1| PREDICTED: 23 kDa integral membrane protein [Drosophila mira
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>gi|1036943550|ref|XP_017146037.1| PREDICTED: protein late bloomer [Drosophila miranda]
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>gi|1037004887|ref|XP_017149736.1| PREDICTED: protein late bloomer [Drosophila miranda]
>gi|1037004870|ref|XP_017149735.1| PREDICTED: polypeptide N-acetylgalactosaminyltransferase 3
>gi|1037014491|ref|XP_017150322.1| PREDICTED: protein late bloomer [Drosophila miranda]
>gi|1037014509|ref|XP_017150323.1| PREDICTED: 23 kDa integral membrane protein-like [Drosophila
>gi|1036946396|ref|XP_017146205.1| PREDICTED: tctex1 domain-containing protein 2 [Drosophila mi
>gi|1036942017|ref|XP_017145942.1| PREDICTED: uncharacterized protein LOC108158222 [Drosophila
>gi|1037006762|ref|XP_017149858.1| PREDICTED: uncharacterized protein LOC108160387 isoform X1
>gi|1037006780|ref|XP_017149859.1| PREDICTED: uncharacterized protein LOC108160387 isoform X2
>gi|1036936314|ref|XP_017145586.1| PREDICTED: uncharacterized protein LOC108157953 [Drosophila
>gi|1037006724|ref|XP_017149856.1| PREDICTED: protein espinas [Drosophila miranda]
>gi|1037006744|ref|XP_017149857.1| PREDICTED: LOW QUALITY PROTEIN: cytochrome P450 9b2-like [Dr
>gi|1036936330|ref|XP_017145588.1| PREDICTED: cytochrome P450 9b2-like [Drosophila miranda]
>gi|1036992567|ref|XP_017148991.1| PREDICTED: uncharacterized protein LOC108159886 isoform X1
>gi|1036992581|ref|XP_017148992.1| PREDICTED: uncharacterized protein LOC108159886 isoform X2
>gi|1036992640|ref|XP_017148996.1| PREDICTED: alaserpin isoform X2 [Drosophila miranda]
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>gi|1036992613|ref|XP_017148994.1| PREDICTED: antichymotrypsin-2 [Drosophila miranda]
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>gi|1036992513|ref|XP_017148988.1| PREDICTED: protein prickle isoform X1 [Drosophila miranda]
>gi|1036992531|ref|XP_017148989.1| PREDICTED: protein prickle isoform X2 [Drosophila miranda]
>gi|1036943146|ref|XP_017146013.1| PREDICTED: odorant receptor 43a [Drosophila miranda]
>gi|1037019133|ref|XP_017150612.1| PREDICTED: adenosine kinase [Drosophila miranda]
>gi|1037022055|ref|XP_017150795.1| PREDICTED: growth arrest and DNA damage-inducible protein GA
>gi|1036943512|ref|XP_017146035.1| PREDICTED: uncharacterized protein LOC108158306 [Drosophila
>gi|1036999902|ref|XP_017149442.1| PREDICTED: peregrin [Drosophila miranda]
>gi|1037002252|ref|XP_017149588.1| PREDICTED: caldesmon isoform X1 [Drosophila miranda]
>gi|1037002278|ref|XP_017149589.1| PREDICTED: caldesmon isoform X2 [Drosophila miranda]
>gi|1037002296|ref|XP_017149590.1| PREDICTED: uncharacterized protein LOC108160242 [Drosophila
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>gi|1037008292|ref|XP_017149953.1| PREDICTED: protein piccolo isoform X1 [Drosophila miranda]
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>gi|1037008358|ref|XP_017149956.1| PREDICTED: protein piccolo isoform X2 [Drosophila miranda]
>gi|1037008419|ref|XP_017149957.1| PREDICTED: protein piccolo isoform X3 [Drosophila miranda]
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>gi|1036956183|ref|XP_017146803.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036956033|ref|XP_017146794.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
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>gi|1036956051|ref|XP_017146795.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
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>gi|1037003642|ref|XP_017149669.1| PREDICTED: putative pre-mRNA-splicing factor ATP-dependent
>gi|1037003658|ref|XP_017149670.1| PREDICTED: putative pre-mRNA-splicing factor ATP-dependent
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>gi|1037003676|ref|XP_017149671.1| PREDICTED: pre-mRNA-splicing factor ATP-dependent RNA helica
>gi|1037003695|ref|XP_017149672.1| PREDICTED: gustatory receptor for sugar taste 43a [Drosophil
>gi|1036994481|ref|XP_017149111.1| PREDICTED: ell-associated factor Eaf [Drosophila miranda]
>gi|1036994499|ref|XP_017149112.1| PREDICTED: ell-associated factor Eaf [Drosophila miranda]
>gi|1036994516|ref|XP_017149113.1| PREDICTED: ell-associated factor Eaf [Drosophila miranda]
>gi|1037012191|ref|XP_017150181.1| PREDICTED: protein sine oculis [Drosophila miranda]
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>gi|1036985753|ref|XP_017148585.1| PREDICTED: nucleolar protein 9 [Drosophila miranda]
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>gi|1036985727|ref|XP_017148583.1| PREDICTED: transcription elongation factor SPT5 [Drosophila
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>gi|1036976383|ref|XP_017148000.1| PREDICTED: benign gonial cell neoplasm protein isoform X2 [
>gi|1036976440|ref|XP_017148003.1| PREDICTED: partner of Y14 and mago [Drosophila miranda]
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>gi|1037002499|ref|XP_017149602.1| PREDICTED: TAR DNA-binding protein 43 isoform X1 [Drosophila
>gi|1037011942|ref|XP_017150164.1| PREDICTED: ceramide phosphoethanolamine synthase [Drosophila
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>gi|1037020202|ref|XP_017150678.1| PREDICTED: uncharacterized protein LOC108160912 [Drosophila
>gi|1037016749|ref|XP_017150454.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcompl
>gi|1037011765|ref|XP_017150153.1| PREDICTED: trifunctional enzyme subunit beta, mitochondrial
>gi|1037006205|ref|XP_017149822.1| PREDICTED: transcription factor Ken isoform X3 [Drosophila
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>gi|1036982511|ref|XP_017148384.1| PREDICTED: uncharacterized protein LOC108159526 isoform X2
>gi|1036982528|ref|XP_017148385.1| PREDICTED: methyltransferase-like protein 9 [Drosophila mir
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>gi|1037009493|ref|XP_017150025.1| PREDICTED: uncharacterized protein LOC108160499 [Drosophila
>gi|1037009475|ref|XP_017150024.1| PREDICTED: nucleosome assembly protein 1-like 1 [Drosophila
>gi|1036962206|ref|XP_017147152.1| PREDICTED: solute carrier family 12 member 6 isoform X1 [Dro
>gi|1036962280|ref|XP_017147157.1| PREDICTED: solute carrier family 12 member 6 isoform X5 [Dro
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>gi|1036962296|ref|XP_017147158.1| PREDICTED: solute carrier family 12 member 6 isoform X6 [Dro
>gi|1036962352|ref|XP_017147161.1| PREDICTED: solute carrier family 12 member 6 isoform X9 [Dro
>gi|1036962246|ref|XP_017147155.1| PREDICTED: solute carrier family 12 member 6 isoform X3 [Dro
>gi|1036962266|ref|XP_017147156.1| PREDICTED: solute carrier family 12 member 6 isoform X4 [Dro
>gi|1036962314|ref|XP_017147159.1| PREDICTED: solute carrier family 12 member 6 isoform X7 [Dro
>gi|1036962370|ref|XP_017147162.1| PREDICTED: solute carrier family 12 member 6 isoform X10 [Di
>gi|1036962471|ref|XP_017147169.1| PREDICTED: WD repeat-containing protein 47 [Drosophila miran
>gi|1036962418|ref|XP_017147166.1| PREDICTED: calcium-transporting ATPase sarcoplasmic/endoplas
>gi|1036962436|ref|XP_017147167.1| PREDICTED: calcium-transporting ATPase sarcoplasmic/endoplas
>gi|1036962388|ref|XP_017147163.1| PREDICTED: calcium-transporting ATPase sarcoplasmic/endoplas
>gi|1036962402|ref|XP_017147165.1| PREDICTED: calcium-transporting ATPase sarcoplasmic/endoplas
>gi|1036962454|ref|XP_017147168.1| PREDICTED: digestive organ expansion factor homolog [Drosoph
>gi|1036958897|ref|XP_017146957.1| PREDICTED: sulfhydryl oxidase 1-like [Drosophila miranda]
>gi|1036958920|ref|XP_017146960.1| PREDICTED: probable palmitoyltransferase ZDHHC24 [Drosophila
>gi|1036958932|ref|XP_017146961.1| PREDICTED: vacuolar protein-sorting-associated protein 25 [D
>gi|1036958841|ref|XP_017146953.1| PREDICTED: latrophilin Cirl isoform X1 [Drosophila miranda]
>gi|1036958859|ref|XP_017146954.1| PREDICTED: latrophilin Cirl isoform X2 [Drosophila miranda]
>gi|1036958870|ref|XP_017146955.1| PREDICTED: latrophilin Cirl isoform X2 [Drosophila miranda]
>gi|1036958884|ref|XP_017146956.1| PREDICTED: latrophilin Cirl isoform X3 [Drosophila miranda]
>gi|1036958907|ref|XP_017146958.1| PREDICTED: uncharacterized protein LOC108158788 [Drosophila
>gi|1036961592|ref|XP_017147115.1| PREDICTED: ecdysone-induced protein 75B isoform X2 [Drosoph
>gi|1036961648|ref|XP_017147118.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036961609|ref|XP_017147116.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036961574|ref|XP_017147114.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036961627|ref|XP_017147117.1| PREDICTED: ecdysone-induced protein 75B isoform X4 [Drosoph
>gi|1036961666|ref|XP_017147119.1| PREDICTED: angiotensin-converting enzyme [Drosophila miranda
>gi|1036961390|ref|XP_017147103.1| PREDICTED: sodium channel protein 60E isoform X1 [Drosophila
>gi|1036961408|ref|XP_017147104.1| PREDICTED: sodium channel protein 60E isoform X2 [Drosophila
>gi|1036961450|ref|XP_017147106.1| PREDICTED: sodium channel protein 60E isoform X4 [Drosophila
>gi|1036961427|ref|XP_017147105.1| PREDICTED: sodium channel protein 60E isoform X3 [Drosophila
>gi|1036961686|ref|XP_017147120.1| PREDICTED: loricrin [Drosophila miranda]
>gi|1036961704|ref|XP_017147121.1| PREDICTED: mitochondrial thiamine pyrophosphate carrier [Dro
>gi|1036961721|ref|XP_017147122.1| PREDICTED: mitochondrial thiamine pyrophosphate carrier [Dro
>gi|1036961743|ref|XP_017147123.1| PREDICTED: mitochondrial thiamine pyrophosphate carrier [Dro
>gi|1036961478|ref|XP_017147107.1| PREDICTED: LOW QUALITY PROTEIN: transient receptor potential
>gi|1036961560|ref|XP_017147113.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 isoform
>gi|1036961514|ref|XP_017147109.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 isoform
>gi|1036961529|ref|XP_017147110.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 isoform
>gi|1036961546|ref|XP_017147112.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 isoform
>gi|1036961496|ref|XP_017147108.1| PREDICTED: fatty acyl-CoA reductase 1 isoform X1 [Drosophila
>gi|1036986030|ref|XP_017148602.1| PREDICTED: protein CDV3 homolog [Drosophila miranda]
>gi|1036986101|ref|XP_017148608.1| PREDICTED: putative gamma-glutamylcyclotransferase CG2811 is
>gi|1036986087|ref|XP_017148606.1| PREDICTED: putative gamma-glutamylcyclotransferase CG2811 is
>gi|1036986073|ref|XP_017148605.1| PREDICTED: troponin C-akin-1 protein [Drosophila miranda]
>gi|1036986057|ref|XP_017148604.1| PREDICTED: wnt inhibitory factor 1 [Drosophila miranda]
>gi|1036986040|ref|XP_017148603.1| PREDICTED: lipoma HMGIC fusion partner-like 2 protein [Droso
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>gi|1036986012|ref|XP_017148601.1| PREDICTED: dnaJ homolog subfamily C member 21 [Drosophila m:
>gi|1037005044|ref|XP_017149746.1| PREDICTED: uncharacterized protein LOC108160328 isoform X1
>gi|1037005075|ref|XP_017149747.1| PREDICTED: uncharacterized protein LOC108160328 isoform X2
>gi|1036992798|ref|XP_017149007.1| PREDICTED: dentin sialophosphoprotein [Drosophila miranda]
>gi|1036992781|ref|XP_017149005.1| PREDICTED: uncharacterized protein LOC108159894 [Drosophila
>gi|1036992816|ref|XP_017149008.1| PREDICTED: 60S ribosomal protein L19 [Drosophila miranda]
>gi|1037007144|ref|XP_017149883.1| PREDICTED: uncharacterized protein C2orf47 homolog, mitocho:
>gi|1037007158|ref|XP_017149884.1| PREDICTED: ejaculatory bulb-specific protein 3 [Drosophila r
>gi|1037007179|ref|XP_017149885.1| PREDICTED: scavenger receptor class B member 1 [Drosophila r
>gi|1036950516|ref|XP_017146466.1| PREDICTED: scavenger receptor class B member 1 isoform X3 [I
>gi|1036950500|ref|XP_017146465.1| PREDICTED: lysosome membrane protein 2 isoform X2 [Drosophil
>gi|1036950454|ref|XP_017146462.1| PREDICTED: scavenger receptor class B member 1 isoform X1 [I
>gi|1036950468|ref|XP_017146463.1| PREDICTED: scavenger receptor class B member 1 isoform X1 [I
>gi|1036950485|ref|XP_017146464.1| PREDICTED: scavenger receptor class B member 1 isoform X1 [I
>gi|1036950422|ref|XP_017146460.1| PREDICTED: scavenger receptor class B member 1 [Drosophila r
>gi|1036950437|ref|XP_017146461.1| PREDICTED: scavenger receptor class B member 1 [Drosophila r
>gi|1036950386|ref|XP_017146457.1| PREDICTED: myosin heavy chain, non-muscle isoform X1 [Drosop
>gi|1036950404|ref|XP_017146458.1| PREDICTED: myosin heavy chain, non-muscle isoform X2 [Drosop
>gi|1036950533|ref|XP_017146467.1| PREDICTED: dolichol kinase isoform X1 [Drosophila miranda]
>gi|1036950550|ref|XP_017146468.1| PREDICTED: dolichol kinase isoform X1 [Drosophila miranda]
>gi|1036950568|ref|XP_017146470.1| PREDICTED: dolichol kinase isoform X2 [Drosophila miranda]
>gi|1036950585|ref|XP_017146471.1| PREDICTED: cyclin-dependent kinase 4 isoform X1 [Drosophila
>gi|1036950603|ref|XP_017146472.1| PREDICTED: cyclin-dependent kinase 4 isoform X1 [Drosophila
>gi|1036950620|ref|XP_017146473.1| PREDICTED: cyclin-dependent kinase 4 isoform X2 [Drosophila
>gi|1036950638|ref|XP_017146474.1| PREDICTED: uncharacterized protein LOC108158560 [Drosophila
>gi|1037003245|ref|XP_017149642.1| PREDICTED: probable ATP-dependent RNA helicase DDX27 [Drosop
>gi|1037003263|ref|XP_017149644.1| PREDICTED: uncharacterized protein LOC108160273 [Drosophila
>gi|1036996378|ref|XP_017149228.1| PREDICTED: ankyrin repeat, SAM and basic leucine zipper doma
>gi|1036996362|ref|XP_017149227.1| PREDICTED: DNA mismatch repair protein Mlh1 [Drosophila mira
>gi|1036996396|ref|XP_017149229.1| PREDICTED: succinate dehydrogenase assembly factor 2-A, mito
>gi|1036996414|ref|XP_017149231.1| PREDICTED: succinate dehydrogenase assembly factor 2-A, mito
>gi|1036965593|ref|XP_017147352.1| PREDICTED: prolow-density lipoprotein receptor-related prote
>gi|1036965631|ref|XP_017147354.1| PREDICTED: accessory gland protein Acp62F-like [Drosophila r
>gi|1036965613|ref|XP_017147353.1| PREDICTED: putative uncharacterized protein DDB_G0274435 [Dr
>gi|1036942678|ref|XP_017145984.1| PREDICTED: uncharacterized protein LOC108158258 [Drosophila
>gi|1037012022|ref|XP_017150169.1| PREDICTED: protein deadpan [Drosophila miranda]
>gi|1037006272|ref|XP_017149827.1| PREDICTED: protein peanut [Drosophila miranda]
>gi|1036936395|ref|XP_017145592.1| PREDICTED: venom serine protease 34 [Drosophila miranda]
>gi|1036936413|ref|XP_017145593.1| PREDICTED: uncharacterized protein LOC108157959 [Drosophila
>gi|1037022005|ref|XP_017150792.1| PREDICTED: uncharacterized protein LOC108160995 [Drosophila
>gi|1036942648|ref|XP_017145982.1| PREDICTED: venom serine protease [Drosophila miranda]
>gi|1036941201|ref|XP_017145892.1| PREDICTED: venom serine protease [Drosophila miranda]
>gi|1036941647|ref|XP_017145916.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1037005320|ref|XP_017149761.1| PREDICTED: alpha-protein kinase 1 isoform X1 [Drosophila mir
>gi|1037005338|ref|XP_017149762.1| PREDICTED: alpha-protein kinase 1 isoform X2 [Drosophila mir
>gi|1037005352|ref|XP_017149763.1| PREDICTED: alpha-protein kinase 1 isoform X3 [Drosophila mir
>gi|1037005376|ref|XP_017149764.1| PREDICTED: alpha-protein kinase 1 isoform X4 [Drosophila mir
>gi|1037005392|ref|XP_017149765.1| PREDICTED: alpha-protein kinase 1 isoform X5 [Drosophila mir
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>gi|1037005410|ref|XP_017149766.1| PREDICTED: alpha-protein kinase 1 isoform X6 [Drosophila mi
>gi|1037005496|ref|XP_017149772.1| PREDICTED: uncharacterized protein LOC108160342 [Drosophila
>gi|1037005444|ref|XP_017149769.1| PREDICTED: uncharacterized protein LOC108160339 [Drosophila
>gi|1037005462|ref|XP_017149770.1| PREDICTED: uncharacterized protein LOC108160340 [Drosophila
>gi|1037005480|ref|XP_017149771.1| PREDICTED: uncharacterized protein LOC108160341 [Drosophila
>gi|1037005428|ref|XP_017149768.1| PREDICTED: POU domain, class 6, transcription factor 2 isofo
>gi|1037017175|ref|XP_017150482.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1037012716|ref|XP_017150213.1| PREDICTED: COP9 signalosome complex subunit 7 [Drosophila mi
>gi|1037020761|ref|XP_017150715.1| PREDICTED: larval cuticle protein 1 [Drosophila miranda]
>gi|1037020777|ref|XP_017150716.1| PREDICTED: larval cuticle protein 2 [Drosophila miranda]
>gi|1036943128|ref|XP_017146012.1| PREDICTED: larval cuticle protein III/IV [Drosophila miranda
>gi|1036943332|ref|XP_017146025.1| PREDICTED: larval cuticle protein III/IV [Drosophila miranda
>gi|1037005617|ref|XP_017149780.1| PREDICTED: probable cytochrome P450 4ad1 [Drosophila miranda
>gi|1037005512|ref|XP_017149773.1| PREDICTED: cytochrome P450 4e2 [Drosophila miranda]
>gi|1037005530|ref|XP_017149774.1| PREDICTED: cytochrome P450 4e2 [Drosophila miranda]
>gi|1037005564|ref|XP_017149777.1| PREDICTED: cytochrome P450 4e2-like isoform X2 [Drosophila m
>gi|1037005546|ref|XP_017149776.1| PREDICTED: cytochrome P450 4e2-like isoform X1 [Drosophila m
>gi|1037005582|ref|XP_017149778.1| PREDICTED: cytochrome P450 4e2-like isoform X3 [Drosophila m
>gi|1037005600|ref|XP_017149779.1| PREDICTED: cytochrome P450 4e2-like [Drosophila miranda]
>gi|1037019663|ref|XP_017150643.1| PREDICTED: maltase A1 [Drosophila miranda]
>gi|1037020392|ref|XP_017150691.1| PREDICTED: maltase A2 [Drosophila miranda]
>gi|1037020941|ref|XP_017150726.1| PREDICTED: maltase A3 [Drosophila miranda]
>gi|1037021143|ref|XP_017150740.1| PREDICTED: maltase A3 [Drosophila miranda]
>gi|1037008500|ref|XP_017149963.1| PREDICTED: maltase A3 [Drosophila miranda]
>gi|1037008516|ref|XP_017149964.1| PREDICTED: maltase A3 [Drosophila miranda]
>gi|1037012090|ref|XP_017150174.1| PREDICTED: maltase A1 [Drosophila miranda]
>gi|1037012107|ref|XP_017150175.1| PREDICTED: maltase A1-like [Drosophila miranda]
>gi|1037012140|ref|XP_017150177.1| PREDICTED: metabotropic glutamate receptor 6 isoform X2 [Dro
>gi|1037012124|ref|XP_017150176.1| PREDICTED: metabotropic glutamate receptor 2 isoform X1 [Dro
>gi|1036944074|ref|XP_017146068.1| PREDICTED: gram-negative bacteria-binding protein 3 [Drosoph
>gi|1037019563|ref|XP_017150638.1| PREDICTED: LOW QUALITY PROTEIN: F-box only protein 39 [Droso
>gi|1037019616|ref|XP_017150640.1| PREDICTED: transient receptor potential cation channel prote
>gi|1036936431|ref|XP_017145594.1| PREDICTED: myosin heavy chain, non-muscle-like [Drosophila m
>gi|1036936449|ref|XP_017145596.1| PREDICTED: Krueppel-like factor 8 [Drosophila miranda]
>gi|1037013353|ref|XP_017150252.1| PREDICTED: protein twist [Drosophila miranda]
>gi|1037011013|ref|XP_017150108.1| PREDICTED: long-chain fatty acid transport protein 1 [Drosop
>gi|1037011027|ref|XP_017150109.1| PREDICTED: long-chain fatty acid transport protein 1 [Drosop
>gi|1037011043|ref|XP_017150110.1| PREDICTED: long-chain fatty acid transport protein 1 [Drosop
>gi|1037011061|ref|XP_017150111.1| PREDICTED: long-chain fatty acid transport protein 1 [Drosop
>gi|1037011126|ref|XP_017150115.1| PREDICTED: uncharacterized protein LOC108160553 [Drosophila
>gi|1037011094|ref|XP_017150113.1| PREDICTED: uncharacterized protein LOC108160552 isoform X2
>gi|1037011108|ref|XP_017150114.1| PREDICTED: uncharacterized protein LOC108160552 isoform X3
>gi|1037011083|ref|XP_017150112.1| PREDICTED: uncharacterized protein LOC108160552 isoform X1
>gi|1037013844|ref|XP_017150283.1| PREDICTED: zinc finger protein 277 [Drosophila miranda]
>gi|1036998279|ref|XP_017149349.1| PREDICTED: putative inorganic phosphate cotransporter [Droso
>gi|1036998221|ref|XP_017149345.1| PREDICTED: putative inorganic phosphate cotransporter isofor
>gi|1036998238|ref|XP_017149346.1| PREDICTED: putative inorganic phosphate cotransporter isofor
>gi|1036998261|ref|XP_017149348.1| PREDICTED: prohibitin-2 [Drosophila miranda]
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>gi|1036998205|ref|XP_017149344.1| PREDICTED: kelch-like protein 17 [Drosophila miranda]
>gi|1037013975|ref|XP_017150290.1| PREDICTED: uncharacterized protein LOC108160661 isoform X1
>gi|1037013990|ref|XP_017150291.1| PREDICTED: uncharacterized protein LOC108160661 isoform X2
>gi|1037014004|ref|XP_017150292.1| PREDICTED: uncharacterized protein LOC108160662 [Drosophila
>gi|1036982070|ref|XP_017148357.1| PREDICTED: uncharacterized protein LOC108159509 [Drosophila
>gi|1036982054|ref|XP_017148356.1| PREDICTED: protein dopey-1 homolog [Drosophila miranda]
>gi|1037004368|ref|XP_017149708.1| PREDICTED: methionine--tRNA ligase, cytoplasmic [Drosophila
>gi|1036964813|ref|XP_017147303.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcompl
>gi|1036964831|ref|XP_017147304.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcompl
>gi|1036964550|ref|XP_017147289.1| PREDICTED: tyrosine-protein kinase transmembrane receptor Ro
>gi|1036964760|ref|XP_017147300.1| PREDICTED: apolipoprotein D [Drosophila miranda]
>gi|1036964777|ref|XP_017147301.1| PREDICTED: apolipoprotein D [Drosophila miranda]
>gi|1036964793|ref|XP_017147302.1| PREDICTED: apolipoprotein D [Drosophila miranda]
>gi|1036964677|ref|XP_017147297.1| PREDICTED: peptidyl-prolyl cis-trans isomerase FKBP8 [Droso
>gi|1036964694|ref|XP_017147299.1| PREDICTED: lysosomal aspartic protease [Drosophila miranda]
>gi|1036964631|ref|XP_017147294.1| PREDICTED: protein ABHD4 isoform X1 [Drosophila miranda]
>gi|1036964649|ref|XP_017147295.1| PREDICTED: protein ABHD4 isoform X2 [Drosophila miranda]
>gi|1036964659|ref|XP_017147296.1| PREDICTED: protein ABHD4 isoform X2 [Drosophila miranda]
>gi|1036964565|ref|XP_017147290.1| PREDICTED: tubulin--tyrosine ligase-like protein 12 isoform
>gi|1036964580|ref|XP_017147291.1| PREDICTED: tubulin--tyrosine ligase-like protein 12 isoform
>gi|1036964599|ref|XP_017147292.1| PREDICTED: activin receptor type-1 isoform X1 [Drosophila m
>gi|1036964615|ref|XP_017147293.1| PREDICTED: activin receptor type-1 isoform X2 [Drosophila m
>gi|1036980452|ref|XP_017148257.1| PREDICTED: protein kintoun isoform X1 [Drosophila miranda]
>gi|1036980470|ref|XP_017148258.1| PREDICTED: protein kintoun isoform X1 [Drosophila miranda]
>gi|1036980486|ref|XP_017148259.1| PREDICTED: protein kintoun isoform X2 [Drosophila miranda]
>gi|1036980398|ref|XP_017148254.1| PREDICTED: DNA polymerase zeta catalytic subunit isoform X1
>gi|1036980418|ref|XP_017148255.1| PREDICTED: DNA polymerase zeta catalytic subunit isoform X2
>gi|1036980434|ref|XP_017148256.1| PREDICTED: DNA polymerase zeta catalytic subunit isoform X3
>gi|1036969950|ref|XP_017147603.1| PREDICTED: kynurenine 3-monooxygenase isoform X1 [Drosophila
>gi|1036969968|ref|XP_017147605.1| PREDICTED: kynurenine 3-monooxygenase isoform X1 [Drosophila
>gi|1036970046|ref|XP_017147610.1| PREDICTED: uncharacterized protein LOC108159129 [Drosophila
>gi|1036970035|ref|XP_017147609.1| PREDICTED: calcineurin subunit B type 2 [Drosophila miranda]
>gi|1036969986|ref|XP_017147606.1| PREDICTED: kynurenine 3-monooxygenase isoform X2 [Drosophila
>gi|1036970003|ref|XP_017147607.1| PREDICTED: kynurenine 3-monooxygenase isoform X2 [Drosophila
>gi|1036970023|ref|XP_017147608.1| PREDICTED: kynurenine 3-monooxygenase isoform X2 [Drosophila
>gi|1036969878|ref|XP_017147599.1| PREDICTED: uncharacterized protein LOC108159125 isoform X1
>gi|1036969900|ref|XP_017147600.1| PREDICTED: uncharacterized protein LOC108159125 isoform X1
>gi|1036969914|ref|XP_017147601.1| PREDICTED: protein FAM46C isoform X2 [Drosophila miranda]
>gi|1036969932|ref|XP_017147602.1| PREDICTED: protein FAM46C isoform X3 [Drosophila miranda]
>gi|1036936467|ref|XP_017145597.1| PREDICTED: ras guanine nucleotide exchange factor P, partial
>gi|1037001045|ref|XP_017149508.1| PREDICTED: protein GDAP2 homolog isoform X1 [Drosophila mira
>gi|1037001065|ref|XP_017149509.1| PREDICTED: protein GDAP2 homolog isoform X2 [Drosophila mira
>gi|1037001080|ref|XP_017149510.1| PREDICTED: protein GDAP2 homolog isoform X3 [Drosophila mira
>gi|1037001103|ref|XP_017149512.1| PREDICTED: protein GDAP2 homolog isoform X4 [Drosophila mira
>gi|1037001118|ref|XP_017149513.1| PREDICTED: diacylglycerol O-acyltransferase 2-like [Drosophi
>gi|1036971670|ref|XP_017147704.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971616|ref|XP_017147701.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971652|ref|XP_017147703.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
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>gi|1036971577|ref|XP_017147699.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971596|ref|XP_017147700.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971634|ref|XP_017147702.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971489|ref|XP_017147694.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971507|ref|XP_017147695.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971525|ref|XP_017147696.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971540|ref|XP_017147697.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971559|ref|XP_017147698.1| PREDICTED: SH3 and multiple ankyrin repeat domains protein
>gi|1036971758|ref|XP_017147709.1| PREDICTED: uncharacterized protein LOC108159174 [Drosophila
>gi|1036971686|ref|XP_017147705.1| PREDICTED: uncharacterized protein LOC108159172 isoform X1
>gi|1036971702|ref|XP_017147706.1| PREDICTED: uncharacterized protein LOC108159172 isoform X2
>gi|1036971722|ref|XP_017147707.1| PREDICTED: uncharacterized protein LOC108159172 isoform X3
>gi|1036971740|ref|XP_017147708.1| PREDICTED: uncharacterized protein LOC108159173 [Drosophila
>gi|1036977316|ref|XP_017148058.1| PREDICTED: uncharacterized protein LOC108159352 isoform X7
>gi|1036977298|ref|XP_017148057.1| PREDICTED: uncharacterized protein LOC108159352 isoform X6
>gi|1036977282|ref|XP_017148056.1| PREDICTED: uncharacterized protein LOC108159352 isoform X5
>gi|1036977266|ref|XP_017148055.1| PREDICTED: uncharacterized protein LOC108159352 isoform X4
>gi|1036977248|ref|XP_017148054.1| PREDICTED: uncharacterized protein LOC108159352 isoform X3
>gi|1036977230|ref|XP_017148053.1| PREDICTED: uncharacterized protein LOC108159352 isoform X2
>gi|1036977212|ref|XP_017148052.1| PREDICTED: uncharacterized protein LOC108159352 isoform X1
>gi|1036977149|ref|XP_017148048.1| PREDICTED: ATP-dependent RNA helicase DHX8 isoform X1 [Droso
>gi|1036977164|ref|XP_017148049.1| PREDICTED: ATP-dependent RNA helicase DHX8 isoform X2 [Droso
>gi|1036977194|ref|XP_017148051.1| PREDICTED: ATP-dependent RNA helicase DHX8 isoform X3 [Droso
>gi|1036977334|ref|XP_017148059.1| PREDICTED: cytochrome c oxidase assembly factor 5 [Drosophil
>gi|1036990039|ref|XP_017148844.1| PREDICTED: transmembrane protein 39A [Drosophila miranda]
>gi|1036990021|ref|XP_017148843.1| PREDICTED: probable cysteine--tRNA ligase, mitochondrial [Dr
>gi|1036990055|ref|XP_017148845.1| PREDICTED: GDP-fucose protein O-fucosyltransferase 1 [Drosop
>gi|1036990075|ref|XP_017148846.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037009623|ref|XP_017150033.1| PREDICTED: DNA excision repair protein ERCC-1 [Drosophila mi
>gi|1037009606|ref|XP_017150032.1| PREDICTED: probable cytosolic iron-sulfur protein assembly p
>gi|1036951924|ref|XP_017146551.1| PREDICTED: uncharacterized protein LOC108158600 [Drosophila
>gi|1036951985|ref|XP_017146553.1| PREDICTED: transmembrane protein 177 [Drosophila miranda]
>gi|1036951851|ref|XP_017146545.1| PREDICTED: protein hu-li tai shao isoform X4 [Drosophila mi
>gi|1036951865|ref|XP_017146546.1| PREDICTED: protein hu-li tai shao isoform X5 [Drosophila mi
>gi|1036951833|ref|XP_017146544.1| PREDICTED: protein hu-li tai shao isoform X3 [Drosophila mi
>gi|1036951692|ref|XP_017146538.1| PREDICTED: uncharacterized protein LOC108158598 isoform X1
>gi|1036951726|ref|XP_017146539.1| PREDICTED: uncharacterized protein LOC108158598 isoform X1
>gi|1036951742|ref|XP_017146540.1| PREDICTED: uncharacterized protein LOC108158598 isoform X1
>gi|1036951759|ref|XP_017146541.1| PREDICTED: uncharacterized protein LOC108158598 isoform X1
>gi|1036951777|ref|XP_017146542.1| PREDICTED: uncharacterized protein LOC108158598 isoform X1
>gi|1036951801|ref|XP_017146543.1| PREDICTED: uncharacterized protein LOC108158598 isoform X2
>gi|1036951997|ref|XP_017146554.1| PREDICTED: protein CTLA-2-alpha [Drosophila miranda]
>gi|1036951878|ref|XP_017146547.1| PREDICTED: succinate dehydrogenase [ubiquinone] flavoprotei
>gi|1036951896|ref|XP_017146548.1| PREDICTED: succinate dehydrogenase [ubiquinone] flavoprotei
>gi|1036951910|ref|XP_017146549.1| PREDICTED: succinate dehydrogenase [ubiquinone] flavoprotei
>gi|1036951970|ref|XP_017146552.1| PREDICTED: cytochrome b5 reductase 4 [Drosophila miranda]
>gi|1037007003|ref|XP_017149874.1| PREDICTED: fatty-acid amide hydrolase 2-A [Drosophila mirand
>gi|1037007021|ref|XP_017149876.1| PREDICTED: sentrin-specific protease 8 isoform X1 [Drosophil
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>gi|1037007037|ref|XP_017149877.1| PREDICTED: sentrin-specific protease 8 isoform X2 [Drosophil
>gi|1037016179|ref|XP_017150419.1| PREDICTED: uncharacterized protein LOC108160739 [Drosophila
>gi|1036987446|ref|XP_017148688.1| PREDICTED: Golgi-specific brefeldin A-resistance guanine nuc
>gi|1036987464|ref|XP_017148689.1| PREDICTED: Golgi-specific brefeldin A-resistance guanine nuc
>gi|1036976018|ref|XP_017147977.1| PREDICTED: protein HID1 [Drosophila miranda]
>gi|1036976035|ref|XP_017147978.1| PREDICTED: protein HID1 [Drosophila miranda]
>gi|1036976339|ref|XP_017147997.1| PREDICTED: uncharacterized protein LOC108159319 isoform X1
>gi|1036976351|ref|XP_017147998.1| PREDICTED: uncharacterized protein LOC108159319 isoform X2
>gi|1036975966|ref|XP_017147974.1| PREDICTED: spondin-1 [Drosophila miranda]
>gi|1036975984|ref|XP_017147975.1| PREDICTED: spondin-1 [Drosophila miranda]
>gi|1036976002|ref|XP_017147976.1| PREDICTED: spondin-1 [Drosophila miranda]
>gi|1036976054|ref|XP_017147979.1| PREDICTED: myosin heavy chain, muscle-like isoform X1 [Dros
>gi|1036976070|ref|XP_017147980.1| PREDICTED: myosin heavy chain, muscle-like isoform X2 [Dros
>gi|1036976136|ref|XP_017147984.1| PREDICTED: myosin heavy chain, muscle-like isoform X4 [Dros
>gi|1036976153|ref|XP_017147985.1| PREDICTED: myosin heavy chain, muscle-like isoform X5 [Dros
>gi|1036976103|ref|XP_017147982.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036976120|ref|XP_017147983.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036976291|ref|XP_017147993.1| PREDICTED: uncharacterized protein LOC108159317 isoform X4
>gi|1036976325|ref|XP_017147995.1| PREDICTED: uncharacterized protein LOC108159317 isoform X6
>gi|1036976309|ref|XP_017147994.1| PREDICTED: uncharacterized protein LOC108159317 isoform X5
>gi|1036976277|ref|XP_017147992.1| PREDICTED: uncharacterized protein LOC108159317 isoform X3
>gi|1036976259|ref|XP_017147991.1| PREDICTED: uncharacterized protein LOC108159317 isoform X2
>gi|1036976170|ref|XP_017147987.1| PREDICTED: uncharacterized protein LOC108159317 isoform X1
>gi|1036976205|ref|XP_017147988.1| PREDICTED: uncharacterized protein LOC108159317 isoform X1
>gi|1036976224|ref|XP_017147989.1| PREDICTED: uncharacterized protein LOC108159317 isoform X1
>gi|1036976243|ref|XP_017147990.1| PREDICTED: uncharacterized protein LOC108159317 isoform X1
>gi|1036990161|ref|XP_017148852.1| PREDICTED: calmodulin [Drosophila miranda]
>gi|1036990143|ref|XP_017148851.1| PREDICTED: V-type proton ATPase subunit D 1 [Drosophila mira
>gi|1036990125|ref|XP_017148850.1| PREDICTED: golgin subfamily A member 6-like protein 22 [Dros
>gi|1036990251|ref|XP_017148858.1| PREDICTED: uncharacterized protein LOC108159803 [Drosophila
>gi|1036990179|ref|XP_017148853.1| PREDICTED: uncharacterized protein DDB_G0271670 [Drosophila
>gi|1036990215|ref|XP_017148855.1| PREDICTED: uncharacterized protein LOC108159801 isoform X2
>gi|1036990197|ref|XP_017148854.1| PREDICTED: uncharacterized protein LOC108159801 isoform X1
>gi|1036990233|ref|XP_017148857.1| PREDICTED: uncharacterized protein DDB_G0271670 [Drosophila
>gi|1036990093|ref|XP_017148848.1| PREDICTED: uncharacterized protein LOC108159796 [Drosophila
>gi|1036990111|ref|XP_017148849.1| PREDICTED: uncharacterized protein LOC108159796 [Drosophila
>gi|1037013221|ref|XP_017150246.1| PREDICTED: 40S ribosomal protein S11 [Drosophila miranda]
>gi|1036969199|ref|XP_017147558.1| PREDICTED: proteasome-associated protein ECM29 homolog [Dros
>gi|1036969215|ref|XP_017147559.1| PREDICTED: probable cytochrome P450 6t3 [Drosophila miranda]
>gi|1036969252|ref|XP_017147561.1| PREDICTED: probable cytochrome P450 6g2 [Drosophila miranda]
>gi|1036969232|ref|XP_017147560.1| PREDICTED: cytochrome P450 6g1 [Drosophila miranda]
>gi|1036969312|ref|XP_017147564.1| PREDICTED: methyltransferase-like protein 25 isoform X3 [Dro
>gi|1036969292|ref|XP_017147563.1| PREDICTED: methyltransferase-like protein 25 isoform X2 [Dro
>gi|1036969270|ref|XP_017147562.1| PREDICTED: methyltransferase-like protein 25 isoform X1 [Dro
>gi|1036969357|ref|XP_017147567.1| PREDICTED: small nuclear ribonucleoprotein F [Drosophila mir
>gi|1036969335|ref|XP_017147566.1| PREDICTED: oral cancer-overexpressed protein 1 homolog [Dros
>gi|1037006512|ref|XP_017149843.1| PREDICTED: WAS protein family homolog 1 [Drosophila miranda]
>gi|1037006575|ref|XP_017149847.1| PREDICTED: uncharacterized protein LOC108160380 [Drosophila
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>gi|1037006562|ref|XP_017149846.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037006530|ref|XP_017149844.1| PREDICTED: plant intracellular Ras-group-related LRR protei
>gi|1037006546|ref|XP_017149845.1| PREDICTED: immune-induced peptides [Drosophila miranda]
>gi|1036993942|ref|XP_017149080.1| PREDICTED: TATA-box-binding protein [Drosophila miranda]
>gi|1036993973|ref|XP_017149082.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036993991|ref|XP_017149083.1| PREDICTED: protein stum isoform X1 [Drosophila miranda]
>gi|1036994009|ref|XP_017149084.1| PREDICTED: protein stum isoform X2 [Drosophila miranda]
>gi|1036993955|ref|XP_017149081.1| PREDICTED: dr1-associated corepressor [Drosophila miranda]
>gi|1037000253|ref|XP_017149463.1| PREDICTED: cx9C motif-containing protein 4 isoform X1 [Droso
>gi|1037000271|ref|XP_017149464.1| PREDICTED: cx9C motif-containing protein 4 isoform X2 [Droso
>gi|1037000235|ref|XP_017149462.1| PREDICTED: uncharacterized protein LOC108160157 [Drosophila
>gi|1037000289|ref|XP_017149465.1| PREDICTED: uncharacterized protein LOC108160159 [Drosophila
>gi|1037000184|ref|XP_017149459.1| PREDICTED: GPI mannosyltransferase 1 [Drosophila miranda]
>gi|1037000200|ref|XP_017149460.1| PREDICTED: uncharacterized protein LOC108160154 [Drosophila
>gi|1037000307|ref|XP_017149467.1| PREDICTED: uncharacterized protein LOC108160160 [Drosophila
>gi|1037000218|ref|XP_017149461.1| PREDICTED: uncharacterized protein LOC108160155 [Drosophila
>gi|1037009044|ref|XP_017149997.1| PREDICTED: mRNA export factor [Drosophila miranda]
>gi|1037009062|ref|XP_017149998.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomple
>gi|1037009088|ref|XP_017149999.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomple
>gi|1036989432|ref|XP_017148806.1| PREDICTED: uncharacterized protein LOC108159760 isoform X1
>gi|1036989451|ref|XP_017148807.1| PREDICTED: uncharacterized protein LOC108159760 isoform X2
>gi|1036989467|ref|XP_017148808.1| PREDICTED: GRAM domain-containing protein 1B [Drosophila mi
>gi|1036965117|ref|XP_017147321.1| PREDICTED: uncharacterized protein LOC108158987 isoform X1
>gi|1036965137|ref|XP_017147322.1| PREDICTED: uncharacterized protein LOC108158987 isoform X2
>gi|1036965171|ref|XP_017147325.1| PREDICTED: uncharacterized protein LOC108158989 [Drosophila
>gi|1036965101|ref|XP_017147320.1| PREDICTED: protein mahjong isoform X2 [Drosophila miranda]
>gi|1036965085|ref|XP_017147319.1| PREDICTED: protein mahjong isoform X1 [Drosophila miranda]
>gi|1036965203|ref|XP_017147327.1| PREDICTED: protein TEX261 [Drosophila miranda]
>gi|1036965187|ref|XP_017147326.1| PREDICTED: ribose-5-phosphate isomerase [Drosophila miranda]
>gi|1036965153|ref|XP_017147324.1| PREDICTED: uncharacterized protein LOC108158988 [Drosophila
>gi|1037020826|ref|XP_017150720.1| PREDICTED: uncharacterized protein LOC108160942 [Drosophila
>gi|1037011301|ref|XP_017150126.1| PREDICTED: aminopeptidase N isoform X1 [Drosophila miranda]
>gi|1037011361|ref|XP_017150129.1| PREDICTED: aminopeptidase N isoform X3 [Drosophila miranda]
>gi|1037011319|ref|XP_017150127.1| PREDICTED: aminopeptidase N isoform X2 [Drosophila miranda]
>gi|1037011337|ref|XP_017150128.1| PREDICTED: aminopeptidase N isoform X2 [Drosophila miranda]
>gi|1036936485|ref|XP_017145598.1| PREDICTED: uncharacterized protein LOC108157963 [Drosophila
>gi|1037014712|ref|XP_017150335.1| PREDICTED: LOW QUALITY PROTEIN: bromodomain-containing prote
>gi|1037014678|ref|XP_017150333.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1037014694|ref|XP_017150334.1| PREDICTED: zinc finger CCCH domain-containing protein 13-lik
>gi|1036941734|ref|XP_017145921.1| PREDICTED: uncharacterized protein LOC108158210 [Drosophila
>gi|1037022203|ref|XP_017150803.1| PREDICTED: proteasome subunit beta type-5 [Drosophila miran
>gi|1036936503|ref|XP_017145599.1| PREDICTED: putative odorant receptor 65c [Drosophila miranda
>gi|1036971964|ref|XP_017147723.1| PREDICTED: putative odorant receptor 65c isoform X2 [Drosoph
>gi|1036971943|ref|XP_017147721.1| PREDICTED: odorant receptor 65a-like isoform X1 [Drosophila
>gi|1036971978|ref|XP_017147724.1| PREDICTED: putative odorant receptor 65c [Drosophila miranda
>gi|1036971995|ref|XP_017147725.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036972009|ref|XP_017147726.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036971915|ref|XP_017147719.1| PREDICTED: protein ORD [Drosophila miranda]
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>gi|1036971927|ref|XP_017147720.1| PREDICTED: protein ORD [Drosophila miranda]
>gi|1036972027|ref|XP_017147728.1| PREDICTED: uncharacterized protein LOC108159185 [Drosophila
>gi|1036936521|ref|XP_017145600.1| PREDICTED: uncharacterized protein LOC108157965 [Drosophila
>gi|1036971901|ref|XP_017147718.1| PREDICTED: splicing factor U2AF 50 kDa subunit [Drosophila
>gi|1036971879|ref|XP_017147717.1| PREDICTED: uncharacterized protein LOC108159179 [Drosophila
>gi|1036943368|ref|XP_017146027.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
>gi|1036945078|ref|XP_017146125.1| PREDICTED: uncharacterized protein LOC108158382 isoform X2
>gi|1036945060|ref|XP_017146124.1| PREDICTED: uncharacterized protein LOC108158382 isoform X1
>gi|1036943827|ref|XP_017146055.1| PREDICTED: uncharacterized protein LOC108158325 [Drosophila
>gi|1036978014|ref|XP_017148102.1| PREDICTED: uncharacterized protein LOC108159382 [Drosophila
>gi|1036978202|ref|XP_017148112.1| PREDICTED: uncharacterized protein LOC108159390 [Drosophila
>gi|1036977971|ref|XP_017148098.1| PREDICTED: facilitated trehalose transporter Tret1-like iso
>gi|1036977955|ref|XP_017148097.1| PREDICTED: facilitated trehalose transporter Tret1-like iso
>gi|1036977985|ref|XP_017148100.1| PREDICTED: facilitated trehalose transporter Tret1-like iso
>gi|1036978073|ref|XP_017148105.1| PREDICTED: glycerol-3-phosphate dehydrogenase [NAD(+)], cyto
>gi|1036977925|ref|XP_017148095.1| PREDICTED: kinesin-like protein Klp59C [Drosophila miranda]
>gi|1036936537|ref|XP_017145601.1| PREDICTED: putative gustatory receptor 59d [Drosophila mira
>gi|1036936555|ref|XP_017145602.1| PREDICTED: putative gustatory receptor 59c [Drosophila mira
>gi|1036978084|ref|XP_017148106.1| PREDICTED: paired mesoderm homeobox protein 2 [Drosophila m
>gi|1036978120|ref|XP_017148109.1| PREDICTED: uncharacterized protein LOC108159387 isoform X2
>gi|1036978100|ref|XP_017148108.1| PREDICTED: uncharacterized protein LOC108159387 isoform X1
>gi|1036978181|ref|XP_017148111.1| PREDICTED: neuropeptide-like protein 31 [Drosophila miranda]
>gi|1036978132|ref|XP_017148110.1| PREDICTED: uncharacterized protein LOC108159388 [Drosophila
>gi|1036978030|ref|XP_017148103.1| PREDICTED: putative gustatory receptor 59b [Drosophila mira
>gi|1036978047|ref|XP_017148104.1| PREDICTED: putative gustatory receptor 59b [Drosophila mira
>gi|1036936572|ref|XP_017145603.1| PREDICTED: putative gustatory receptor 59b [Drosophila mira
>gi|1036936590|ref|XP_017145604.1| PREDICTED: putative gustatory receptor 59b [Drosophila mira
>gi|1036978216|ref|XP_017148113.1| PREDICTED: uncharacterized protein LOC108159391 [Drosophila
>gi|1036977941|ref|XP_017148096.1| PREDICTED: protein arginine N-methyltransferase 7 [Drosophil
>gi|1036977999|ref|XP_017148101.1| PREDICTED: tektin-B1 [Drosophila miranda]
>gi|1037000359|ref|XP_017149470.1| PREDICTED: rRNA 2'-O-methyltransferase fibrillarin [Drosophi
>gi|1037000325|ref|XP_017149468.1| PREDICTED: transmembrane protein 53 [Drosophila miranda]
>gi|1037000341|ref|XP_017149469.1| PREDICTED: transmembrane protein 53 [Drosophila miranda]
>gi|1036993728|ref|XP_017149064.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036993746|ref|XP_017149065.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1036993762|ref|XP_017149066.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1036993712|ref|XP_017149063.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1036993676|ref|XP_017149061.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1036993694|ref|XP_017149062.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037015174|ref|XP_017150364.1| PREDICTED: probable 3-hydroxyisobutyrate dehydrogenase, mit
>gi|1037012489|ref|XP_017150198.1| PREDICTED: multiple C2 and transmembrane domain-containing
>gi|1037012473|ref|XP_017150197.1| PREDICTED: multiple C2 and transmembrane domain-containing
>gi|1037012508|ref|XP_017150199.1| PREDICTED: multiple C2 and transmembrane domain-containing
>gi|1037012522|ref|XP_017150201.1| PREDICTED: multiple C2 and transmembrane domain-containing
>gi|1037002915|ref|XP_017149619.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X
>gi|1037002939|ref|XP_017149622.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X3
>gi|1037002953|ref|XP_017149623.1| PREDICTED: uncharacterized protein LOC108160262 isoform X4
>gi|1037002928|ref|XP_017149620.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X2
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>gi|1037002967|ref|XP_017149624.1| PREDICTED: uncharacterized protein LOC108160262 isoform X5
>gi|1037002982|ref|XP_017149625.1| PREDICTED: uncharacterized protein LOC108160262 isoform X6
>gi|1037003002|ref|XP_017149626.1| PREDICTED: probable cytochrome P450 12b2, mitochondrial [Dro
>gi|1037003017|ref|XP_017149627.1| PREDICTED: uncharacterized protein LOC108160264 [Drosophila
>gi|1037001696|ref|XP_017149551.1| PREDICTED: uncharacterized protein LOC108160214 [Drosophila
>gi|1037001713|ref|XP_017149552.1| PREDICTED: uncharacterized protein LOC108160214 [Drosophila
>gi|1037001790|ref|XP_017149557.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037001773|ref|XP_017149556.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037001844|ref|XP_017149560.1| PREDICTED: probable salivary secreted peptide [Drosophila m
>gi|1037001826|ref|XP_017149559.1| PREDICTED: histone-like protein 18C [Drosophila miranda]
>gi|1037001808|ref|XP_017149558.1| PREDICTED: histone-like protein 18C [Drosophila miranda]
>gi|1037001727|ref|XP_017149553.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1037001741|ref|XP_017149554.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1037001755|ref|XP_017149555.1| PREDICTED: LOW QUALITY PROTEIN: odorant receptor 98a-like [
>gi|1036992915|ref|XP_017149013.1| PREDICTED: vacuolar protein sorting-associated protein 51 h
>gi|1036992933|ref|XP_017149014.1| PREDICTED: transcription factor grauzone [Drosophila mirand
>gi|1036992951|ref|XP_017149015.1| PREDICTED: caspase-1-like isoform X1 [Drosophila miranda]
>gi|1036992967|ref|XP_017149016.1| PREDICTED: caspase-1-like isoform X2 [Drosophila miranda]
>gi|1036992981|ref|XP_017149018.1| PREDICTED: caspase-1-like isoform X3 [Drosophila miranda]
>gi|1036977468|ref|XP_017148068.1| PREDICTED: uncharacterized protein LOC108159356 isoform X2
>gi|1036977454|ref|XP_017148067.1| PREDICTED: uncharacterized protein LOC108159356 isoform X1
>gi|1036977482|ref|XP_017148069.1| PREDICTED: uncharacterized protein LOC108159357 [Drosophila
>gi|1036977529|ref|XP_017148073.1| PREDICTED: uncharacterized protein LOC108159360 [Drosophila
>gi|1036977498|ref|XP_017148070.1| PREDICTED: uncharacterized protein LOC108159358 [Drosophila
>gi|1036977514|ref|XP_017148071.1| PREDICTED: uncharacterized protein LOC108159359 [Drosophila
>gi|1036977567|ref|XP_017148075.1| PREDICTED: serine protease easter-like isoform X2 [Drosophil
>gi|1036977549|ref|XP_017148074.1| PREDICTED: serine protease easter-like isoform X1 [Drosophil
>gi|1036977585|ref|XP_017148076.1| PREDICTED: serine protease easter-like isoform X3 [Drosophil
>gi|1036977599|ref|XP_017148077.1| PREDICTED: coagulation factor X-like [Drosophila miranda]
>gi|1036977617|ref|XP_017148078.1| PREDICTED: uncharacterized protein LOC108159364 [Drosophila
>gi|1036947671|ref|XP_017146285.1| PREDICTED: uncharacterized protein LOC108158488 [Drosophila
>gi|1036947582|ref|XP_017146279.1| PREDICTED: UBX domain-containing protein 6 [Drosophila mira
>gi|1036947654|ref|XP_017146284.1| PREDICTED: protein N-terminal asparagine amidohydrolase iso
>gi|1036947634|ref|XP_017146283.1| PREDICTED: protein N-terminal asparagine amidohydrolase iso
>gi|1036947500|ref|XP_017146273.1| PREDICTED: maternal protein tudor [Drosophila miranda]
>gi|1036947616|ref|XP_017146282.1| PREDICTED: asporin [Drosophila miranda]
>gi|1036947600|ref|XP_017146281.1| PREDICTED: uncharacterized protein LOC108158484 [Drosophila
>gi|1036947532|ref|XP_017146276.1| PREDICTED: glycogenin-1 isoform X2 [Drosophila miranda]
>gi|1036947516|ref|XP_017146274.1| PREDICTED: E1A-binding protein p400 isoform X1 [Drosophila
>gi|1036947564|ref|XP_017146278.1| PREDICTED: UDP-glucuronosyltransferase 2C1 [Drosophila mira
>gi|1036947548|ref|XP_017146277.1| PREDICTED: UDP-glucuronosyltransferase 2B13 [Drosophila mir
>gi|1036991455|ref|XP_017148922.1| PREDICTED: apoptosis-stimulating of p53 protein 2 [Drosophil
>gi|1037015598|ref|XP_017150386.1| PREDICTED: uncharacterized protein LOC108160721 isoform X1
>gi|1037015614|ref|XP_017150387.1| PREDICTED: GTP-binding protein RAD isoform X2 [Drosophila m
>gi|1037015630|ref|XP_017150388.1| PREDICTED: uncharacterized protein LOC108160722 [Drosophila
>gi|1037015644|ref|XP_017150389.1| PREDICTED: uncharacterized protein LOC108160723 [Drosophila
>gi|1036936608|ref|XP_017145605.1| PREDICTED: glucose-6-phosphate exchanger SLC37A2 [Drosophila
>gi|1036985965|ref|XP_017148599.1| PREDICTED: probable multidrug resistance-associated protein
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>gi|1036985977|ref|XP_017148600.1| PREDICTED: ceramide-1-phosphate transfer protein [Drosophila
>gi|1036940891|ref|XP_017145877.1| PREDICTED: uncharacterized protein LOC108158168 [Drosophila
>gi|1036967161|ref|XP_017147440.1| PREDICTED: SAGA-associated factor 29 [Drosophila miranda]
>gi|1036967207|ref|XP_017147443.1| PREDICTED: 60S ribosomal protein L29 [Drosophila miranda]
>gi|1036967139|ref|XP_017147439.1| PREDICTED: UPF0553 protein C9orf64 homolog [Drosophila mira
>gi|1036966936|ref|XP_017147429.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967101|ref|XP_017147437.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036966991|ref|XP_017147432.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967119|ref|XP_017147438.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967187|ref|XP_017147441.1| PREDICTED: uncharacterized protein LOC108159032 [Drosophila
>gi|1036966961|ref|XP_017147430.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967042|ref|XP_017147434.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967060|ref|XP_017147435.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967024|ref|XP_017147433.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036966975|ref|XP_017147431.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036967082|ref|XP_017147436.1| PREDICTED: kinase D-interacting substrate of 220 kDa isoform
>gi|1036968251|ref|XP_017147502.1| PREDICTED: glycogen debranching enzyme [Drosophila miranda]
>gi|1036968391|ref|XP_017147511.1| PREDICTED: uncharacterized protein LOC108159072 [Drosophila
>gi|1036968269|ref|XP_017147503.1| PREDICTED: putative sodium-coupled neutral amino acid transp
>gi|1036968283|ref|XP_017147504.1| PREDICTED: putative sodium-coupled neutral amino acid transp
>gi|1036968403|ref|XP_017147512.1| PREDICTED: 40S ribosomal protein S15 [Drosophila miranda]
>gi|1036968357|ref|XP_017147508.1| PREDICTED: serine/threonine-protein kinase SBK1 isoform X2
>gi|1036968375|ref|XP_017147509.1| PREDICTED: sperm motility kinase isoform X3 [Drosophila mira
>gi|1036968319|ref|XP_017147506.1| PREDICTED: sperm motility kinase isoform X1 [Drosophila mira
>gi|1036968339|ref|XP_017147507.1| PREDICTED: sperm motility kinase isoform X1 [Drosophila mira
>gi|1036968301|ref|XP_017147505.1| PREDICTED: sodium-dependent dopamine transporter [Drosophila
>gi|1037021406|ref|XP_017150755.1| PREDICTED: gamma-1-syntrophin [Drosophila miranda]
>gi|1036978243|ref|XP_017148116.1| PREDICTED: semaphorin-2A isoform X2 [Drosophila miranda]
>gi|1036978230|ref|XP_017148115.1| PREDICTED: semaphorin-2A isoform X1 [Drosophila miranda]
>gi|1036978256|ref|XP_017148117.1| PREDICTED: axoneme-associated protein mst101(2) [Drosophila
>gi|1036978271|ref|XP_017148118.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1036978283|ref|XP_017148119.1| PREDICTED: chymotrypsin-C-like [Drosophila miranda]
>gi|1036978301|ref|XP_017148120.1| PREDICTED: uncharacterized protein LOC108159396 [Drosophila
>gi|1036941079|ref|XP_017145884.1| PREDICTED: accessory gland protein Acp29AB-like [Drosophila
>gi|1037022358|ref|XP_017150812.1| PREDICTED: proton-coupled amino acid transporter 1-like iso
>gi|1037022394|ref|XP_017150815.1| PREDICTED: proton-coupled amino acid transporter 1-like iso
>gi|1037022376|ref|XP_017150814.1| PREDICTED: proton-coupled amino acid transporter 1-like iso
>gi|1036942259|ref|XP_017145957.1| PREDICTED: uncharacterized protein LOC108158236 [Drosophila
>gi|1037002670|ref|XP_017149611.1| PREDICTED: uncharacterized protein LOC108160256 [Drosophila
>gi|1037002652|ref|XP_017149610.1| PREDICTED: NADP-dependent malic enzyme [Drosophila miranda]
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>gi|1037002846|ref|XP_017149614.1| PREDICTED: semaphorin-2A isoform X2 [Drosophila miranda]
>gi|1037002808|ref|XP_017149613.1| PREDICTED: semaphorin-2A isoform X1 [Drosophila miranda]
>gi|1037002864|ref|XP_017149615.1| PREDICTED: semaphorin-2A isoform X3 [Drosophila miranda]
>gi|1037002881|ref|XP_017149617.1| PREDICTED: NADP-dependent malic enzyme [Drosophila miranda]
>gi|1037002896|ref|XP_017149618.1| PREDICTED: NADP-dependent malic enzyme-like [Drosophila mira
>gi|1036943076|ref|XP_017146009.1| PREDICTED: uncharacterized protein LOC108158282 isoform X2
>gi|1036943040|ref|XP_017146007.1| PREDICTED: uncharacterized protein LOC108158282 isoform X1
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>gi|1036943058|ref|XP_017146008.1| PREDICTED: uncharacterized protein LOC108158282 isoform X1
>gi|1037015348|ref|XP_017150375.1| PREDICTED: uncharacterized protein LOC108160713 isoform X1
>gi|1037015382|ref|XP_017150377.1| PREDICTED: uncharacterized protein LOC108160713 isoform X3
>gi|1037015366|ref|XP_017150376.1| PREDICTED: uncharacterized protein LOC108160713 isoform X2
>gi|1036942439|ref|XP_017145969.1| PREDICTED: uncharacterized protein LOC108158246 [Drosophila
>gi|1036936626|ref|XP_017145607.1| PREDICTED: uncharacterized protein LOC108157972 [Drosophila
>gi|1036957478|ref|XP_017146877.1| PREDICTED: U3 small nucleolar RNA-associated protein 18 hom
>gi|1036957604|ref|XP_017146885.1| PREDICTED: nuclear pore glycoprotein p62 [Drosophila miranda
>gi|1036957538|ref|XP_017146881.1| PREDICTED: alpha-N-acetylgalactosaminidase [Drosophila miran
>gi|1036957560|ref|XP_017146883.1| PREDICTED: alpha-N-acetylgalactosaminidase [Drosophila miran
>gi|1036957574|ref|XP_017146884.1| PREDICTED: alpha-N-acetylgalactosaminidase [Drosophila miran
>gi|1036957490|ref|XP_017146878.1| PREDICTED: hydroxymethylglutaryl-CoA synthase 1 [Drosophila
>gi|1036957506|ref|XP_017146879.1| PREDICTED: hydroxymethylglutaryl-CoA synthase 1 [Drosophila
>gi|1036957523|ref|XP_017146880.1| PREDICTED: hydroxymethylglutaryl-CoA synthase 1 [Drosophila
>gi|1036957446|ref|XP_017146875.1| PREDICTED: V-type proton ATPase subunit C isoform X3 [Drosop
>gi|1036957464|ref|XP_017146876.1| PREDICTED: V-type proton ATPase subunit C isoform X4 [Drosop
>gi|1036957400|ref|XP_017146873.1| PREDICTED: V-type proton ATPase subunit C isoform X1 [Drosop
>gi|1036957417|ref|XP_017146874.1| PREDICTED: V-type proton ATPase subunit C isoform X2 [Drosop
>gi|1036957383|ref|XP_017146872.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036957620|ref|XP_017146886.1| PREDICTED: uncharacterized protein LOC108158770 [Drosophila
>gi|1036999247|ref|XP_017149405.1| PREDICTED: inhibitor of Bruton tyrosine kinase [Drosophila
>gi|1036999229|ref|XP_017149404.1| PREDICTED: NADPH oxidase 5 isoform X2 [Drosophila miranda]
>gi|1036999195|ref|XP_017149403.1| PREDICTED: uncharacterized protein LOC108160121 isoform X1
>gi|1036970381|ref|XP_017147630.1| PREDICTED: autophagy-related protein 9A [Drosophila miranda]
>gi|1036970399|ref|XP_017147631.1| PREDICTED: peptidyl-prolyl cis-trans isomerase-like 2 [Dros
>gi|1036970526|ref|XP_017147639.1| PREDICTED: probable peptide chain release factor C12orf65 h
>gi|1036970545|ref|XP_017147641.1| PREDICTED: uncharacterized protein LOC108159154 [Drosophila
>gi|1036970435|ref|XP_017147634.1| PREDICTED: uncharacterized protein LOC108159148 [Drosophila
>gi|1036970417|ref|XP_017147633.1| PREDICTED: uncharacterized protein LOC108159147 [Drosophila
>gi|1036970490|ref|XP_017147637.1| PREDICTED: uncharacterized protein LOC108159151 [Drosophila
>gi|1036970508|ref|XP_017147638.1| PREDICTED: uncharacterized protein LOC108159152 [Drosophila
>gi|1036970453|ref|XP_017147635.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit gamma, mit
>gi|1036970564|ref|XP_017147642.1| PREDICTED: uncharacterized protein LOC108159155 [Drosophila
>gi|1036970473|ref|XP_017147636.1| PREDICTED: protein atonal homolog 1 [Drosophila miranda]
>gi|1036970343|ref|XP_017147628.1| PREDICTED: uncharacterized protein LOC108159143 isoform X1
>gi|1036970361|ref|XP_017147629.1| PREDICTED: uncharacterized protein LOC108159143 isoform X2
>gi|1036936644|ref|XP_017145608.1| PREDICTED: uncharacterized protein LOC108157973 [Drosophila
>gi|1036975020|ref|XP_017147917.1| PREDICTED: kinesin heavy chain [Drosophila miranda]
>gi|1036975081|ref|XP_017147919.1| PREDICTED: LOW QUALITY PROTEIN: protein arginine N-methyltra
>gi|1036975061|ref|XP_017147918.1| PREDICTED: coiled-coil domain-containing protein 93 [Drosoph
>gi|1036936662|ref|XP_017145609.1| PREDICTED: coiled-coil domain-containing protein 93-like [Dr
>gi|1037019511|ref|XP_017150634.1| PREDICTED: cyclic nucleotide-gated cation channel subunit A
>gi|1037019528|ref|XP_017150636.1| PREDICTED: cyclic nucleotide-gated cation channel subunit A
>gi|1037019545|ref|XP_017150637.1| PREDICTED: cyclic nucleotide-gated cation channel subunit A
>gi|1037000621|ref|XP_017149485.1| PREDICTED: uncharacterized protein LOC108160172 [Drosophila
>gi|1037000666|ref|XP_017149488.1| PREDICTED: uncharacterized protein LOC108160175 [Drosophila
>gi|1037000803|ref|XP_017149499.1| PREDICTED: uncharacterized protein LOC108160183 [Drosophila
>gi|1037000648|ref|XP_017149487.1| PREDICTED: uncharacterized protein LOC108160174 [Drosophila
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>gi|1037000612|ref|XP_017149484.1| PREDICTED: probable cytochrome P450 6d2 [Drosophila miranda]
>gi|1037000697|ref|XP_017149490.1| PREDICTED: uncharacterized protein LOC108160178 [Drosophila
>gi|1037000785|ref|XP_017149498.1| PREDICTED: uncharacterized protein LOC108160182 isoform X2
>gi|1037000768|ref|XP_017149497.1| PREDICTED: uncharacterized protein LOC108160182 isoform X1
>gi|1037000683|ref|XP_017149489.1| PREDICTED: uncharacterized protein LOC108160176 [Drosophila
>gi|1037000754|ref|XP_017149496.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037000740|ref|XP_017149495.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037000710|ref|XP_017149491.1| PREDICTED: cell death-inducing p53-target protein 1 homolog
>gi|1037000724|ref|XP_017149492.1| PREDICTED: cell death-inducing p53-target protein 1 homolog
>gi|1037009442|ref|XP_017150021.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1036936680|ref|XP_017145610.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037009424|ref|XP_017150020.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037009406|ref|XP_017150019.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037009389|ref|XP_017150018.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037009372|ref|XP_017150017.1| PREDICTED: lipopolysaccharide-induced tumor necrosis factor-
>gi|1037013368|ref|XP_017150253.1| PREDICTED: YY1-associated factor 2 [Drosophila miranda]
>gi|1036950673|ref|XP_017146475.1| PREDICTED: F-box/LRR-repeat protein 14 [Drosophila miranda]
>gi|1036950844|ref|XP_017146486.1| PREDICTED: roundabout homolog 2 isoform X3 [Drosophila miranda]
>gi|1036950831|ref|XP_017146485.1| PREDICTED: protein sax-3 isoform X2 [Drosophila miranda]
>gi|1036950781|ref|XP_017146482.1| PREDICTED: roundabout homolog 2 isoform X1 [Drosophila miranda]
>gi|1036950797|ref|XP_017146483.1| PREDICTED: roundabout homolog 2 isoform X1 [Drosophila miranda]
>gi|1036950815|ref|XP_017146484.1| PREDICTED: roundabout homolog 2 isoform X1 [Drosophila miranda]
>gi|1036950928|ref|XP_017146491.1| PREDICTED: coiled-coil domain-containing protein 65 [Drosophila
>gi|1036950946|ref|XP_017146492.1| PREDICTED: general odorant-binding protein 71 [Drosophila miranda]
>gi|1036950981|ref|XP_017146494.1| PREDICTED: general odorant-binding protein 68 [Drosophila miranda]
>gi|1036950999|ref|XP_017146496.1| PREDICTED: uncharacterized protein LOC108158571 [Drosophila
>gi|1036950964|ref|XP_017146493.1| PREDICTED: uncharacterized protein LOC108158569 [Drosophila
>gi|1036950859|ref|XP_017146488.1| PREDICTED: cilia- and flagella-associated protein 61 isoform
>gi|1036950877|ref|XP_017146489.1| PREDICTED: cilia- and flagella-associated protein 61 isoform
>gi|1036950910|ref|XP_017146490.1| PREDICTED: cilia- and flagella-associated protein 61 [Drosophila
>gi|1036950728|ref|XP_017146479.1| PREDICTED: filamin-B isoform X2 [Drosophila miranda]
>gi|1036950692|ref|XP_017146476.1| PREDICTED: filamin-B isoform X1 [Drosophila miranda]
>gi|1036950710|ref|XP_017146478.1| PREDICTED: filamin-B isoform X1 [Drosophila miranda]
>gi|1036950747|ref|XP_017146480.1| PREDICTED: filamin-C isoform X3 [Drosophila miranda]
>gi|1036950764|ref|XP_017146481.1| PREDICTED: filamin-A isoform X4 [Drosophila miranda]
>gi|1037013566|ref|XP_017150265.1| PREDICTED: cyclic nucleotide-gated cation channel beta-1 iso
>gi|1037013548|ref|XP_017150264.1| PREDICTED: cyclic nucleotide-gated cation channel beta-1 iso
>gi|1037013584|ref|XP_017150266.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037013636|ref|XP_017150269.1| PREDICTED: uncharacterized protein LOC108160648 [Drosophila
>gi|1037013600|ref|XP_017150267.1| PREDICTED: uncharacterized protein LOC108160645 [Drosophila
>gi|1037013618|ref|XP_017150268.1| PREDICTED: uncharacterized protein LOC108160646 [Drosophila
>gi|1037006159|ref|XP_017149819.1| PREDICTED: ATP synthase subunit alpha, mitochondrial [Drosop
>gi|1037001018|ref|XP_017149507.1| PREDICTED: G2/mitotic-specific cyclin-B isoform X2 [Drosophi
>gi|1037001000|ref|XP_017149506.1| PREDICTED: G2/mitotic-specific cyclin-B isoform X1 [Drosophi
>gi|1037018218|ref|XP_017150549.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1037018229|ref|XP_017150550.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1037018242|ref|XP_017150551.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
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>gi|1037018965|ref|XP_017150601.1| PREDICTED: uncharacterized protein LOC108160854 isoform X1
>gi|1037018981|ref|XP_017150602.1| PREDICTED: high mobility group nucleosome-binding domain-co
>gi|1037019012|ref|XP_017150604.1| PREDICTED: uncharacterized protein LOC108160854 isoform X4
>gi|1037018995|ref|XP_017150603.1| PREDICTED: uncharacterized protein LOC108160854 isoform X3
>gi|1037019029|ref|XP_017150605.1| PREDICTED: arginine kinase-like isoform X5 [Drosophila mira
>gi|1037016835|ref|XP_017150459.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036941804|ref|XP_017145926.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036990407|ref|XP_017148868.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036990424|ref|XP_017148869.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036990391|ref|XP_017148867.1| PREDICTED: spatacsin [Drosophila miranda]
>gi|1036982431|ref|XP_017148378.1| PREDICTED: protein Shroom isoform X1 [Drosophila miranda]
>gi|1036982445|ref|XP_017148379.1| PREDICTED: protein Shroom isoform X2 [Drosophila miranda]
>gi|1036982459|ref|XP_017148380.1| PREDICTED: protein Shroom isoform X3 [Drosophila miranda]
>gi|1037014626|ref|XP_017150330.1| PREDICTED: PRKR-interacting protein 1 homolog [Drosophila m
>gi|1036959468|ref|XP_017146994.1| PREDICTED: lissencephaly-1 homolog [Drosophila miranda]
>gi|1036959584|ref|XP_017147000.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1036959514|ref|XP_017146995.1| PREDICTED: phosphatidylinositol phosphatase PTPRQ [Drosophi
>gi|1036959530|ref|XP_017146996.1| PREDICTED: phosphatidylinositol phosphatase PTPRQ [Drosophi
>gi|1036959548|ref|XP_017146997.1| PREDICTED: phosphatidylinositol phosphatase PTPRQ [Drosophi
>gi|1036959602|ref|XP_017147001.1| PREDICTED: coiled-coil domain-containing protein 94 [Drosop
>gi|1036959566|ref|XP_017146998.1| PREDICTED: leucine-rich repeats and immunoglobulin-like dom
>gi|1036984261|ref|XP_017148491.1| PREDICTED: ATP synthase subunit s, mitochondrial [Drosophila
>gi|1036984227|ref|XP_017148489.1| PREDICTED: exostosin-2 [Drosophila miranda]
>gi|1036984211|ref|XP_017148488.1| PREDICTED: cysteine--tRNA ligase, cytoplasmic [Drosophila m
>gi|1036984245|ref|XP_017148490.1| PREDICTED: alpha N-terminal protein methyltransferase 1 [Dr
>gi|1036984279|ref|XP_017148492.1| PREDICTED: uncharacterized protein LOC108159566 [Drosophila
>gi|1037017270|ref|XP_017150489.1| PREDICTED: zinc finger protein 423 homolog isoform X1 [Dros
>gi|1037017289|ref|XP_017150490.1| PREDICTED: zinc finger protein 423 homolog isoform X2 [Dros
>gi|1037017302|ref|XP_017150491.1| PREDICTED: zinc finger protein 423 homolog isoform X3 [Dros
>gi|1037008679|ref|XP_017149974.1| PREDICTED: protein phyllopod [Drosophila miranda]
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>gi|1037006971|ref|XP_017149872.1| PREDICTED: stress-activated map kinase-interacting protein
>gi|1037006987|ref|XP_017149873.1| PREDICTED: zinc finger protein 271 [Drosophila miranda]
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>gi|1037006706|ref|XP_017149854.1| PREDICTED: uncharacterized protein LOC108160384 isoform X2
>gi|1037006688|ref|XP_017149853.1| PREDICTED: uncharacterized protein LOC108160384 isoform X1
>gi|1037006658|ref|XP_017149851.1| PREDICTED: uncharacterized protein LOC108160383 isoform X1
>gi|1037006674|ref|XP_017149852.1| PREDICTED: uncharacterized protein LOC108160383 isoform X2
>gi|1037006607|ref|XP_017149848.1| PREDICTED: general transcription factor IIH subunit 1 [Dros
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>gi|1037006622|ref|XP_017149849.1| PREDICTED: general transcription factor IIH subunit 1 [Dros
>gi|1037006640|ref|XP_017149850.1| PREDICTED: general transcription factor IIH subunit 1 [Dros
>gi|1037017648|ref|XP_017150512.1| PREDICTED: uncharacterized protein LOC108160802 [Drosophila
>gi|1036943024|ref|XP_017146005.1| PREDICTED: uncharacterized protein LOC108158281 [Drosophila
>gi|1037004746|ref|XP_017149727.1| PREDICTED: uncharacterized protein LOC108160315 isoform X1
>gi|1037004766|ref|XP_017149728.1| PREDICTED: uncharacterized protein LOC108160315 isoform X2
>gi|1037004714|ref|XP_017149724.1| PREDICTED: spermatogenesis-associated protein 20 isoform X1
>gi|1037004732|ref|XP_017149726.1| PREDICTED: spermatogenesis-associated protein 20 isoform X2
>gi|1036979919|ref|XP_017148222.1| PREDICTED: protein clueless isoform X1 [Drosophila miranda]
>gi|1036979935|ref|XP_017148224.1| PREDICTED: protein clueless isoform X2 [Drosophila miranda]
>gi|1036979950|ref|XP_017148225.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase calypso [Dr
>gi|1036979962|ref|XP_017148226.1| PREDICTED: uncharacterized protein LOC108159447 isoform X1
>gi|1036979977|ref|XP_017148227.1| PREDICTED: uncharacterized protein LOC108159447 isoform X1
>gi|1036979990|ref|XP_017148228.1| PREDICTED: uncharacterized protein LOC108159447 isoform X2
>gi|1036991930|ref|XP_017148952.1| PREDICTED: uncharacterized protein LOC108159862 [Drosophila
>gi|1036991912|ref|XP_017148949.1| PREDICTED: lipoyltransferase 1, mitochondrial [Drosophila mi
>gi|1036991896|ref|XP_017148948.1| PREDICTED: tyrosine-protein kinase shark [Drosophila miranda
>gi|1036995322|ref|XP_017149164.1| PREDICTED: dnaJ homolog subfamily B member 6 isoform X1 [Dro
>gi|1036995340|ref|XP_017149165.1| PREDICTED: dnaJ homolog subfamily B member 6 isoform X1 [Dro
>gi|1036995354|ref|XP_017149166.1| PREDICTED: dnaJ homolog subfamily B member 6 isoform X1 [Dro
>gi|1036995370|ref|XP_017149167.1| PREDICTED: dnaJ homolog subfamily B member 6 isoform X2 [Dro
>gi|1036995304|ref|XP_017149162.1| PREDICTED: uncharacterized protein LOC108159980 [Drosophila
>gi|1036995294|ref|XP_017149161.1| PREDICTED: homeobox protein HMX3 [Drosophila miranda]
>gi|1037010995|ref|XP_017150107.1| PREDICTED: aquaporin [Drosophila miranda]
>gi|1037013891|ref|XP_017150284.1| PREDICTED: aquaporin AQPcic [Drosophila miranda]
>gi|1037016044|ref|XP_017150411.1| PREDICTED: protein quiver isoform X2 [Drosophila miranda]
>gi|1037016026|ref|XP_017150410.1| PREDICTED: protein quiver isoform X1 [Drosophila miranda]
>gi|1037016062|ref|XP_017150412.1| PREDICTED: protein quiver isoform X3 [Drosophila miranda]
>gi|1037016078|ref|XP_017150413.1| PREDICTED: protein quiver isoform X4 [Drosophila miranda]
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>gi|1037012734|ref|XP_017150214.1| PREDICTED: homeobox protein invected [Drosophila miranda]
>gi|1037004676|ref|XP_017149722.1| PREDICTED: segmentation polarity homeobox protein engrailed
>gi|1037004696|ref|XP_017149723.1| PREDICTED: serine/threonine-protein kinase polo [Drosophila
>gi|1036965763|ref|XP_017147361.1| PREDICTED: bromodomain adjacent to zinc finger domain prote
>gi|1036965651|ref|XP_017147355.1| PREDICTED: uncharacterized protein LOC108159002 isoform X1
>gi|1036965671|ref|XP_017147356.1| PREDICTED: uncharacterized protein LOC108159002 isoform X1
>gi|1036965689|ref|XP_017147357.1| PREDICTED: uncharacterized protein LOC108159002 isoform X1
>gi|1036965707|ref|XP_017147358.1| PREDICTED: uncharacterized protein LOC108159002 isoform X1
>gi|1036965725|ref|XP_017147359.1| PREDICTED: uncharacterized protein LOC108159002 isoform X1
>gi|1036965743|ref|XP_017147360.1| PREDICTED: uncharacterized protein LOC108159002 isoform X1
>gi|1037011507|ref|XP_017150138.1| PREDICTED: acyl-CoA dehydrogenase family member 9, mitochond
>gi|1036987482|ref|XP_017148690.1| PREDICTED: protein FAM214A [Drosophila miranda]
>gi|1036987500|ref|XP_017148691.1| PREDICTED: protein FAM214A [Drosophila miranda]
>gi|1036998489|ref|XP_017149362.1| PREDICTED: F-box/LRR-repeat protein 20 isoform X3 [Drosophil
>gi|1036998507|ref|XP_017149363.1| PREDICTED: F-box/LRR-repeat protein 20 isoform X4 [Drosophil
>gi|1036998471|ref|XP_017149361.1| PREDICTED: F-box/LRR-repeat protein 20 isoform X2 [Drosophil
>gi|1036998455|ref|XP_017149360.1| PREDICTED: F-box/LRR-repeat protein 20 isoform X1 [Drosophil
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>gi|1036996910|ref|XP_017149262.1| PREDICTED: small integral membrane protein 12-A [Drosophila
>gi|1036996876|ref|XP_017149260.1| PREDICTED: RING-box protein 2 isoform X1 [Drosophila miranda
>gi|1036996928|ref|XP_017149264.1| PREDICTED: uncharacterized protein LOC108160034 [Drosophila
>gi|1036996893|ref|XP_017149261.1| PREDICTED: RING-box protein 2 isoform X2 [Drosophila miranda
>gi|1036996845|ref|XP_017149258.1| PREDICTED: facilitated trehalose transporter Tret1 isoform X
>gi|1036996861|ref|XP_017149259.1| PREDICTED: facilitated trehalose transporter Tret1 isoform X
>gi|1036943476|ref|XP_017146033.1| PREDICTED: uncharacterized protein LOC108158304 [Drosophila
>gi|1037019185|ref|XP_017150615.1| PREDICTED: probable serine/threonine-protein kinase kinX [Dr
>gi|1036988133|ref|XP_017148727.1| PREDICTED: proteasome subunit alpha type-5 [Drosophila miran
>gi|1036988151|ref|XP_017148728.1| PREDICTED: proteasome subunit alpha type-5 [Drosophila miran
>gi|1036988119|ref|XP_017148726.1| PREDICTED: uncharacterized protein LOC108159717 [Drosophila
>gi|1036994567|ref|XP_017149117.1| PREDICTED: uridine-cytidine kinase-like 1 isoform X3 [Drosop
>gi|1036994552|ref|XP_017149116.1| PREDICTED: uridine-cytidine kinase-like 1 isoform X2 [Drosop
>gi|1036994534|ref|XP_017149114.1| PREDICTED: uridine-cytidine kinase-like 1 isoform X1 [Drosop
>gi|1036994588|ref|XP_017149118.1| PREDICTED: threonylcarbamoyladenosine tRNA methylthiotransfe
>gi|1036975709|ref|XP_017147958.1| PREDICTED: S-methyl-5'-thioadenosine phosphorylase [Drosophi
>gi|1036975673|ref|XP_017147955.1| PREDICTED: actin-like protein 6B [Drosophila miranda]
>gi|1036975691|ref|XP_017147956.1| PREDICTED: uncharacterized protein LOC108159298 [Drosophila
>gi|1036975655|ref|XP_017147954.1| PREDICTED: myotubularin-related protein 14 [Drosophila miran
>gi|1036975727|ref|XP_017147959.1| PREDICTED: glucose-induced degradation protein 8 homolog [Dr
>gi|1037019729|ref|XP_017150649.1| PREDICTED: dynein light chain roadblock-type 1 [Drosophila m
>gi|1037019766|ref|XP_017150650.1| PREDICTED: dynein light chain roadblock-type 2 [Drosophila m
>gi|1036965964|ref|XP_017147373.1| PREDICTED: protein MTL1 [Drosophila miranda]
>gi|1036965944|ref|XP_017147372.1| PREDICTED: uncharacterized protein LOC108159009 [Drosophila
>gi|1037006256|ref|XP_017149825.1| PREDICTED: LOW QUALITY PROTEIN: probable WRKY transcription
>gi|1037003094|ref|XP_017149632.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1037003033|ref|XP_017149628.1| PREDICTED: protein 5NUC [Drosophila miranda]
>gi|1037003050|ref|XP_017149629.1| PREDICTED: protein 5NUC [Drosophila miranda]
>gi|1037003064|ref|XP_017149630.1| PREDICTED: protein 5NUC [Drosophila miranda]
>gi|1037003077|ref|XP_017149631.1| PREDICTED: protein 5NUC [Drosophila miranda]
>gi|1037012648|ref|XP_017150209.1| PREDICTED: uncharacterized protein LOC108160609 isoform X1
>gi|1037012666|ref|XP_017150210.1| PREDICTED: uncharacterized protein LOC108160609 isoform X2
>gi|1037012698|ref|XP_017150212.1| PREDICTED: uncharacterized protein LOC108160609 isoform X4
>gi|1037012684|ref|XP_017150211.1| PREDICTED: serine-aspartate repeat-containing protein I iso
>gi|1036943973|ref|XP_017146063.1| PREDICTED: uncharacterized protein LOC108158334 [Drosophila
>gi|1037020613|ref|XP_017150705.1| PREDICTED: cathepsin L1 isoform X1 [Drosophila miranda]
>gi|1037020627|ref|XP_017150706.1| PREDICTED: cathepsin L1 isoform X2 [Drosophila miranda]
>gi|1036966718|ref|XP_017147419.1| PREDICTED: ras-GEF domain-containing family member 1B-B [Dro
>gi|1036966737|ref|XP_017147420.1| PREDICTED: LOW QUALITY PROTEIN: anaphase-promoting complex s
>gi|1036966755|ref|XP_017147421.1| PREDICTED: U3 small nucleolar ribonucleoprotein protein IMP3
>gi|1036966700|ref|XP_017147418.1| PREDICTED: patronin isoform X37 [Drosophila miranda]
>gi|1036966676|ref|XP_017147417.1| PREDICTED: patronin isoform X36 [Drosophila miranda]
>gi|1036966640|ref|XP_017147415.1| PREDICTED: patronin isoform X34 [Drosophila miranda]
>gi|1036966450|ref|XP_017147402.1| PREDICTED: patronin isoform X22 [Drosophila miranda]
>gi|1036966626|ref|XP_017147414.1| PREDICTED: patronin isoform X33 [Drosophila miranda]
>gi|1036966658|ref|XP_017147416.1| PREDICTED: patronin isoform X35 [Drosophila miranda]
>gi|1036966285|ref|XP_017147393.1| PREDICTED: patronin isoform X13 [Drosophila miranda]
>gi|1036966267|ref|XP_017147392.1| PREDICTED: patronin isoform X12 [Drosophila miranda]
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>gi|1036966612|ref|XP_017147413.1| PREDICTED: patronin isoform X32 [Drosophila miranda]
>gi|1036966465|ref|XP_017147403.1| PREDICTED: patronin isoform X23 [Drosophila miranda]
>gi|1036966557|ref|XP_017147410.1| PREDICTED: patronin isoform X29 [Drosophila miranda]
>gi|1036966340|ref|XP_017147396.1| PREDICTED: patronin isoform X16 [Drosophila miranda]
>gi|1036966376|ref|XP_017147398.1| PREDICTED: patronin isoform X18 [Drosophila miranda]
>gi|1036966358|ref|XP_017147397.1| PREDICTED: patronin isoform X17 [Drosophila miranda]
>gi|1036966144|ref|XP_017147384.1| PREDICTED: patronin isoform X5 [Drosophila miranda]
>gi|1036966126|ref|XP_017147383.1| PREDICTED: patronin isoform X4 [Drosophila miranda]
>gi|1036966396|ref|XP_017147399.1| PREDICTED: patronin isoform X19 [Drosophila miranda]
>gi|1036966253|ref|XP_017147390.1| PREDICTED: patronin isoform X11 [Drosophila miranda]
>gi|1036966090|ref|XP_017147381.1| PREDICTED: patronin isoform X2 [Drosophila miranda]
>gi|1036966596|ref|XP_017147412.1| PREDICTED: patronin isoform X31 [Drosophila miranda]
>gi|1036966578|ref|XP_017147411.1| PREDICTED: patronin isoform X30 [Drosophila miranda]
>gi|1036966431|ref|XP_017147401.1| PREDICTED: patronin isoform X21 [Drosophila miranda]
>gi|1036966542|ref|XP_017147409.1| PREDICTED: patronin isoform X28 [Drosophila miranda]
>gi|1036966414|ref|XP_017147400.1| PREDICTED: patronin isoform X20 [Drosophila miranda]
>gi|1036966235|ref|XP_017147389.1| PREDICTED: patronin isoform X10 [Drosophila miranda]
>gi|1036966529|ref|XP_017147408.1| PREDICTED: patronin isoform X27 [Drosophila miranda]
>gi|1036966179|ref|XP_017147386.1| PREDICTED: patronin isoform X7 [Drosophila miranda]
>gi|1036966516|ref|XP_017147406.1| PREDICTED: patronin isoform X26 [Drosophila miranda]
>gi|1036966498|ref|XP_017147405.1| PREDICTED: patronin isoform X25 [Drosophila miranda]
>gi|1036966306|ref|XP_017147394.1| PREDICTED: patronin isoform X14 [Drosophila miranda]
>gi|1036966161|ref|XP_017147385.1| PREDICTED: patronin isoform X6 [Drosophila miranda]
>gi|1036966482|ref|XP_017147404.1| PREDICTED: patronin isoform X24 [Drosophila miranda]
>gi|1036966323|ref|XP_017147395.1| PREDICTED: patronin isoform X15 [Drosophila miranda]
>gi|1036966217|ref|XP_017147388.1| PREDICTED: patronin isoform X9 [Drosophila miranda]
>gi|1036966108|ref|XP_017147382.1| PREDICTED: patronin isoform X3 [Drosophila miranda]
>gi|1036966199|ref|XP_017147387.1| PREDICTED: patronin isoform X8 [Drosophila miranda]
>gi|1036966072|ref|XP_017147380.1| PREDICTED: patronin isoform X1 [Drosophila miranda]
>gi|1037001527|ref|XP_017149540.1| PREDICTED: uncharacterized protein LOC108160208 isoform X1
>gi|1037001547|ref|XP_017149541.1| PREDICTED: uncharacterized protein LOC108160208 isoform X2
>gi|1037001473|ref|XP_017149537.1| PREDICTED: transcription factor collier isoform X6 [Drosophi
>gi|1037001439|ref|XP_017149534.1| PREDICTED: transcription factor collier isoform X4 [Drosophi
>gi|1037001509|ref|XP_017149539.1| PREDICTED: transcription factor collier isoform X8 [Drosophi
>gi|1037001425|ref|XP_017149533.1| PREDICTED: transcription factor collier isoform X3 [Drosophi
>gi|1037001455|ref|XP_017149536.1| PREDICTED: transcription factor collier isoform X5 [Drosophi
>gi|1037001413|ref|XP_017149532.1| PREDICTED: transcription factor collier isoform X2 [Drosophi
>gi|1037001397|ref|XP_017149531.1| PREDICTED: transcription factor collier isoform X1 [Drosophi
>gi|1037001491|ref|XP_017149538.1| PREDICTED: transcription factor collier isoform X7 [Drosophi
>gi|1036943164|ref|XP_017146014.1| PREDICTED: extracellular glycosidase CRH11 [Drosophila mirar
>gi|1037021286|ref|XP_017150747.1| PREDICTED: phosphoglucomutase-2 [Drosophila miranda]
>gi|1036941837|ref|XP_017145928.1| PREDICTED: uncharacterized protein LOC108158216 isoform X1
>gi|1036941853|ref|XP_017145929.1| PREDICTED: uncharacterized protein LOC108158216 isoform X2
>gi|1036974123|ref|XP_017147860.1| PREDICTED: sialin [Drosophila miranda]
>gi|1036974141|ref|XP_017147861.1| PREDICTED: hormone receptor 4 [Drosophila miranda]
>gi|1036974105|ref|XP_017147859.1| PREDICTED: structural maintenance of chromosomes protein 2
>gi|1037004923|ref|XP_017149738.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037004939|ref|XP_017149739.1| PREDICTED: actin-related protein 4-like isoform X1 [Drosophi
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>gi|1037004973|ref|XP_017149741.1| PREDICTED: actin-like protein 6B isoform X3 [Drosophila mira
>gi|1037004957|ref|XP_017149740.1| PREDICTED: actin-related protein 4-like isoform X2 [Drosophi
>gi|1037004990|ref|XP_017149742.1| PREDICTED: actin-related protein 4-like isoform X1 [Drosophi
>gi|1037005026|ref|XP_017149744.1| PREDICTED: actin-like protein 6B isoform X3 [Drosophila mira
>gi|1037005008|ref|XP_017149743.1| PREDICTED: actin-related protein 4-like isoform X2 [Drosophi
>gi|1036970103|ref|XP_017147614.1| PREDICTED: uncharacterized protein LOC108159131 [Drosophila
>gi|1036970124|ref|XP_017147615.1| PREDICTED: uncharacterized protein LOC108159132 [Drosophila
>gi|1036970162|ref|XP_017147617.1| PREDICTED: uncharacterized protein LOC108159134 [Drosophila
>gi|1036970180|ref|XP_017147618.1| PREDICTED: ATPase inhibitor A, mitochondrial [Drosophila mi
>gi|1036970144|ref|XP_017147616.1| PREDICTED: 5-formyltetrahydrofolate cyclo-ligase [Drosophila
>gi|1036970069|ref|XP_017147611.1| PREDICTED: protein virilizer isoform X1 [Drosophila miranda]
>gi|1036970085|ref|XP_017147612.1| PREDICTED: protein virilizer isoform X2 [Drosophila miranda]
>gi|1036975481|ref|XP_017147945.1| PREDICTED: ataxin-2 homolog [Drosophila miranda]
>gi|1036975447|ref|XP_017147942.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036975465|ref|XP_017147943.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036987518|ref|XP_017148692.1| PREDICTED: polycomb protein Asx [Drosophila miranda]
>gi|1037020454|ref|XP_017150694.1| PREDICTED: cuticle protein 21 [Drosophila miranda]
>gi|1037020483|ref|XP_017150695.1| PREDICTED: uncharacterized protein LOC108160923 [Drosophila
>gi|1036941663|ref|XP_017145917.1| PREDICTED: facilitated trehalose transporter Tret1-like [Dro
>gi|1036986476|ref|XP_017148628.1| PREDICTED: exostosin-1 [Drosophila miranda]
>gi|1036986530|ref|XP_017148631.1| PREDICTED: uncharacterized protein LOC108159664 [Drosophila
>gi|1036986510|ref|XP_017148630.1| PREDICTED: lamin-C isoform X2 [Drosophila miranda]
>gi|1036986492|ref|XP_017148629.1| PREDICTED: lamin-C isoform X1 [Drosophila miranda]
>gi|1037022113|ref|XP_017150798.1| PREDICTED: cocaine esterase [Drosophila miranda]
>gi|1036950118|ref|XP_017146439.1| PREDICTED: DNA-directed RNA polymerase I subunit RPA1 [Droso
>gi|1036950188|ref|XP_017146443.1| PREDICTED: biogenesis of lysosome-related organelles comple
>gi|1036936745|ref|XP_017145614.1| PREDICTED: transformer-2 sex-determining protein [Drosophila
>gi|1036950136|ref|XP_017146440.1| PREDICTED: zinc finger CCHC domain-containing protein 4 [Dr
>gi|1036950156|ref|XP_017146441.1| PREDICTED: uncharacterized protein LOC108158543 [Drosophila
>gi|1036950172|ref|XP_017146442.1| PREDICTED: uncharacterized protein LOC108158544 [Drosophila
>gi|1036950102|ref|XP_017146438.1| PREDICTED: titin isoform X2 [Drosophila miranda]
>gi|1036950084|ref|XP_017146437.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1037016197|ref|XP_017150420.1| PREDICTED: protein transport protein Sec61 subunit beta [Dro
>gi|1036942597|ref|XP_017145978.1| PREDICTED: lambda-crystallin homolog [Drosophila miranda]
>gi|1037022324|ref|XP_017150810.1| PREDICTED: adenosine deaminase CECR1 [Drosophila miranda]
>gi|1037021720|ref|XP_017150774.1| PREDICTED: uncharacterized protein LOC108160982 [Drosophila
>gi|1037021702|ref|XP_017150773.1| PREDICTED: uncharacterized protein LOC108160981 [Drosophila
>gi|1037013722|ref|XP_017150275.1| PREDICTED: uncharacterized protein LOC108160652 isoform X1
>gi|1037013755|ref|XP_017150277.1| PREDICTED: uncharacterized protein LOC108160652 isoform X3
>gi|1037013738|ref|XP_017150276.1| PREDICTED: uncharacterized protein LOC108160652 isoform X2
>gi|1037013770|ref|XP_017150278.1| PREDICTED: uncharacterized protein LOC108160652 isoform X4
>gi|1037013808|ref|XP_017150280.1| PREDICTED: uncharacterized protein LOC108160655 [Drosophila
>gi|1037013826|ref|XP_017150282.1| PREDICTED: formin-like protein 5 [Drosophila miranda]
>gi|1037013788|ref|XP_017150279.1| PREDICTED: proline-rich protein 36 [Drosophila miranda]
>gi|1037019201|ref|XP_017150616.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1037021525|ref|XP_017150759.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1037021537|ref|XP_017150760.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1037021553|ref|XP_017150761.1| PREDICTED: prominin-like protein isoform X3 [Drosophila mira
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>gi|1036994139|ref|XP_017149089.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036994155|ref|XP_017149090.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036994175|ref|XP_017149091.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036994271|ref|XP_017149097.1| PREDICTED: drosocin [Drosophila miranda]
>gi|1036994239|ref|XP_017149095.1| PREDICTED: attacin-A-like [Drosophila miranda]
>gi|1036994223|ref|XP_017149094.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 11
>gi|1036994252|ref|XP_017149096.1| PREDICTED: protein aveugle [Drosophila miranda]
>gi|1036994209|ref|XP_017149093.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila n
>gi|1036994191|ref|XP_017149092.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila n
>gi|1037014834|ref|XP_017150343.1| PREDICTED: uncharacterized protein LOC108160699 [Drosophila
>gi|1037014816|ref|XP_017150342.1| PREDICTED: trafficking protein particle complex subunit 5 [I
>gi|1037016717|ref|XP_017150451.1| PREDICTED: kelch-like protein 40b [Drosophila miranda]
>gi|1037016735|ref|XP_017150452.1| PREDICTED: kelch-like protein 40b [Drosophila miranda]
>gi|1036978317|ref|XP_017148121.1| PREDICTED: LOW QUALITY PROTEIN: sprouty-related, EVH1 domair
>gi|1036978365|ref|XP_017148125.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036978335|ref|XP_017148123.1| PREDICTED: coiled-coil and C2 domain-containing protein 2A i
>gi|1036978351|ref|XP_017148124.1| PREDICTED: coiled-coil and C2 domain-containing protein 2A i
>gi|1036936761|ref|XP_017145615.1| PREDICTED: uncharacterized protein LOC108157981 [Drosophila
>gi|1036945317|ref|XP_017146139.1| PREDICTED: peptidyl-prolyl cis-trans isomerase E [Drosophila
>gi|1036945370|ref|XP_017146142.1| PREDICTED: 60S ribosomal protein L18a [Drosophila miranda]
>gi|1036945127|ref|XP_017146127.1| PREDICTED: uncharacterized protein LOC108158385 [Drosophila
>gi|1036945248|ref|XP_017146135.1| PREDICTED: kinesin-like protein KIF12 [Drosophila miranda]
>gi|1036945405|ref|XP_017146145.1| PREDICTED: guanine nucleotide-binding protein subunit gamma-
>gi|1036945335|ref|XP_017146140.1| PREDICTED: protein FAM192A [Drosophila miranda]
>gi|1036945266|ref|XP_017146136.1| PREDICTED: nucleolar GTP-binding protein 2 [Drosophila miran
>gi|1036945212|ref|XP_017146132.1| PREDICTED: E3 ubiquitin-protein ligase SH3RF1 isoform X1 [Dr
>gi|1036945230|ref|XP_017146134.1| PREDICTED: SH3 domain-containing RING finger protein 3 isofc
>gi|1036945423|ref|XP_017146146.1| PREDICTED: uncharacterized protein LOC108158399 [Drosophila
>gi|1036945387|ref|XP_017146143.1| PREDICTED: uncharacterized protein LOC108158397 [Drosophila
>gi|1036945353|ref|XP_017146141.1| PREDICTED: ras-related protein Rab-4B [Drosophila miranda]
>gi|1036945179|ref|XP_017146130.1| PREDICTED: endoribonuclease Dicer isoform X2 [Drosophila mir
>gi|1036945282|ref|XP_017146137.1| PREDICTED: fibrinogen-like protein 1 isoform X1 [Drosophila
>gi|1036945300|ref|XP_017146138.1| PREDICTED: angiopoietin-related protein 7 isoform X2 [Drosop
>gi|1036945145|ref|XP_017146128.1| PREDICTED: endoribonuclease Dicer isoform X1 [Drosophila mir
>gi|1036945163|ref|XP_017146129.1| PREDICTED: endoribonuclease Dicer isoform X1 [Drosophila mir
>gi|1036945196|ref|XP_017146131.1| PREDICTED: endoribonuclease Dicer isoform X1 [Drosophila mir
>gi|1037006830|ref|XP_017149862.1| PREDICTED: uncharacterized protein LOC108160388 isoform X3
>gi|1037006814|ref|XP_017149861.1| PREDICTED: uncharacterized protein LOC108160388 isoform X2
>gi|1037006796|ref|XP_017149860.1| PREDICTED: uncharacterized protein LOC108160388 isoform X1
>gi|1037013704|ref|XP_017150274.1| PREDICTED: 39S ribosomal protein L41, mitochondrial [Drosopl
>gi|1037013686|ref|XP_017150273.1| PREDICTED: probable splicing factor, arginine/serine-rich 7
>gi|1037020331|ref|XP_017150686.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1036987390|ref|XP_017148685.1| PREDICTED: uncharacterized protein LOC108159689 [Drosophila
>gi|1036987408|ref|XP_017148686.1| PREDICTED: uncharacterized protein LOC108159690 isoform X1
>gi|1036987428|ref|XP_017148687.1| PREDICTED: uncharacterized protein LOC108159690 isoform X2
>gi|1037010040|ref|XP_017150053.1| PREDICTED: uncharacterized protein LOC108160514 isoform X2
>gi|1037010020|ref|XP_017150052.1| PREDICTED: uncharacterized protein LOC108160514 isoform X1
>gi|1037010060|ref|XP_017150054.1| PREDICTED: nischarin [Drosophila miranda]
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>gi|1037013670|ref|XP_017150271.1| PREDICTED: Golgi pH regulator isoform X2 [Drosophila miranda
>gi|1037013654|ref|XP_017150270.1| PREDICTED: Golgi pH regulator isoform X1 [Drosophila miranda
>gi|1036936777|ref|XP_017145616.1| PREDICTED: uncharacterized protein LOC108157982 [Drosophila
>gi|1036941511|ref|XP_017145907.1| PREDICTED: uncharacterized protein LOC108158199 isoform X1
>gi|1036941529|ref|XP_017145908.1| PREDICTED: uncharacterized protein LOC108158199 isoform X1
>gi|1036941563|ref|XP_017145910.1| PREDICTED: uncharacterized protein LOC108158199 isoform X3
>gi|1036941545|ref|XP_017145909.1| PREDICTED: uncharacterized protein LOC108158199 isoform X2
>gi|1036941599|ref|XP_017145913.1| PREDICTED: uncharacterized protein LOC108158201 [Drosophila
>gi|1036941479|ref|XP_017145906.1| PREDICTED: methyltransferase-like protein 16 homolog [Drosop
>gi|1036941376|ref|XP_017145901.1| PREDICTED: SH3 domain-binding protein 5 homolog [Drosophila
>gi|1036941396|ref|XP_017145902.1| PREDICTED: SAM50-like protein CG7639 [Drosophila miranda]
>gi|1036941340|ref|XP_017145899.1| PREDICTED: protein RRNAD1 [Drosophila miranda]
>gi|1036941358|ref|XP_017145900.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1036941412|ref|XP_017145903.1| PREDICTED: septin-2 [Drosophila miranda]
>gi|1036941289|ref|XP_017145897.1| PREDICTED: putative RNA-binding protein 15 [Drosophila miran
>gi|1036941581|ref|XP_017145912.1| PREDICTED: tctex1 domain-containing protein 2 [Drosophila m
>gi|1036941322|ref|XP_017145898.1| PREDICTED: PX domain-containing protein kinase-like protein
>gi|1036941430|ref|XP_017145904.1| PREDICTED: COP9 signalosome complex subunit 4 [Drosophila m
>gi|1036941254|ref|XP_017145895.1| PREDICTED: acetyl-CoA carboxylase isoform X2 [Drosophila mi
>gi|1036941218|ref|XP_017145893.1| PREDICTED: acetyl-CoA carboxylase isoform X1 [Drosophila mi
>gi|1036941236|ref|XP_017145894.1| PREDICTED: acetyl-CoA carboxylase isoform X1 [Drosophila mi
>gi|1036941271|ref|XP_017145896.1| PREDICTED: acetyl-CoA carboxylase isoform X3 [Drosophila mi
>gi|1036941446|ref|XP_017145905.1| PREDICTED: nucleoporin seh1 [Drosophila miranda]
>gi|1036941613|ref|XP_017145914.1| PREDICTED: mitochondrial dicarboxylate carrier [Drosophila
>gi|1037021818|ref|XP_017150780.1| PREDICTED: hairy/enhancer-of-split related with YRPW motif p
>gi|1036942070|ref|XP_017145945.1| PREDICTED: uncharacterized protein LOC108158226 [Drosophila
>gi|1037004249|ref|XP_017149701.1| PREDICTED: ornithine decarboxylase 1 [Drosophila miranda]
>gi|1037004233|ref|XP_017149700.1| PREDICTED: slit homolog 2 protein isoform X2 [Drosophila mi
>gi|1037004152|ref|XP_017149698.1| PREDICTED: slit homolog 2 protein isoform X1 [Drosophila mi
>gi|1037004215|ref|XP_017149699.1| PREDICTED: slit homolog 2 protein isoform X1 [Drosophila mi
>gi|1037004267|ref|XP_017149702.1| PREDICTED: LOW QUALITY PROTEIN: biogenesis of lysosome-relat
>gi|1036936793|ref|XP_017145618.1| PREDICTED: transformer-2 sex-determining protein-like, parti
>gi|1036936809|ref|XP_017145619.1| PREDICTED: Golgi pH regulator-like [Drosophila miranda]
>gi|1037011750|ref|XP_017150152.1| PREDICTED: protein Optix [Drosophila miranda]
>gi|1037015824|ref|XP_017150399.1| PREDICTED: putative uncharacterized protein DDB_G0277255 iso
>gi|1037015740|ref|XP_017150396.1| PREDICTED: DNA-binding protein Ikaros isoform X1 [Drosophila
>gi|1037015792|ref|XP_017150397.1| PREDICTED: DNA-binding protein Ikaros isoform X2 [Drosophila
>gi|1037015806|ref|XP_017150398.1| PREDICTED: DNA-binding protein Ikaros isoform X2 [Drosophila
>gi|1036979871|ref|XP_017148219.1| PREDICTED: uncharacterized protein LOC108159442 [Drosophila
>gi|1036979713|ref|XP_017148208.1| PREDICTED: protein lingerer isoform X1 [Drosophila miranda]
>gi|1036979759|ref|XP_017148212.1| PREDICTED: protein lingerer isoform X4 [Drosophila miranda]
>gi|1036979727|ref|XP_017148210.1| PREDICTED: protein lingerer isoform X2 [Drosophila miranda]
>gi|1036979779|ref|XP_017148213.1| PREDICTED: protein lingerer isoform X5 [Drosophila miranda]
>gi|1036979796|ref|XP_017148214.1| PREDICTED: protein lingerer isoform X6 [Drosophila miranda]
>gi|1036979815|ref|XP_017148215.1| PREDICTED: protein lingerer isoform X7 [Drosophila miranda]
>gi|1036979849|ref|XP_017148217.1| PREDICTED: protein lingerer isoform X9 [Drosophila miranda]
>gi|1036979741|ref|XP_017148211.1| PREDICTED: protein lingerer isoform X3 [Drosophila miranda]
>gi|1036979833|ref|XP_017148216.1| PREDICTED: protein lingerer isoform X8 [Drosophila miranda]
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>gi|1036979863|ref|XP_017148218.1| PREDICTED: vacuolar protein sorting-associated protein 28 h
>gi|1037015988|ref|XP_017150407.1| PREDICTED: sugar transporter SWEET1 [Drosophila miranda]
>gi|1036942954|ref|XP_017146001.1| PREDICTED: uncharacterized protein LOC108158276 [Drosophila
>gi|1037014852|ref|XP_017150344.1| PREDICTED: multidrug resistance protein homolog 49 [Drosophi
>gi|1037015192|ref|XP_017150365.1| PREDICTED: uncharacterized protein LOC108160706 [Drosophila
>gi|1037015228|ref|XP_017150367.1| PREDICTED: uncharacterized protein LOC108160707 isoform X2
>gi|1037015210|ref|XP_017150366.1| PREDICTED: uncharacterized protein LOC108160707 isoform X1
>gi|1037015244|ref|XP_017150368.1| PREDICTED: uncharacterized protein LOC108160709 isoform X1
>gi|1037015262|ref|XP_017150369.1| PREDICTED: uncharacterized protein LOC108160709 isoform X2
>gi|1036999263|ref|XP_017149406.1| PREDICTED: polycomb group protein Psc [Drosophila miranda]
>gi|1037012155|ref|XP_017150178.1| PREDICTED: protein suppressor 2 of zeste [Drosophila mirand
>gi|1037020505|ref|XP_017150697.1| PREDICTED: probable salivary secreted peptide [Drosophila m
>gi|1037020520|ref|XP_017150698.1| PREDICTED: probable salivary secreted peptide [Drosophila m
>gi|1037017193|ref|XP_017150483.1| PREDICTED: tyrosine-protein kinase RYK [Drosophila miranda]
>gi|1037017205|ref|XP_017150484.1| PREDICTED: sideroflexin-1-like [Drosophila miranda]
>gi|1037019115|ref|XP_017150611.1| PREDICTED: abscisic-aldehyde oxidase-like [Drosophila miran
>gi|1036947068|ref|XP_017146248.1| PREDICTED: uncharacterized protein LOC108158468 [Drosophila
>gi|1036946994|ref|XP_017146244.1| PREDICTED: uncharacterized protein LOC108158464 [Drosophila
>gi|1036946958|ref|XP_017146242.1| PREDICTED: condensin complex subunit 3 isoform X1 [Drosophi
>gi|1036946976|ref|XP_017146243.1| PREDICTED: condensin complex subunit 3 isoform X2 [Drosophi
>gi|1036946904|ref|XP_017146238.1| PREDICTED: tyrosine-protein kinase PR2 isoform X2 [Drosophi
>gi|1036946886|ref|XP_017146237.1| PREDICTED: tyrosine-protein kinase PR2 isoform X1 [Drosophi
>gi|1036946940|ref|XP_017146240.1| PREDICTED: tripeptidyl-peptidase 2 isoform X2 [Drosophila m
>gi|1036946922|ref|XP_017146239.1| PREDICTED: tripeptidyl-peptidase 2 isoform X1 [Drosophila m
>gi|1036947029|ref|XP_017146246.1| PREDICTED: coiled-coil domain-containing protein 130 homolog
>gi|1036947050|ref|XP_017146247.1| PREDICTED: microtubule-associated protein RP/EB family membe
>gi|1036947011|ref|XP_017146245.1| PREDICTED: juvenile hormone epoxide hydrolase 1 [Drosophila
>gi|1037013383|ref|XP_017150254.1| PREDICTED: juvenile hormone epoxide hydrolase 1 [Drosophila
>gi|1037019310|ref|XP_017150622.1| PREDICTED: juvenile hormone epoxide hydrolase 1 [Drosophila
>gi|1036998921|ref|XP_017149389.1| PREDICTED: diptericin-like [Drosophila miranda]
>gi|1036998937|ref|XP_017149390.1| PREDICTED: diptericin-like [Drosophila miranda]
>gi|1036998904|ref|XP_017149388.1| PREDICTED: diptericin [Drosophila miranda]
>gi|1036998955|ref|XP_017149391.1| PREDICTED: uncharacterized protein LOC108160110 [Drosophila
>gi|1036998854|ref|XP_017149384.1| PREDICTED: V-type proton ATPase 116 kDa subunit a [Drosophil
>gi|1036998869|ref|XP_017149385.1| PREDICTED: uncharacterized protein LOC108160106 [Drosophila
>gi|1036998886|ref|XP_017149387.1| PREDICTED: uncharacterized protein LOC108160106 [Drosophila
>gi|1037007674|ref|XP_017149914.1| PREDICTED: uncharacterized protein LOC108160423 [Drosophila
>gi|1037007689|ref|XP_017149915.1| PREDICTED: uncharacterized protein LOC108160424 [Drosophila
>gi|1036999793|ref|XP_017149437.1| PREDICTED: LOW QUALITY PROTEIN: E3 ubiquitin-protein ligase
>gi|1037011930|ref|XP_017150163.1| PREDICTED: RING finger protein nhl-1 isoform X8 [Drosophila
>gi|1037011785|ref|XP_017150154.1| PREDICTED: RING finger protein nhl-1 isoform X1 [Drosophila
>gi|1037011824|ref|XP_017150155.1| PREDICTED: RING finger protein nhl-1 isoform X1 [Drosophila
>gi|1037011873|ref|XP_017150159.1| PREDICTED: RING finger protein nhl-1 isoform X4 [Drosophila
>gi|1037011841|ref|XP_017150156.1| PREDICTED: RING finger protein nhl-1 isoform X2 [Drosophila
>gi|1037011857|ref|XP_017150157.1| PREDICTED: RING finger protein nhl-1 isoform X3 [Drosophila
>gi|1037011888|ref|XP_017150160.1| PREDICTED: RING finger protein nhl-1 isoform X5 [Drosophila
>gi|1037011900|ref|XP_017150161.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1037011913|ref|XP_017150162.1| PREDICTED: cell wall protein IFF6 isoform X7 [Drosophila mi
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>gi|1037017489|ref|XP_017150503.1| PREDICTED: 5-hydroxytryptamine receptor 2B isoform X1 [Dros
>gi|1037017507|ref|XP_017150504.1| PREDICTED: 5-hydroxytryptamine receptor 2B isoform X2 [Dros
>gi|1037017524|ref|XP_017150505.1| PREDICTED: uncharacterized protein LOC108160798 [Drosophila
>gi|1037017542|ref|XP_017150506.1| PREDICTED: uncharacterized protein LOC108160798 [Drosophila
>gi|1036956788|ref|XP_017146840.1| PREDICTED: uncharacterized protein LOC108158742 [Drosophila
>gi|1036956770|ref|XP_017146838.1| PREDICTED: probable phospholipid hydroperoxide glutathione p
>gi|1036956738|ref|XP_017146836.1| PREDICTED: uncharacterized protein LOC108158739 [Drosophila
>gi|1036956573|ref|XP_017146825.1| PREDICTED: 5-hydroxytryptamine receptor 2A [Drosophila miran
>gi|1036956801|ref|XP_017146841.1| PREDICTED: RING-box protein 1A-like [Drosophila miranda]
>gi|1036956606|ref|XP_017146827.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036956660|ref|XP_017146831.1| PREDICTED: uncharacterized protein LOC108158737 [Drosophila
>gi|1036956672|ref|XP_017146832.1| PREDICTED: uncharacterized protein LOC108158737 [Drosophila
>gi|1036956690|ref|XP_017146833.1| PREDICTED: uncharacterized protein LOC108158737 [Drosophila
>gi|1036956753|ref|XP_017146837.1| PREDICTED: cyclin-dependent kinase 5 homolog [Drosophila mi
>gi|1036956624|ref|XP_017146828.1| PREDICTED: flotillin-1 isoform X1 [Drosophila miranda]
>gi|1036956642|ref|XP_017146829.1| PREDICTED: flotillin-1 isoform X2 [Drosophila miranda]
>gi|1036956588|ref|XP_017146826.1| PREDICTED: transitional endoplasmic reticulum ATPase TER94
>gi|1036956708|ref|XP_017146834.1| PREDICTED: cAMP-dependent protein kinase type II regulatory
>gi|1036956725|ref|XP_017146835.1| PREDICTED: cAMP-dependent protein kinase type II regulatory
>gi|1036961261|ref|XP_017147094.1| PREDICTED: palmitoyltransferase ZDHHC15 isoform X5 [Drosophi
>gi|1036961279|ref|XP_017147095.1| PREDICTED: palmitoyltransferase ZDHHC2 isoform X6 [Drosophil
>gi|1036961298|ref|XP_017147096.1| PREDICTED: palmitoyltransferase ZDHHC2 isoform X7 [Drosophil
>gi|1036961192|ref|XP_017147089.1| PREDICTED: palmitoyltransferase ZDHHC15 isoform X1 [Drosophi
>gi|1036961240|ref|XP_017147091.1| PREDICTED: palmitoyltransferase ZDHHC15 isoform X3 [Drosophi
>gi|1036961210|ref|XP_017147090.1| PREDICTED: palmitoyltransferase ZDHHC15 isoform X2 [Drosophi
>gi|1036961249|ref|XP_017147092.1| PREDICTED: palmitoyltransferase ZDHHC15 isoform X4 [Drosophi
>gi|1036961178|ref|XP_017147088.1| PREDICTED: putative methyltransferase C9orf114 [Drosophila r
>gi|1036961316|ref|XP_017147097.1| PREDICTED: activating signal cointegrator 1 complex subunit
>gi|1036961163|ref|XP_017147087.1| PREDICTED: lysophospholipid acyltransferase 1 [Drosophila m
>gi|1036961324|ref|XP_017147098.1| PREDICTED: mucin-2-like [Drosophila miranda]
>gi|1036961356|ref|XP_017147100.1| PREDICTED: cytochrome c oxidase subunit 7C, mitochondrial [
>gi|1036961374|ref|XP_017147101.1| PREDICTED: cytochrome c oxidase subunit 7C, mitochondrial [
>gi|1036961338|ref|XP_017147099.1| PREDICTED: protein canopy homolog 2 [Drosophila miranda]
>gi|1037020747|ref|XP_017150714.1| PREDICTED: defensin [Drosophila miranda]
>gi|1036946766|ref|XP_017146230.1| PREDICTED: SPARC-related modular calcium-binding protein 2
>gi|1036946782|ref|XP_017146231.1| PREDICTED: SPARC-related modular calcium-binding protein 2
>gi|1036946800|ref|XP_017146232.1| PREDICTED: SPARC-related modular calcium-binding protein 2
>gi|1036946816|ref|XP_017146233.1| PREDICTED: SPARC-related modular calcium-binding protein 2
>gi|1036946677|ref|XP_017146224.1| PREDICTED: GPI ethanolamine phosphate transferase 1 [Drosop
>gi|1036946712|ref|XP_017146226.1| PREDICTED: GPI ethanolamine phosphate transferase 1-like iso
>gi|1036946694|ref|XP_017146225.1| PREDICTED: GPI ethanolamine phosphate transferase 1-like iso
>gi|1036946870|ref|XP_017146236.1| PREDICTED: 39S ribosomal protein L42, mitochondrial [Drosop
>gi|1036946834|ref|XP_017146234.1| PREDICTED: cyclin-dependent kinase 10 [Drosophila miranda]
>gi|1036946852|ref|XP_017146235.1| PREDICTED: uncharacterized protein LOC108158457 [Drosophila
>gi|1036946593|ref|XP_017146219.1| PREDICTED: nodal modulator 2 [Drosophila miranda]
>gi|1036946629|ref|XP_017146221.1| PREDICTED: uncharacterized protein LOC108158449 isoform X1
>gi|1036946661|ref|XP_017146223.1| PREDICTED: uncharacterized protein KIAA0930 homolog isoform
>gi|1036946645|ref|XP_017146222.1| PREDICTED: uncharacterized protein LOC108158449 isoform X2
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>gi|1036946748|ref|XP_017146229.1| PREDICTED: spondin-1 [Drosophila miranda]
>gi|1036946611|ref|XP_017146220.1| PREDICTED: protein teflon [Drosophila miranda]
>gi|1036946730|ref|XP_017146227.1| PREDICTED: general transcription factor 3C polypeptide 3 [D
>gi|1036980764|ref|XP_017148276.1| PREDICTED: putative helicase mov-10-B.2 [Drosophila miranda]
>gi|1036980696|ref|XP_017148272.1| PREDICTED: uncharacterized protein LOC108159466 isoform X4
>gi|1036980712|ref|XP_017148273.1| PREDICTED: uncharacterized protein LOC108159466 isoform X4
>gi|1036980679|ref|XP_017148271.1| PREDICTED: uncharacterized protein LOC108159466 isoform X3
>gi|1036980662|ref|XP_017148270.1| PREDICTED: uncharacterized protein LOC108159466 isoform X2
>gi|1036980642|ref|XP_017148269.1| PREDICTED: uncharacterized protein LOC108159466 isoform X1
>gi|1036980748|ref|XP_017148275.1| PREDICTED: uncharacterized protein LOC108159466 isoform X6
>gi|1036980730|ref|XP_017148274.1| PREDICTED: uncharacterized protein LOC108159466 isoform X5
>gi|1037007414|ref|XP_017149899.1| PREDICTED: sphingosine-1-phosphate lyase [Drosophila miranda
>gi|1037007432|ref|XP_017149900.1| PREDICTED: sphingosine-1-phosphate lyase [Drosophila miranda
>gi|1037010724|ref|XP_017150092.1| PREDICTED: phenoloxidase 2 [Drosophila miranda]
>gi|1036936825|ref|XP_017145620.1| PREDICTED: uncharacterized protein LOC108157986 [Drosophila
>gi|1036936841|ref|XP_017145621.1| PREDICTED: uncharacterized protein LOC108157987 [Drosophila
>gi|1037010600|ref|XP_017150088.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1037010617|ref|XP_017150089.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1037010694|ref|XP_017150090.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1037010710|ref|XP_017150091.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1037017903|ref|XP_017150529.1| PREDICTED: E3 ubiquitin-protein ligase Topors-like [Drosoph
>gi|1037017943|ref|XP_017150530.1| PREDICTED: E3 ubiquitin-protein ligase Topors-like [Drosoph
>gi|1037017959|ref|XP_017150531.1| PREDICTED: E3 ubiquitin-protein ligase Topors-like [Drosoph
>gi|1037017850|ref|XP_017150526.1| PREDICTED: uncharacterized protein LOC108160814 [Drosophila
>gi|1037017868|ref|XP_017150527.1| PREDICTED: uncharacterized protein LOC108160814 [Drosophila
>gi|1037017885|ref|XP_017150528.1| PREDICTED: uncharacterized protein LOC108160814 [Drosophila
>gi|1037021838|ref|XP_017150781.1| PREDICTED: uncharacterized protein LOC108160988 [Drosophila
>gi|1037020007|ref|XP_017150667.1| PREDICTED: uncharacterized protein LOC108160905 [Drosophila
>gi|1036942710|ref|XP_017145986.1| PREDICTED: ras-interacting protein RIP3 [Drosophila miranda]
>gi|1037020715|ref|XP_017150712.1| PREDICTED: nucleoporin NSP1 [Drosophila miranda]
>gi|1036944156|ref|XP_017146074.1| PREDICTED: uncharacterized protein LOC108158346 [Drosophila
>gi|1036975880|ref|XP_017147968.1| PREDICTED: putative GPI-anchored protein PB15E9.01c [Drosoph
>gi|1036975948|ref|XP_017147973.1| PREDICTED: uncharacterized protein LOC108159310 [Drosophila
>gi|1036975932|ref|XP_017147972.1| PREDICTED: uncharacterized protein LOC108159309 [Drosophila
>gi|1036975846|ref|XP_017147966.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036975830|ref|XP_017147965.1| PREDICTED: putative uncharacterized protein YHR219W isoform
>gi|1036975814|ref|XP_017147964.1| PREDICTED: lisH domain-containing protein C1711.05 isoform
>gi|1036975862|ref|XP_017147967.1| PREDICTED: uncharacterized protein LOC108159306 [Drosophila
>gi|1036975798|ref|XP_017147963.1| PREDICTED: uncharacterized protein LOC108159302 [Drosophila
>gi|1036975898|ref|XP_017147970.1| PREDICTED: thymidylate kinase [Drosophila miranda]
>gi|1036975915|ref|XP_017147971.1| PREDICTED: thymidylate kinase [Drosophila miranda]
>gi|1036975745|ref|XP_017147960.1| PREDICTED: uncharacterized protein CG5098 isoform X1 [Drosop
>gi|1036975763|ref|XP_017147961.1| PREDICTED: uncharacterized protein CG5098 isoform X2 [Drosop
>gi|1036975780|ref|XP_017147962.1| PREDICTED: uncharacterized protein CG5098 isoform X3 [Drosop
>gi|1036936859|ref|XP_017145622.1| PREDICTED: uncharacterized protein YMR317W [Drosophila mira
>gi|1036946479|ref|XP_017146211.1| PREDICTED: uncharacterized protein LOC108158439 [Drosophila
>gi|1036991469|ref|XP_017148923.1| PREDICTED: maternal effect protein staufen [Drosophila mira
>gi|1036953390|ref|XP_017146640.1| PREDICTED: leukocyte elastase inhibitor [Drosophila miranda]
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>gi|1036953474|ref|XP_017146646.1| PREDICTED: retinol dehydrogenase 13 [Drosophila miranda]
>gi|1036953374|ref|XP_017146639.1| PREDICTED: dynactin subunit 4 [Drosophila miranda]
>gi|1036953492|ref|XP_017146647.1| PREDICTED: tubulin-folding cofactor B [Drosophila miranda]
>gi|1036953335|ref|XP_017146637.1| PREDICTED: MAP/microtubule affinity-regulating kinase 3 isof
>gi|1036953303|ref|XP_017146635.1| PREDICTED: MAP/microtubule affinity-regulating kinase 3 isof
>gi|1036953320|ref|XP_017146636.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X12
>gi|1036953280|ref|XP_017146634.1| PREDICTED: MAP/microtubule affinity-regulating kinase 3 isof
>gi|1036953252|ref|XP_017146632.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X8
>gi|1036953236|ref|XP_017146631.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X7
>gi|1036953204|ref|XP_017146629.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X5
>gi|1036953170|ref|XP_017146627.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X3
>gi|1036953222|ref|XP_017146630.1| PREDICTED: MAP/microtubule affinity-regulating kinase 3 isof
>gi|1036953186|ref|XP_017146628.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X4
>gi|1036953152|ref|XP_017146625.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X2
>gi|1036953116|ref|XP_017146623.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X1
>gi|1036953134|ref|XP_017146624.1| PREDICTED: serine/threonine-protein kinase MARK2 isoform X1
>gi|1036953264|ref|XP_017146633.1| PREDICTED: MAP/microtubule affinity-regulating kinase 3 isof
>gi|1036953353|ref|XP_017146638.1| PREDICTED: uncharacterized protein LOC108158650 isoform X14
>gi|1036953456|ref|XP_017146645.1| PREDICTED: meiotic recombination protein W68 [Drosophila mir
>gi|1036953402|ref|XP_017146642.1| PREDICTED: diacylglycerol O-acyltransferase 2-like [Drosophi
>gi|1036953422|ref|XP_017146643.1| PREDICTED: diacylglycerol O-acyltransferase 2-like [Drosophi
>gi|1036953440|ref|XP_017146644.1| PREDICTED: diacylglycerol O-acyltransferase 2-like [Drosophi
>gi|1036996148|ref|XP_017149214.1| PREDICTED: tyrosine-protein kinase receptor torso [Drosophil
>gi|1036936873|ref|XP_017145623.1| PREDICTED: uncharacterized protein LOC108157989 [Drosophila
>gi|1036996182|ref|XP_017149216.1| PREDICTED: ubiquinone biosynthesis protein COQ9, mitochondri
>gi|1036996202|ref|XP_017149218.1| PREDICTED: ubiquinone biosynthesis protein COQ9, mitochondri
>gi|1036996164|ref|XP_017149215.1| PREDICTED: glomulin [Drosophila miranda]
>gi|1036981733|ref|XP_017148338.1| PREDICTED: F-actin-uncapping protein LRRC16A isoform X3 [Dro
>gi|1036981751|ref|XP_017148339.1| PREDICTED: F-actin-uncapping protein LRRC16A isoform X4 [Dro
>gi|1036981798|ref|XP_017148340.1| PREDICTED: F-actin-uncapping protein LRRC16A isoform X5 [Dro
>gi|1036981695|ref|XP_017148335.1| PREDICTED: F-actin-uncapping protein LRRC16A isoform X1 [Dro
>gi|1036981715|ref|XP_017148336.1| PREDICTED: F-actin-uncapping protein LRRC16A isoform X2 [Dro
>gi|1036981811|ref|XP_017148341.1| PREDICTED: F-actin-uncapping protein LRRC16A isoform X6 [Dro
>gi|1036981837|ref|XP_017148342.1| PREDICTED: uncharacterized protein LOC108159501 isoform X7
>gi|1036981855|ref|XP_017148343.1| PREDICTED: mucin-21 isoform X8 [Drosophila miranda]
>gi|1037010925|ref|XP_017150102.1| PREDICTED: probable U2 small nuclear ribonucleoprotein A' [D
>gi|1037010944|ref|XP_017150103.1| PREDICTED: uncharacterized protein C20orf24 homolog [Drosoph
>gi|1037019047|ref|XP_017150606.1| PREDICTED: 39S ribosomal protein L52, mitochondrial [Drosoph
>gi|1036943423|ref|XP_017146031.1| PREDICTED: uncharacterized protein LOC108158301 [Drosophila
>gi|1036976760|ref|XP_017148022.1| PREDICTED: WD repeat-containing protein 19 isoform X2 [Droso
>gi|1036976746|ref|XP_017148021.1| PREDICTED: WD repeat-containing protein 19 isoform X1 [Droso
>gi|1036976789|ref|XP_017148025.1| PREDICTED: rab proteins geranylgeranyltransferase component
>gi|1036976775|ref|XP_017148023.1| PREDICTED: serine/threonine-protein kinase hippo [Drosophila
>gi|1036974601|ref|XP_017147889.1| PREDICTED: uncharacterized protein LOC108159260 isoform X1
>gi|1036974633|ref|XP_017147890.1| PREDICTED: uncharacterized protein LOC108159260 isoform X2
>gi|1036974652|ref|XP_017147891.1| PREDICTED: sister chromatid cohesion protein PDS5 homolog B
>gi|1036974670|ref|XP_017147893.1| PREDICTED: sister chromatid cohesion protein PDS5 homolog B
>gi|1036974739|ref|XP_017147897.1| PREDICTED: metallophosphoesterase 1 homolog isoform X1 [Dros
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>gi|1036974757|ref|XP_017147898.1| PREDICTED: metallophosphoesterase 1 isoform X2 [Drosophila r
>gi|1036974688|ref|XP_017147894.1| PREDICTED: tyrosine-protein kinase-like otk [Drosophila mira
>gi|1036974705|ref|XP_017147895.1| PREDICTED: tyrosine-protein kinase-like otk [Drosophila mira
>gi|1036974721|ref|XP_017147896.1| PREDICTED: tyrosine-protein kinase-like otk [Drosophila mira
>gi|1037005183|ref|XP_017149753.1| PREDICTED: tyrosine-protein kinase-like otk [Drosophila mira
>gi|1036950323|ref|XP_017146453.1| PREDICTED: glycerol kinase [Drosophila miranda]
>gi|1036950337|ref|XP_017146454.1| PREDICTED: uncharacterized protein LOC108158551 [Drosophila
>gi|1036950291|ref|XP_017146450.1| PREDICTED: heat shock factor protein isoform X4 [Drosophila
>gi|1036950273|ref|XP_017146449.1| PREDICTED: heat shock factor protein isoform X3 [Drosophila
>gi|1036950255|ref|XP_017146448.1| PREDICTED: heat shock factor protein isoform X2 [Drosophila
>gi|1036950237|ref|XP_017146447.1| PREDICTED: heat shock factor protein isoform X1 [Drosophila
>gi|1036950205|ref|XP_017146444.1| PREDICTED: polycomb protein Pcl [Drosophila miranda]
>gi|1036950221|ref|XP_017146445.1| PREDICTED: polycomb protein Pcl [Drosophila miranda]
>gi|1036950299|ref|XP_017146451.1| PREDICTED: polyadenylate-binding protein [Drosophila miranda
>gi|1036950311|ref|XP_017146452.1| PREDICTED: polyadenylate-binding protein [Drosophila miranda
>gi|1036950368|ref|XP_017146456.1| PREDICTED: protein EMRE homolog, mitochondrial [Drosophila r
>gi|1036950352|ref|XP_017146455.1| PREDICTED: ankyrin repeat domain-containing protein 13C [Dro
>gi|1036986619|ref|XP_017148637.1| PREDICTED: WD and tetratricopeptide repeats protein 1 [Droso
>gi|1036986654|ref|XP_017148639.1| PREDICTED: longitudinals lacking protein-like [Drosophila mi
>gi|1036986671|ref|XP_017148641.1| PREDICTED: longitudinals lacking protein-like [Drosophila mi
>gi|1036986636|ref|XP_017148638.1| PREDICTED: nitric oxide-associated protein 1 [Drosophila mir
>gi|1036988713|ref|XP_017148761.1| PREDICTED: uncharacterized protein CG10915 isoform X1 [Droso
>gi|1036988731|ref|XP_017148762.1| PREDICTED: uncharacterized protein CG10915 isoform X2 [Droso
>gi|1036988672|ref|XP_017148760.1| PREDICTED: nuclear pore complex protein Nup85 isoform X2 [Di
>gi|1036988653|ref|XP_017148759.1| PREDICTED: nuclear pore complex protein Nup85 isoform X1 [Di
>gi|1036988748|ref|XP_017148763.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037020973|ref|XP_017150729.1| PREDICTED: uncharacterized protein LOC108160949 [Drosophila
>gi|1037020955|ref|XP_017150728.1| PREDICTED: hepatic leukemia factor [Drosophila miranda]
>gi|1036943685|ref|XP_017146047.1| PREDICTED: uncharacterized protein LOC108158316 [Drosophila
>gi|1037021686|ref|XP_017150772.1| PREDICTED: aquaporin-12 isoform X2 [Drosophila miranda]
>gi|1037021672|ref|XP_017150771.1| PREDICTED: aquaporin-12 isoform X1 [Drosophila miranda]
>gi|1036967373|ref|XP_017147453.1| PREDICTED: transmembrane protein 186 [Drosophila miranda]
>gi|1036967312|ref|XP_017147449.1| PREDICTED: organic cation transporter protein isoform X2 [Di
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>gi|1036967435|ref|XP_017147455.1| PREDICTED: UPF0587 protein GA18326 [Drosophila miranda]
>gi|1036967455|ref|XP_017147456.1| PREDICTED: myeloma-overexpressed gene 2 protein homolog [Dro
>gi|1036967360|ref|XP_017147452.1| PREDICTED: F-box/SPRY domain-containing protein 1 [Drosophil
>gi|1036967327|ref|XP_017147450.1| PREDICTED: transcription factor Dp [Drosophila miranda]
>gi|1036967343|ref|XP_017147451.1| PREDICTED: probable peroxisomal membrane protein PEX13 [Dros
>gi|1036997074|ref|XP_017149272.1| PREDICTED: WD repeat-containing protein CG11141 [Drosophila
>gi|1036997089|ref|XP_017149273.1| PREDICTED: inositol-3-phosphate synthase [Drosophila miranda
>gi|1037011152|ref|XP_017150117.1| PREDICTED: dihydroceramide fatty acyl 2-hydroxylase FAH2 [Di
>gi|1037011166|ref|XP_017150118.1| PREDICTED: dihydroceramide fatty acyl 2-hydroxylase FAH2 [Di
>gi|1037011140|ref|XP_017150116.1| PREDICTED: uncharacterized protein LOC108160555 [Drosophila
>gi|1037009659|ref|XP_017150035.1| PREDICTED: probable elongation factor 1-beta [Drosophila mir
>gi|1037009677|ref|XP_017150036.1| PREDICTED: lysozyme [Drosophila miranda]
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>gi|1036942526|ref|XP_017145974.1| PREDICTED: lysozyme [Drosophila miranda]
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>gi|1037016299|ref|XP_017150427.1| PREDICTED: lysozyme 3 [Drosophila miranda]
>gi|1037016267|ref|XP_017150425.1| PREDICTED: lateral signaling target protein 2 homolog [Droso
>gi|1037016281|ref|XP_017150426.1| PREDICTED: pancreatic triacylglycerol lipase [Drosophila mir
>gi|1037019678|ref|XP_017150644.1| PREDICTED: protein kinase C, eye isozyme [Drosophila miranda
>gi|1037013109|ref|XP_017150238.1| PREDICTED: protein kinase C, brain isozyme isoform X1 [Droso
>gi|1037013124|ref|XP_017150239.1| PREDICTED: protein kinase C, brain isozyme isoform X1 [Droso
>gi|1037013138|ref|XP_017150240.1| PREDICTED: protein kinase C, brain isozyme isoform X1 [Droso
>gi|1037013152|ref|XP_017150241.1| PREDICTED: protein kinase C, brain isozyme isoform X1 [Droso
>gi|1037013168|ref|XP_017150242.1| PREDICTED: protein kinase C, brain isozyme isoform X2 [Droso
>gi|1037013186|ref|XP_017150244.1| PREDICTED: protein kinase C, brain isozyme isoform X3 [Droso
>gi|1037013202|ref|XP_017150245.1| PREDICTED: protein kinase C, brain isozyme isoform X4 [Droso
>gi|1036972074|ref|XP_017147731.1| PREDICTED: uncharacterized protein LOC108159188 [Drosophila
>gi|1036972202|ref|XP_017147737.1| PREDICTED: uncharacterized protein LOC108159193 [Drosophila
>gi|1036972168|ref|XP_017147736.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit
>gi|1036972058|ref|XP_017147729.1| PREDICTED: uncharacterized protein LOC108159187 [Drosophila
>gi|1036972220|ref|XP_017147739.1| PREDICTED: ubiquitin-fold modifier 1 [Drosophila miranda]
>gi|1036972092|ref|XP_017147732.1| PREDICTED: GPI mannosyltransferase 2 [Drosophila miranda]
>gi|1036972112|ref|XP_017147733.1| PREDICTED: GPI mannosyltransferase 2 [Drosophila miranda]
>gi|1036972144|ref|XP_017147735.1| PREDICTED: uncharacterized protein C1orf43 homolog [Drosophi
>gi|1036972128|ref|XP_017147734.1| PREDICTED: glyceraldehyde-3-phosphate dehydrogenase [Drosoph
>gi|1037018719|ref|XP_017150584.1| PREDICTED: calbindin-32 isoform X1 [Drosophila miranda]
>gi|1037018733|ref|XP_017150585.1| PREDICTED: calbindin-32 isoform X2 [Drosophila miranda]
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>gi|1037008015|ref|XP_017149935.1| PREDICTED: CAAX prenyl protease 1 homolog [Drosophila miran
>gi|1037008032|ref|XP_017149936.1| PREDICTED: CAAX prenyl protease 1 homolog isoform X1 [Drosop
>gi|1037008048|ref|XP_017149938.1| PREDICTED: CAAX prenyl protease 1 homolog isoform X2 [Drosop
>gi|1037008062|ref|XP_017149939.1| PREDICTED: CAAX prenyl protease 1 homolog [Drosophila miran
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>gi|1036965274|ref|XP_017147332.1| PREDICTED: uncharacterized protein LOC108158993 isoform X2
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>gi|1036965329|ref|XP_017147335.1| PREDICTED: probable serine/threonine-protein kinase cdc7 iso
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>gi|1036965575|ref|XP_017147350.1| PREDICTED: uncharacterized protein LOC108158998 isoform X2
>gi|1036936889|ref|XP_017145624.1| PREDICTED: uncharacterized protein LOC108157992 [Drosophila
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>gi|1036965505|ref|XP_017147346.1| PREDICTED: potassium voltage-gated channel subfamily KQT mem
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>gi|1036965397|ref|XP_017147340.1| PREDICTED: potassium voltage-gated channel subfamily KQT mem
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>gi|1036946413|ref|XP_017146207.1| PREDICTED: uncharacterized protein LOC108158435 [Drosophila
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>gi|1036993520|ref|XP_017149051.1| PREDICTED: probable nuclear hormone receptor HR3 isoform X4
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>gi|1036942543|ref|XP_017145975.1| PREDICTED: keratin, type I cytoskeletal 9 isoform X1 [Droso
>gi|1036942561|ref|XP_017145976.1| PREDICTED: keratin, type I cytoskeletal 9 isoform X2 [Droso
>gi|1036959923|ref|XP_017147015.1| PREDICTED: uncharacterized protein LOC108158827 isoform X3
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>gi|1036959941|ref|XP_017147016.1| PREDICTED: uncharacterized protein LOC108158827 isoform X4
>gi|1036959975|ref|XP_017147018.1| PREDICTED: protein enabled homolog isoform X6 [Drosophila m
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>gi|1036960011|ref|XP_017147021.1| PREDICTED: E3 ubiquitin-protein ligase MYLIP [Drosophila mi
>gi|1037019697|ref|XP_017150647.1| PREDICTED: uncharacterized protein LOC108160889 [Drosophila
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>gi|1037014577|ref|XP_017150328.1| PREDICTED: charged multivesicular body protein 6-A [Drosoph
>gi|1037014559|ref|XP_017150327.1| PREDICTED: cilia- and flagella-associated protein 57 [Droso
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>gi|1036979241|ref|XP_017148177.1| PREDICTED: alkylated DNA repair protein alkB homolog 8 [Dro
>gi|1036979259|ref|XP_017148178.1| PREDICTED: cyclin-dependent kinase 9 [Drosophila miranda]
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>gi|1036979276|ref|XP_017148179.1| PREDICTED: zinc finger Ran-binding domain-containing protei
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>gi|1037006288|ref|XP_017149828.1| PREDICTED: ribonuclease Z, mitochondrial [Drosophila miranda
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>gi|1036988505|ref|XP_017148747.1| PREDICTED: small subunit processome component 20 homolog [Dr
>gi|1036975555|ref|XP_017147949.1| PREDICTED: uncharacterized protein LOC108159292 isoform X4
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>gi|1036992398|ref|XP_017148980.1| PREDICTED: uncharacterized protein LOC108159879 isoform X1
>gi|1036992416|ref|XP_017148981.1| PREDICTED: uncharacterized protein LOC108159879 isoform X2
>gi|1036974477|ref|XP_017147882.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1036974531|ref|XP_017147885.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1036974547|ref|XP_017147886.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1036974495|ref|XP_017147883.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1036974513|ref|XP_017147884.1| PREDICTED: polyadenylate-binding protein-interacting protei
>gi|1036974429|ref|XP_017147879.1| PREDICTED: EH domain-binding protein 1 isoform X2 [Drosophil
>gi|1036974370|ref|XP_017147875.1| PREDICTED: EH domain-binding protein 1 isoform X1 [Drosophil
>gi|1036974388|ref|XP_017147876.1| PREDICTED: EH domain-binding protein 1 isoform X1 [Drosophil
>gi|1036974415|ref|XP_017147877.1| PREDICTED: EH domain-binding protein 1 isoform X1 [Drosophil
>gi|1036974441|ref|XP_017147880.1| PREDICTED: EH domain-binding protein 1 isoform X3 [Drosophil
>gi|1036974459|ref|XP_017147881.1| PREDICTED: EH domain-binding protein 1 isoform X4 [Drosophil
>gi|1036974583|ref|XP_017147888.1| PREDICTED: phosphatidylinositol 4,5-bisphosphate 5-phosphata
>gi|1036974565|ref|XP_017147887.1| PREDICTED: phosphatidylinositol 4,5-bisphosphate 5-phosphata
>gi|1036988599|ref|XP_017148755.1| PREDICTED: nuclear inhibitor of protein phosphatase 1 [Droso
>gi|1036988589|ref|XP_017148754.1| PREDICTED: probable asparagine--tRNA ligase, mitochondrial
>gi|1036988617|ref|XP_017148756.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036988636|ref|XP_017148758.1| PREDICTED: uncharacterized protein NCBP2-AS2 homolog [Droso
>gi|1036988556|ref|XP_017148751.1| PREDICTED: GTP-binding protein 1 [Drosophila miranda]
>gi|1036988573|ref|XP_017148752.1| PREDICTED: GTP-binding protein 1 [Drosophila miranda]
>gi|1036972240|ref|XP_017147740.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila m
>gi|1036972258|ref|XP_017147741.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila m
>gi|1036972275|ref|XP_017147742.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila m
>gi|1036972304|ref|XP_017147744.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X3 [Droso
>gi|1036972320|ref|XP_017147745.1| PREDICTED: uncharacterized protein LOC108159195 isoform X4
>gi|1036972289|ref|XP_017147743.1| PREDICTED: serine/arginine repetitive matrix protein 5 isofo
>gi|1036972533|ref|XP_017147760.1| PREDICTED: uncharacterized protein LOC108159201 isoform X1
>gi|1036972553|ref|XP_017147761.1| PREDICTED: uncharacterized protein LOC108159201 isoform X2
>gi|1036972505|ref|XP_017147758.1| PREDICTED: uncharacterized protein LOC108159200 isoform X2
>gi|1036972491|ref|XP_017147757.1| PREDICTED: uncharacterized protein LOC108159200 isoform X1
>gi|1036972518|ref|XP_017147759.1| PREDICTED: uncharacterized protein LOC108159200 isoform X3
>gi|1036972397|ref|XP_017147750.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972350|ref|XP_017147747.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972368|ref|XP_017147748.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972383|ref|XP_017147749.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972336|ref|XP_017147746.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972413|ref|XP_017147751.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972427|ref|XP_017147753.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972439|ref|XP_017147754.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036972457|ref|XP_017147755.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
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>gi|1036972475|ref|XP_017147756.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1037019063|ref|XP_017150607.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 [Drosopl
>gi|1037020126|ref|XP_017150675.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1037020144|ref|XP_017150676.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1037022218|ref|XP_017150804.1| PREDICTED: endoplasmic reticulum metallopeptidase 1-like [Dr
>gi|1036936907|ref|XP_017145625.1| PREDICTED: endoplasmic reticulum metallopeptidase 1-like [Dr
>gi|1036963230|ref|XP_017147216.1| PREDICTED: uncharacterized protein LOC108158924 isoform X1
>gi|1036963248|ref|XP_017147217.1| PREDICTED: uncharacterized protein LOC108158924 isoform X2
>gi|1036963264|ref|XP_017147218.1| PREDICTED: endoplasmic reticulum metallopeptidase 1-like [Dr
>gi|1036963350|ref|XP_017147224.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 [Drosopl
>gi|1036963314|ref|XP_017147221.1| PREDICTED: endoplasmic reticulum metallopeptidase 1-like iso
>gi|1036963332|ref|XP_017147222.1| PREDICTED: endoplasmic reticulum metallopeptidase 1-like iso
>gi|1036963366|ref|XP_017147225.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 [Drosopl
>gi|1036963280|ref|XP_017147219.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036963298|ref|XP_017147220.1| PREDICTED: endoplasmic reticulum metallopeptidase 1 isoform
>gi|1036963407|ref|XP_017147227.1| PREDICTED: integumentary mucin C.1 [Drosophila miranda]
>gi|1036963384|ref|XP_017147226.1| PREDICTED: endocuticle structural glycoprotein ABD-4 [Drosop
>gi|1037011569|ref|XP_017150142.1| PREDICTED: larval cuticle protein LCP-30 [Drosophila miranda
>gi|1037011585|ref|XP_017150143.1| PREDICTED: uncharacterized protein LOC108160570 [Drosophila
>gi|1037011601|ref|XP_017150145.1| PREDICTED: uncharacterized protein LOC108160571 [Drosophila
>gi|1037011525|ref|XP_017150139.1| PREDICTED: larval cuticle protein LCP-30 isoform X1 [Drosopl
>gi|1037011541|ref|XP_017150140.1| PREDICTED: larval cuticle protein LCP-30 isoform X2 [Drosopl
>gi|1037011557|ref|XP_017150141.1| PREDICTED: larval cuticle protein LCP-30 isoform X3 [Drosopl
>gi|1036945856|ref|XP_017146171.1| PREDICTED: endocuticle structural glycoprotein SgAbd-2 [Dros
>gi|1036936925|ref|XP_017145626.1| PREDICTED: odorant receptor 49a [Drosophila miranda]
>gi|1037019082|ref|XP_017150609.1| PREDICTED: endocuticle structural glycoprotein ABD-4 [Drosop
>gi|1037021509|ref|XP_017150758.1| PREDICTED: endocuticle structural glycoprotein SgAbd-3 [Dros
>gi|1037021491|ref|XP_017150757.1| PREDICTED: endocuticle structural glycoprotein SgAbd-3 [Dros
>gi|1036936943|ref|XP_017145627.1| PREDICTED: uncharacterized protein LOC108157995 [Drosophila
>gi|1036936961|ref|XP_017145628.1| PREDICTED: uncharacterized protein LOC108157996 [Drosophila
>gi|1036946496|ref|XP_017146212.1| PREDICTED: uncharacterized protein LOC108158440 [Drosophila
>gi|1036942724|ref|XP_017145987.1| PREDICTED: endocuticle structural glycoprotein SgAbd-2 [Dros
>gi|1037022429|ref|XP_017150817.1| PREDICTED: uncharacterized protein LOC108161016 [Drosophila
>gi|1036944007|ref|XP_017146065.1| PREDICTED: 4-coumarate--CoA ligase 1-like [Drosophila mirand
>gi|1037020842|ref|XP_017150721.1| PREDICTED: 4-coumarate--CoA ligase-like 7 [Drosophila mirand
>gi|1036969709|ref|XP_017147588.1| PREDICTED: 5-aminolevulinate synthase, erythroid-specific, r
>gi|1036969858|ref|XP_017147598.1| PREDICTED: nuclear receptor 2C2-associated protein isoform 
>gi|1036969838|ref|XP_017147597.1| PREDICTED: nuclear receptor 2C2-associated protein isoform 
>gi|1036969818|ref|XP_017147596.1| PREDICTED: AP-3 complex subunit sigma-2 [Drosophila miranda]
>gi|1036969800|ref|XP_017147594.1| PREDICTED: uncharacterized protein LOC108159122 [Drosophila
>gi|1036969764|ref|XP_017147592.1| PREDICTED: zinc finger protein 225 [Drosophila miranda]
>gi|1036969727|ref|XP_017147589.1| PREDICTED: anaphase-promoting complex subunit 2 [Drosophila
>gi|1036969746|ref|XP_017147591.1| PREDICTED: LOW QUALITY PROTEIN: anaphase-promoting complex s
>gi|1036969782|ref|XP_017147593.1| PREDICTED: uncharacterized protein LOC108159121 [Drosophila
>gi|1037003404|ref|XP_017149652.1| PREDICTED: putative transcription factor SOX-14 [Drosophila
>gi|1037014730|ref|XP_017150337.1| PREDICTED: peptidylglycine alpha-hydroxylating monooxygenase
>gi|1037014748|ref|XP_017150338.1| PREDICTED: peptidylglycine alpha-hydroxylating monooxygenase
>gi|1037014764|ref|XP_017150339.1| PREDICTED: paxillin homolog 1 [Drosophila miranda]
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>gi|1037014782|ref|XP_017150340.1| PREDICTED: paxillin homolog 1 [Drosophila miranda]
>gi|1037014800|ref|XP_017150341.1| PREDICTED: paxillin homolog 1 [Drosophila miranda]
>gi|1037015278|ref|XP_017150370.1| PREDICTED: PAS domain-containing serine/threonine-protein k
>gi|1037015302|ref|XP_017150371.1| PREDICTED: oxysterol-binding protein-related protein 2 [Dros
>gi|1037015316|ref|XP_017150373.1| PREDICTED: oxysterol-binding protein-related protein 2 [Dros
>gi|1037003143|ref|XP_017149636.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X2
>gi|1037003157|ref|XP_017149637.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X3
>gi|1037003125|ref|XP_017149635.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X1
>gi|1037003175|ref|XP_017149638.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X4
>gi|1037003211|ref|XP_017149640.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X6
>gi|1037003193|ref|XP_017149639.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X5
>gi|1037003227|ref|XP_017149641.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X7
>gi|1037007725|ref|XP_017149917.1| PREDICTED: synaptic vesicle 2-related protein isoform X1 [Dr
>gi|1037007741|ref|XP_017149918.1| PREDICTED: synaptic vesicle 2-related protein isoform X1 [Dr
>gi|1037007758|ref|XP_017149919.1| PREDICTED: synaptic vesicle 2-related protein isoform X2 [Dr
>gi|1037007707|ref|XP_017149916.1| PREDICTED: radial spoke head protein 4 homolog A [Drosophila
>gi|1037007780|ref|XP_017149920.1| PREDICTED: uncharacterized protein LOC108160428 [Drosophila
>gi|1037021125|ref|XP_017150739.1| PREDICTED: xyloside xylosyltransferase 1 [Drosophila miranda
>gi|1037014886|ref|XP_017150346.1| PREDICTED: adenylate kinase [Drosophila miranda]
>gi|1037008898|ref|XP_017149987.1| PREDICTED: ejaculatory bulb-specific protein 3 [Drosophila m
>gi|1037008916|ref|XP_017149988.1| PREDICTED: probable protein BRICK1-B [Drosophila miranda]
>gi|1036956912|ref|XP_017146847.1| PREDICTED: uncharacterized protein LOC108158748 [Drosophila
>gi|1036956962|ref|XP_017146850.1| PREDICTED: uncharacterized protein LOC108158751 [Drosophila
>gi|1036956894|ref|XP_017146846.1| PREDICTED: LIM domain-binding protein 2 [Drosophila miranda]
>gi|1036956865|ref|XP_017146843.1| PREDICTED: inner nuclear membrane protein Man1 [Drosophila m
>gi|1036956878|ref|XP_017146844.1| PREDICTED: inner nuclear membrane protein Man1 [Drosophila m
>gi|1036956930|ref|XP_017146848.1| PREDICTED: CCR4-NOT transcription complex subunit 11 [Drosop
>gi|1036956844|ref|XP_017146842.1| PREDICTED: integrator complex subunit 1 [Drosophila miranda]
>gi|1036956945|ref|XP_017146849.1| PREDICTED: cofilin/actin-depolymerizing factor homolog [Dros
>gi|1036981965|ref|XP_017148350.1| PREDICTED: gamma-soluble NSF attachment protein [Drosophila
>gi|1036981873|ref|XP_017148344.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1036981889|ref|XP_017148345.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1036981907|ref|XP_017148346.1| PREDICTED: potassium voltage-gated channel subfamily H membe
>gi|1036981929|ref|XP_017148348.1| PREDICTED: acid-sensing ion channel 5 isoform X1 [Drosophila
>gi|1036981947|ref|XP_017148349.1| PREDICTED: uncharacterized protein LOC108159503 isoform X2
>gi|1036982001|ref|XP_017148352.1| PREDICTED: eukaryotic translation initiation factor 5A [Dros
>gi|1036982019|ref|XP_017148353.1| PREDICTED: eukaryotic translation initiation factor 5A [Dros
>gi|1036981983|ref|XP_017148351.1| PREDICTED: 60S ribosomal protein L12 [Drosophila miranda]
>gi|1036982036|ref|XP_017148354.1| PREDICTED: 60S ribosomal protein L39 [Drosophila miranda]
>gi|1037006896|ref|XP_017149867.1| PREDICTED: ras-related protein Rap-2c [Drosophila miranda]
>gi|1037006926|ref|XP_017149869.1| PREDICTED: transcriptional coactivator yorkie isoform X2 [Dr
>gi|1037006910|ref|XP_017149868.1| PREDICTED: transcriptional coactivator yorkie isoform X1 [Dr
>gi|1036998087|ref|XP_017149336.1| PREDICTED: glycerol-3-phosphate acyltransferase 3 isoform X1
>gi|1036998100|ref|XP_017149337.1| PREDICTED: glycerol-3-phosphate acyltransferase 3 isoform X2
>gi|1036998117|ref|XP_017149339.1| PREDICTED: glycerol-3-phosphate acyltransferase 3-like isofo
>gi|1036998187|ref|XP_017149343.1| PREDICTED: muscle LIM protein Mlp84B isoform X2 [Drosophila
>gi|1036998153|ref|XP_017149341.1| PREDICTED: muscle LIM protein 1 isoform X1 [Drosophila miran
>gi|1036998171|ref|XP_017149342.1| PREDICTED: muscle LIM protein 1 isoform X1 [Drosophila miran
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>gi|1036998135|ref|XP_017149340.1| PREDICTED: carboxylesterase 5A [Drosophila miranda]
>gi|1037013005|ref|XP_017150232.1| PREDICTED: MICOS complex subunit Mic10 [Drosophila miranda]
>gi|1037012989|ref|XP_017150231.1| PREDICTED: uncharacterized protein LOC108160623 [Drosophila
>gi|1036951336|ref|XP_017146517.1| PREDICTED: E3 ubiquitin-protein ligase RBBP6 isoform X1 [Dro
>gi|1036951350|ref|XP_017146518.1| PREDICTED: E3 ubiquitin-protein ligase RBBP6 isoform X1 [Dro
>gi|1036951368|ref|XP_017146519.1| PREDICTED: E3 ubiquitin-protein ligase RBBP6 isoform X1 [Dro
>gi|1036951384|ref|XP_017146520.1| PREDICTED: E3 ubiquitin-protein ligase RBBP6 isoform X2 [Dro
>gi|1036951320|ref|XP_017146515.1| PREDICTED: serine/threonine-protein kinase Genghis Khan [Dro
>gi|1036951404|ref|XP_017146521.1| PREDICTED: histone acetyltransferase KAT6A isoform X1 [Droso
>gi|1036951450|ref|XP_017146524.1| PREDICTED: histone acetyltransferase KAT6A isoform X4 [Droso
>gi|1036951436|ref|XP_017146523.1| PREDICTED: histone acetyltransferase KAT6A isoform X3 [Droso
>gi|1036951422|ref|XP_017146522.1| PREDICTED: histone acetyltransferase KAT6A isoform X2 [Droso
>gi|1036962504|ref|XP_017147170.1| PREDICTED: helicase ARIP4 [Drosophila miranda]
>gi|1036936981|ref|XP_017145629.1| PREDICTED: beta-1,4-glucuronyltransferase 1 [Drosophila mira
>gi|1036936996|ref|XP_017145630.1| PREDICTED: uncharacterized protein LOC108157998 [Drosophila
>gi|1036962535|ref|XP_017147173.1| PREDICTED: myb-like protein P [Drosophila miranda]
>gi|1036962549|ref|XP_017147174.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036962524|ref|XP_017147171.1| PREDICTED: protein tamozhennic [Drosophila miranda]
>gi|1037000551|ref|XP_017149479.1| PREDICTED: LOW QUALITY PROTEIN: programmed cell death prote
>gi|1037000535|ref|XP_017149478.1| PREDICTED: N-alpha-acetyltransferase, 35 NatC auxiliary subu
>gi|1037016989|ref|XP_017150469.1| PREDICTED: dopamine N-acetyltransferase isoform X1 [Drosophi
>gi|1037017015|ref|XP_017150470.1| PREDICTED: dopamine N-acetyltransferase isoform X2 [Drosophi
>gi|1036937014|ref|XP_017145631.1| PREDICTED: uncharacterized protein LOC108157999 [Drosophila
>gi|1036992833|ref|XP_017149009.1| PREDICTED: RNA polymerase II-associated protein 3 [Drosophi
>gi|1036992851|ref|XP_017149010.1| PREDICTED: uncharacterized protein LOC108159898 [Drosophila
>gi|1036992897|ref|XP_017149012.1| PREDICTED: 28S ribosomal protein S17, mitochondrial [Drosop
>gi|1036992881|ref|XP_017149011.1| PREDICTED: ras-related GTP-binding protein C [Drosophila mi
>gi|1037017063|ref|XP_017150474.1| PREDICTED: lactoylglutathione lyase [Drosophila miranda]
>gi|1036980607|ref|XP_017148267.1| PREDICTED: PDZ domain-containing protein GIPC3 [Drosophila
>gi|1036980558|ref|XP_017148263.1| PREDICTED: phosphatidate phosphatase LPIN3 isoform X3 [Dros
>gi|1036980592|ref|XP_017148266.1| PREDICTED: phosphatidate phosphatase LPIN3 isoform X5 [Dros
>gi|1036980504|ref|XP_017148260.1| PREDICTED: phosphatidate phosphatase LPIN1 isoform X1 [Dros
>gi|1036980522|ref|XP_017148261.1| PREDICTED: phosphatidate phosphatase LPIN1 isoform X1 [Dros
>gi|1036980576|ref|XP_017148265.1| PREDICTED: phosphatidate phosphatase LPIN1 isoform X4 [Dros
>gi|1036980540|ref|XP_017148262.1| PREDICTED: phosphatidate phosphatase LPIN1 isoform X2 [Dros
>gi|1036980624|ref|XP_017148268.1| PREDICTED: general odorant-binding protein 99a [Drosophila
>gi|1036954610|ref|XP_017146718.1| PREDICTED: polyadenylate-binding protein 2 isoform X2 [Dros
>gi|1036954593|ref|XP_017146717.1| PREDICTED: polyadenylate-binding protein 2 isoform X1 [Dros
>gi|1036954640|ref|XP_017146719.1| PREDICTED: uncharacterized protein LOC108158695 [Drosophila
>gi|1036954522|ref|XP_017146713.1| PREDICTED: snRNA-activating protein complex subunit 3 [Dros
>gi|1036954540|ref|XP_017146714.1| PREDICTED: uncharacterized protein LOC108158692 [Drosophila
>gi|1036954575|ref|XP_017146715.1| PREDICTED: uncharacterized protein LOC108158693 [Drosophila
>gi|1036954506|ref|XP_017146712.1| PREDICTED: nuclear pore complex protein Nup50 [Drosophila m
>gi|1036954482|ref|XP_017146710.1| PREDICTED: arf-GAP with SH3 domain, ANK repeat and PH domai
>gi|1036954492|ref|XP_017146711.1| PREDICTED: arf-GAP with SH3 domain, ANK repeat and PH domai
>gi|1036960787|ref|XP_017147061.1| PREDICTED: cullin-4A [Drosophila miranda]
>gi|1036960838|ref|XP_017147062.1| PREDICTED: CSC1-like protein 2 [Drosophila miranda]
>gi|1036961006|ref|XP_017147075.1| PREDICTED: uncharacterized protein LOC108158855 [Drosophila
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>gi|1036960976|ref|XP_017147073.1| PREDICTED: TWiK family of potassium channels protein 7 isofc
>gi|1036960994|ref|XP_017147074.1| PREDICTED: TWiK family of potassium channels protein 7 isofc
>gi|1036960924|ref|XP_017147068.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036960956|ref|XP_017147070.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036960938|ref|XP_017147069.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036960854|ref|XP_017147063.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036960872|ref|XP_017147064.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036960908|ref|XP_017147067.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036960892|ref|XP_017147065.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1037016215|ref|XP_017150422.1| PREDICTED: uncharacterized protein LOC108160741 [Drosophila
>gi|1036942872|ref|XP_017145996.1| PREDICTED: selenoprotein BthD [Drosophila miranda]
>gi|1037017662|ref|XP_017150513.1| PREDICTED: D-beta-hydroxybutyrate dehydrogenase, mitochondri
>gi|1037022463|ref|XP_017150819.1| PREDICTED: uncharacterized protein LOC108161018 [Drosophila
>gi|1037010961|ref|XP_017150104.1| PREDICTED: uncharacterized protein LOC108160548 [Drosophila
>gi|1036952659|ref|XP_017146593.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC2 [Dro
>gi|1036952774|ref|XP_017146600.1| PREDICTED: GTP-binding protein 128up [Drosophila miranda]
>gi|1036952812|ref|XP_017146604.1| PREDICTED: uncharacterized protein LOC108158634 [Drosophila
>gi|1036952725|ref|XP_017146597.1| PREDICTED: protein kinase C-binding protein 1 isoform X2 [Di
>gi|1036952707|ref|XP_017146596.1| PREDICTED: protein kinase C-binding protein 1 isoform X1 [Di
>gi|1036952742|ref|XP_017146598.1| PREDICTED: protein kinase C-binding protein 1 isoform X3 [Di
>gi|1036952756|ref|XP_017146599.1| PREDICTED: zinc finger protein 551 [Drosophila miranda]
>gi|1036952632|ref|XP_017146591.1| PREDICTED: uncharacterized protein LOC108158626 isoform X1
>gi|1036952646|ref|XP_017146592.1| PREDICTED: uncharacterized protein LOC108158626 isoform X2
>gi|1036952790|ref|XP_017146601.1| PREDICTED: tigger transposable element-derived protein 6 [Di
>gi|1036952671|ref|XP_017146595.1| PREDICTED: zinc finger protein 471 [Drosophila miranda]
>gi|1036952800|ref|XP_017146602.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1037017049|ref|XP_017150473.1| PREDICTED: uncharacterized protein LOC108160780 [Drosophila
>gi|1037017031|ref|XP_017150471.1| PREDICTED: uncharacterized protein LOC108160779 [Drosophila
>gi|1037015524|ref|XP_017150381.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037015561|ref|XP_017150384.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037015543|ref|XP_017150383.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X2
>gi|1037018481|ref|XP_017150568.1| PREDICTED: centrosomal protein of 131 kDa isoform X3 [Drosop
>gi|1037018446|ref|XP_017150566.1| PREDICTED: centrosomal protein of 131 kDa isoform X1 [Drosop
>gi|1037018464|ref|XP_017150567.1| PREDICTED: centrosomal protein of 131 kDa isoform X2 [Drosop
>gi|1037018498|ref|XP_017150569.1| PREDICTED: C-type lectin domain family 2 member D3-like [Dro
>gi|1036937032|ref|XP_017145632.1| PREDICTED: uncharacterized protein LOC108158000 [Drosophila
>gi|1037013432|ref|XP_017150257.1| PREDICTED: uncharacterized protein LOC108160639 isoform X1
>gi|1037013450|ref|XP_017150258.1| PREDICTED: uncharacterized protein LOC108160639 isoform X2
>gi|1037013468|ref|XP_017150259.1| PREDICTED: uncharacterized protein LOC108160639 isoform X3
>gi|1037013496|ref|XP_017150261.1| PREDICTED: uncharacterized protein LOC108160640 [Drosophila
>gi|1037013317|ref|XP_017150250.1| PREDICTED: uncharacterized protein LOC108160633 isoform X2
>gi|1037013299|ref|XP_017150249.1| PREDICTED: uncharacterized protein LOC108160633 isoform X1
>gi|1037013335|ref|XP_017150251.1| PREDICTED: uncharacterized protein LOC108160633 isoform X3
>gi|1037008484|ref|XP_017149962.1| PREDICTED: phosphoenolpyruvate carboxykinase [GTP] [Drosophi
>gi|1036999022|ref|XP_017149394.1| PREDICTED: phosphoenolpyruvate carboxykinase [GTP] [Drosophi
>gi|1036998973|ref|XP_017149392.1| PREDICTED: sodium-dependent neutral amino acid transporter l
>gi|1036998991|ref|XP_017149393.1| PREDICTED: sodium-dependent neutral amino acid transporter l
>gi|1037010292|ref|XP_017150068.1| PREDICTED: E3 ubiquitin-protein ligase RNF25 [Drosophila mi
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>gi|1036948552|ref|XP_017146339.1| PREDICTED: dystonin isoform X34 [Drosophila miranda]
>gi|1036948570|ref|XP_017146340.1| PREDICTED: dystonin isoform X35 [Drosophila miranda]
>gi|1036948519|ref|XP_017146337.1| PREDICTED: dystonin isoform X32 [Drosophila miranda]
>gi|1036948534|ref|XP_017146338.1| PREDICTED: dystonin isoform X33 [Drosophila miranda]
>gi|1036948484|ref|XP_017146335.1| PREDICTED: dystonin isoform X30 [Drosophila miranda]
>gi|1036948466|ref|XP_017146334.1| PREDICTED: dystonin isoform X29 [Drosophila miranda]
>gi|1036948242|ref|XP_017146320.1| PREDICTED: dystonin isoform X16 [Drosophila miranda]
>gi|1036948501|ref|XP_017146336.1| PREDICTED: dystonin isoform X31 [Drosophila miranda]
>gi|1036948224|ref|XP_017146319.1| PREDICTED: uncharacterized protein LOC108158501 isoform X15
>gi|1036948209|ref|XP_017146318.1| PREDICTED: uncharacterized protein LOC108158501 isoform X14
>gi|1036948638|ref|XP_017146345.1| PREDICTED: dystonin isoform X39 [Drosophila miranda]
>gi|1036948378|ref|XP_017146329.1| PREDICTED: dystonin isoform X24 [Drosophila miranda]
>gi|1036948346|ref|XP_017146327.1| PREDICTED: dystonin isoform X22 [Drosophila miranda]
>gi|1036948364|ref|XP_017146328.1| PREDICTED: dystonin isoform X23 [Drosophila miranda]
>gi|1036948313|ref|XP_017146324.1| PREDICTED: dystonin isoform X20 [Drosophila miranda]
>gi|1036948295|ref|XP_017146323.1| PREDICTED: dystonin isoform X19 [Drosophila miranda]
>gi|1036948115|ref|XP_017146313.1| PREDICTED: uncharacterized protein LOC108158501 isoform X9
>gi|1036948031|ref|XP_017146307.1| PREDICTED: uncharacterized protein LOC108158501 isoform X4
>gi|1036948622|ref|XP_017146344.1| PREDICTED: dystonin isoform X38 [Drosophila miranda]
>gi|1036948604|ref|XP_017146343.1| PREDICTED: dystonin isoform X37 [Drosophila miranda]
>gi|1036948586|ref|XP_017146342.1| PREDICTED: dystonin isoform X36 [Drosophila miranda]
>gi|1036948331|ref|XP_017146326.1| PREDICTED: dystonin isoform X21 [Drosophila miranda]
>gi|1036948278|ref|XP_017146322.1| PREDICTED: dystonin isoform X18 [Drosophila miranda]
>gi|1036948142|ref|XP_017146315.1| PREDICTED: uncharacterized protein LOC108158501 isoform X11
>gi|1036947996|ref|XP_017146305.1| PREDICTED: uncharacterized protein LOC108158501 isoform X2
>gi|1036948159|ref|XP_017146316.1| PREDICTED: uncharacterized protein LOC108158501 isoform X12
>gi|1036948081|ref|XP_017146311.1| PREDICTED: uncharacterized protein LOC108158501 isoform X7
>gi|1036948098|ref|XP_017146312.1| PREDICTED: uncharacterized protein LOC108158501 isoform X8
>gi|1036948013|ref|XP_017146306.1| PREDICTED: uncharacterized protein LOC108158501 isoform X3
>gi|1036948066|ref|XP_017146310.1| PREDICTED: uncharacterized protein LOC108158501 isoform X6
>gi|1036948128|ref|XP_017146314.1| PREDICTED: uncharacterized protein LOC108158501 isoform X10
>gi|1036947978|ref|XP_017146304.1| PREDICTED: uncharacterized protein LOC108158501 isoform X1
>gi|1036948049|ref|XP_017146308.1| PREDICTED: uncharacterized protein LOC108158501 isoform X5
>gi|1036948676|ref|XP_017146347.1| PREDICTED: dystonin isoform X41 [Drosophila miranda]
>gi|1036948656|ref|XP_017146346.1| PREDICTED: dystonin isoform X40 [Drosophila miranda]
>gi|1036948449|ref|XP_017146333.1| PREDICTED: dystonin isoform X28 [Drosophila miranda]
>gi|1036948432|ref|XP_017146332.1| PREDICTED: dystonin isoform X27 [Drosophila miranda]
>gi|1036948396|ref|XP_017146330.1| PREDICTED: dystonin isoform X25 [Drosophila miranda]
>gi|1036948414|ref|XP_017146331.1| PREDICTED: dystonin isoform X26 [Drosophila miranda]
>gi|1036948712|ref|XP_017146349.1| PREDICTED: uncharacterized protein LOC108158501 isoform X43
>gi|1036948192|ref|XP_017146317.1| PREDICTED: uncharacterized protein LOC108158501 isoform X13
>gi|1036948694|ref|XP_017146348.1| PREDICTED: dystonin isoform X42 [Drosophila miranda]
>gi|1036948260|ref|XP_017146321.1| PREDICTED: uncharacterized protein LOC108158501 isoform X17
>gi|1036965819|ref|XP_017147364.1| PREDICTED: protein argonaute-2 isoform X2 [Drosophila mirand
>gi|1036965782|ref|XP_017147362.1| PREDICTED: protein argonaute-2 isoform X1 [Drosophila mirand
>gi|1036965800|ref|XP_017147363.1| PREDICTED: protein argonaute-2 isoform X1 [Drosophila mirand
>gi|1036965833|ref|XP_017147366.1| PREDICTED: protein argonaute-2 isoform X3 [Drosophila mirand
>gi|1036965901|ref|XP_017147370.1| PREDICTED: 39S ribosomal protein L53, mitochondrial [Drosoph
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>gi|1036965922|ref|XP_017147371.1| PREDICTED: uncharacterized protein LOC108159008 [Drosophila
>gi|1036965867|ref|XP_017147368.1| PREDICTED: proteasomal ubiquitin receptor ADRM1 homolog iso
>gi|1036965849|ref|XP_017147367.1| PREDICTED: proteasomal ubiquitin receptor ADRM1 homolog iso
>gi|1036965883|ref|XP_017147369.1| PREDICTED: putative nuclease HARBI1 [Drosophila miranda]
>gi|1036979409|ref|XP_017148189.1| PREDICTED: golgin subfamily A member 1 [Drosophila miranda]
>gi|1036979341|ref|XP_017148184.1| PREDICTED: centrosomin isoform X1 [Drosophila miranda]
>gi|1036979358|ref|XP_017148185.1| PREDICTED: centrosomin isoform X2 [Drosophila miranda]
>gi|1036979375|ref|XP_017148186.1| PREDICTED: centrosomin isoform X2 [Drosophila miranda]
>gi|1036979425|ref|XP_017148190.1| PREDICTED: lysozyme 1 [Drosophila miranda]
>gi|1036979392|ref|XP_017148187.1| PREDICTED: centrosomin isoform X3 [Drosophila miranda]
>gi|1036987117|ref|XP_017148668.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036987133|ref|XP_017148669.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036987151|ref|XP_017148670.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036987169|ref|XP_017148672.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036980875|ref|XP_017148284.1| PREDICTED: LOW QUALITY PROTEIN: protein CLP1 homolog [Drosop
>gi|1036980889|ref|XP_017148285.1| PREDICTED: histone H3-like centromeric protein cid [Drosoph
>gi|1036980977|ref|XP_017148290.1| PREDICTED: cholinephosphotransferase 1 isoform X4 [Drosophil
>gi|1036980915|ref|XP_017148287.1| PREDICTED: cholinephosphotransferase 1 isoform X2 [Drosophil
>gi|1036980943|ref|XP_017148288.1| PREDICTED: cholinephosphotransferase 1 isoform X3 [Drosophil
>gi|1036980903|ref|XP_017148286.1| PREDICTED: cholinephosphotransferase 1 isoform X1 [Drosophil
>gi|1036980961|ref|XP_017148289.1| PREDICTED: cholinephosphotransferase 1 isoform X1 [Drosophil
>gi|1037009641|ref|XP_017150034.1| PREDICTED: protein enhancer of sevenless 2B [Drosophila mir
>gi|1036937050|ref|XP_017145633.1| PREDICTED: uncharacterized protein LOC108158002 [Drosophila
>gi|1036963963|ref|XP_017147258.1| PREDICTED: guanylate kinase-associated protein mars isoform
>gi|1036963982|ref|XP_017147259.1| PREDICTED: guanylate kinase-associated protein mars isoform
>gi|1036963944|ref|XP_017147256.1| PREDICTED: protein lin-54 homolog isoform X4 [Drosophila mi
>gi|1036963910|ref|XP_017147254.1| PREDICTED: protein lin-54 homolog isoform X3 [Drosophila mi
>gi|1036963928|ref|XP_017147255.1| PREDICTED: protein lin-54 homolog isoform X3 [Drosophila mi
>gi|1036963889|ref|XP_017147253.1| PREDICTED: protein lin-54 homolog isoform X2 [Drosophila mi
>gi|1036964102|ref|XP_017147265.1| PREDICTED: elongation factor Tu [Drosophila miranda]
>gi|1036963871|ref|XP_017147252.1| PREDICTED: protein lin-54 homolog isoform X1 [Drosophila mi
>gi|1036964058|ref|XP_017147263.1| PREDICTED: uncharacterized protein LOC108158948 isoform X1
>gi|1036964084|ref|XP_017147264.1| PREDICTED: protein split ends isoform X2 [Drosophila miranda
>gi|1036964037|ref|XP_017147262.1| PREDICTED: synaptic vesicular amine transporter isoform X2
>gi|1036964000|ref|XP_017147260.1| PREDICTED: synaptic vesicular amine transporter isoform X1
>gi|1036964019|ref|XP_017147261.1| PREDICTED: synaptic vesicular amine transporter isoform X1
>gi|1036964119|ref|XP_017147267.1| PREDICTED: uncharacterized protein LOC108158950 [Drosophila
>gi|1036964133|ref|XP_017147268.1| PREDICTED: uncharacterized protein LOC108158951 [Drosophila
>gi|1036974213|ref|XP_017147865.1| PREDICTED: NAD kinase isoform X4 [Drosophila miranda]
>gi|1036974177|ref|XP_017147863.1| PREDICTED: NAD kinase isoform X2 [Drosophila miranda]
>gi|1036974159|ref|XP_017147862.1| PREDICTED: NAD kinase isoform X1 [Drosophila miranda]
>gi|1036974227|ref|XP_017147866.1| PREDICTED: NAD kinase isoform X5 [Drosophila miranda]
>gi|1036974195|ref|XP_017147864.1| PREDICTED: NAD kinase isoform X3 [Drosophila miranda]
>gi|1036974245|ref|XP_017147867.1| PREDICTED: NAD kinase isoform X6 [Drosophila miranda]
>gi|1036974317|ref|XP_017147872.1| PREDICTED: NAD kinase-like isoform X4 [Drosophila miranda]
>gi|1036974281|ref|XP_017147869.1| PREDICTED: NAD kinase-like isoform X2 [Drosophila miranda]
>gi|1036974263|ref|XP_017147868.1| PREDICTED: NAD kinase-like isoform X1 [Drosophila miranda]
>gi|1036974299|ref|XP_017147871.1| PREDICTED: NAD kinase-like isoform X3 [Drosophila miranda]
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>gi|1036974352|ref|XP_017147874.1| PREDICTED: grpE protein homolog, mitochondrial [Drosophila r
>gi|1036974334|ref|XP_017147873.1| PREDICTED: uncharacterized protein LOC108159254 [Drosophila
>gi|1036944974|ref|XP_017146118.1| PREDICTED: uncharacterized protein LOC108158375 [Drosophila
>gi|1036945009|ref|XP_017146120.1| PREDICTED: uncharacterized protein LOC108158379 [Drosophila
>gi|1037015330|ref|XP_017150374.1| PREDICTED: retinal homeobox protein Rx-B [Drosophila miranda
>gi|1037004295|ref|XP_017149705.1| PREDICTED: uncharacterized protein LOC108160303 [Drosophila
>gi|1037004281|ref|XP_017149703.1| PREDICTED: actin-57B [Drosophila miranda]
>gi|1037019712|ref|XP_017150648.1| PREDICTED: retinal homeobox protein Rx [Drosophila miranda]
>gi|1036943266|ref|XP_017146021.1| PREDICTED: LOW QUALITY PROTEIN: carbonic anhydrase-related p
>gi|1036994804|ref|XP_017149132.1| PREDICTED: homeobox protein orthopedia [Drosophila miranda]
>gi|1036994772|ref|XP_017149130.1| PREDICTED: calcium and integrin-binding family member 2 iso
>gi|1036994786|ref|XP_017149131.1| PREDICTED: calcium and integrin-binding family member 2 iso
>gi|1036994750|ref|XP_017149129.1| PREDICTED: calcium and integrin-binding family member 2 iso
>gi|1036994634|ref|XP_017149122.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036994653|ref|XP_017149123.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036994668|ref|XP_017149124.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036994680|ref|XP_017149125.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036994700|ref|XP_017149126.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036994718|ref|XP_017149127.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036994734|ref|XP_017149128.1| PREDICTED: inositol-pentakisphosphate 2-kinase [Drosophila r
>gi|1036942854|ref|XP_017145995.1| PREDICTED: immune-induced peptide 14 [Drosophila miranda]
>gi|1037020657|ref|XP_017150708.1| PREDICTED: immune-induced peptide 4 [Drosophila miranda]
>gi|1037020645|ref|XP_017150707.1| PREDICTED: carbonic anhydrase 1 [Drosophila miranda]
>gi|1037022150|ref|XP_017150799.1| PREDICTED: luciferin 4-monooxygenase [Drosophila miranda]
>gi|1037021583|ref|XP_017150764.1| PREDICTED: luciferin 4-monooxygenase [Drosophila miranda]
>gi|1036988537|ref|XP_017148750.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036988520|ref|XP_017148749.1| PREDICTED: uncharacterized protein LOC108159728 [Drosophila
>gi|1036946428|ref|XP_017146208.1| PREDICTED: uncharacterized protein LOC108158436 [Drosophila
>gi|1036937068|ref|XP_017145634.1| PREDICTED: extensin-like [Drosophila miranda]
>gi|1036937085|ref|XP_017145635.1| PREDICTED: uncharacterized protein LOC108158004 [Drosophila
>gi|1036937100|ref|XP_017145637.1| PREDICTED: uncharacterized protein PB18E9.04c-like [Drosophi
>gi|1036937115|ref|XP_017145638.1| PREDICTED: uncharacterized protein LOC108158006 [Drosophila
>gi|1036937133|ref|XP_017145639.1| PREDICTED: ras-associated and pleckstrin homology domains-co
>gi|1036937147|ref|XP_017145640.1| PREDICTED: histone-lysine N-methyltransferase SETD1A [Drosop
>gi|1037013055|ref|XP_017150235.1| PREDICTED: uncharacterized protein LOC108160626 isoform X2
>gi|1037013039|ref|XP_017150234.1| PREDICTED: Down syndrome cell adhesion molecule homolog iso
>gi|1037013091|ref|XP_017150237.1| PREDICTED: tyrosine-protein kinase receptor TYRO3 isoform X4
>gi|1037013073|ref|XP_017150236.1| PREDICTED: tyrosine-protein kinase receptor TYRO3 isoform X3
>gi|1037013023|ref|XP_017150233.1| PREDICTED: E3 ubiquitin-protein ligase MARCH4 [Drosophila mi
>gi|1036943791|ref|XP_017146052.1| PREDICTED: uncharacterized protein LOC108158323 [Drosophila
>gi|1036937167|ref|XP_017145641.1| PREDICTED: uncharacterized protein LOC108158009 [Drosophila
>gi|1036937189|ref|XP_017145642.1| PREDICTED: putative gustatory receptor 57a [Drosophila mirar
>gi|1036960713|ref|XP_017147056.1| PREDICTED: MIP18 family protein CG30152 [Drosophila miranda]
>gi|1036960646|ref|XP_017147052.1| PREDICTED: maternal protein exuperantia-1 [Drosophila mirand
>gi|1036960660|ref|XP_017147053.1| PREDICTED: maternal protein exuperantia-1 [Drosophila mirand
>gi|1036960612|ref|XP_017147050.1| PREDICTED: protein fem-1 homolog B isoform X1 [Drosophila mi
>gi|1036960630|ref|XP_017147051.1| PREDICTED: protein fem-1 homolog B isoform X2 [Drosophila mi
>gi|1036960769|ref|XP_017147060.1| PREDICTED: uncharacterized protein LOC108158848 [Drosophila
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>gi|1036960595|ref|XP_017147049.1| PREDICTED: slit homolog 3 protein [Drosophila miranda]
>gi|1036960689|ref|XP_017147055.1| PREDICTED: lysozyme [Drosophila miranda]
>gi|1036960735|ref|XP_017147057.1| PREDICTED: lysozyme isoform X1 [Drosophila miranda]
>gi|1036960751|ref|XP_017147059.1| PREDICTED: lysozyme isoform X2 [Drosophila miranda]
>gi|1036960675|ref|XP_017147054.1| PREDICTED: DNA methyltransferase 1-associated protein 1 [Dro
>gi|1036952586|ref|XP_017146589.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC10 [Di
>gi|1036952536|ref|XP_017146585.1| PREDICTED: yrdC domain-containing protein, mitochondrial [Di
>gi|1036952553|ref|XP_017146587.1| PREDICTED: mitochondrial inner membrane protease subunit 2
>gi|1036952570|ref|XP_017146588.1| PREDICTED: transcription factor MafG [Drosophila miranda]
>gi|1036952604|ref|XP_017146590.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036952437|ref|XP_017146579.1| PREDICTED: protein rigor mortis [Drosophila miranda]
>gi|1036952484|ref|XP_017146582.1| PREDICTED: UPF0704 protein C6orf165 homolog [Drosophila mira
>gi|1036952500|ref|XP_017146583.1| PREDICTED: uncharacterized protein LOC108158619 [Drosophila
>gi|1036952420|ref|XP_017146578.1| PREDICTED: WASH complex subunit FAM21 homolog [Drosophila mi
>gi|1036952454|ref|XP_017146580.1| PREDICTED: PIN2/TERF1-interacting telomerase inhibitor 1 iso
>gi|1036952469|ref|XP_017146581.1| PREDICTED: PIN2/TERF1-interacting telomerase inhibitor 1 iso
>gi|1036952518|ref|XP_017146584.1| PREDICTED: DDB1- and CUL4-associated factor 11 [Drosophila n
>gi|1036980230|ref|XP_017148244.1| PREDICTED: uncharacterized protein LOC108159457 isoform X1
>gi|1036980250|ref|XP_017148245.1| PREDICTED: uncharacterized protein LOC108159457 isoform X1
>gi|1036980272|ref|XP_017148246.1| PREDICTED: uncharacterized protein LOC108159457 isoform X1
>gi|1036980290|ref|XP_017148247.1| PREDICTED: kyphoscoliosis peptidase isoform X2 [Drosophila n
>gi|1036980308|ref|XP_017148248.1| PREDICTED: kyphoscoliosis peptidase isoform X3 [Drosophila n
>gi|1036980328|ref|XP_017148249.1| PREDICTED: heterogeneous nuclear ribonucleoprotein K isoform
>gi|1036980362|ref|XP_017148252.1| PREDICTED: glycine-rich cell wall structural protein 1 isofo
>gi|1036980380|ref|XP_017148253.1| PREDICTED: glycine-rich cell wall structural protein 1 isofo
>gi|1036980345|ref|XP_017148251.1| PREDICTED: heterogeneous nuclear ribonucleoprotein K isoform
>gi|1037016765|ref|XP_017150455.1| PREDICTED: uncharacterized protein LOC108160765 [Drosophila
>gi|1037009946|ref|XP_017150047.1| PREDICTED: uncharacterized protein LOC108160513 isoform X1
>gi|1037009964|ref|XP_017150048.1| PREDICTED: uncharacterized protein LOC108160513 isoform X2
>gi|1037009982|ref|XP_017150049.1| PREDICTED: uncharacterized protein LOC108160513 isoform X2
>gi|1037010002|ref|XP_017150050.1| PREDICTED: uncharacterized protein LOC108160513 isoform X3
>gi|1036964980|ref|XP_017147314.1| PREDICTED: proteasome subunit alpha type-6 [Drosophila miran
>gi|1036964994|ref|XP_017147315.1| PREDICTED: proteasome subunit alpha type-6 [Drosophila miran
>gi|1036964936|ref|XP_017147310.1| PREDICTED: deoxyribodipyrimidine photo-lyase [Drosophila mir
>gi|1036964918|ref|XP_017147309.1| PREDICTED: exocyst complex component 3 [Drosophila miranda]
>gi|1036964849|ref|XP_017147305.1| PREDICTED: ubiquitin-like modifier-activating enzyme ATG7 is
>gi|1036964887|ref|XP_017147307.1| PREDICTED: ubiquitin-like modifier-activating enzyme ATG7 is
>gi|1036964865|ref|XP_017147306.1| PREDICTED: ubiquitin-like modifier-activating enzyme ATG7 is
>gi|1036964967|ref|XP_017147313.1| PREDICTED: LOW QUALITY PROTEIN: 32 kDa beta-galactoside-bind
>gi|1036964903|ref|XP_017147308.1| PREDICTED: cysteine dioxygenase type 1-like isoform X4 [Dros
>gi|1036965059|ref|XP_017147318.1| PREDICTED: protein FAM136A [Drosophila miranda]
>gi|1036965011|ref|XP_017147316.1| PREDICTED: protein FAM136A [Drosophila miranda]
>gi|1036965041|ref|XP_017147317.1| PREDICTED: 28S ribosomal protein S28, mitochondrial [Drosoph
>gi|1036964951|ref|XP_017147312.1| PREDICTED: probable inactive tRNA-specific adenosine deamina
>gi|1036944139|ref|XP_017146073.1| PREDICTED: uncharacterized protein LOC108158345 [Drosophila
>gi|1036990509|ref|XP_017148874.1| PREDICTED: uncharacterized protein LOC108159815 isoform X1
>gi|1036990535|ref|XP_017148875.1| PREDICTED: uncharacterized protein LOC108159815 isoform X2
>gi|1037001679|ref|XP_017149550.1| PREDICTED: uncharacterized protein LOC108160213 isoform X2
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>gi|1037001612|ref|XP_017149545.1| PREDICTED: skin secretory protein xP2 isoform X4 [Drosophila
>gi|1037001597|ref|XP_017149544.1| PREDICTED: skin secretory protein xP2 isoform X3 [Drosophila
>gi|1037001579|ref|XP_017149543.1| PREDICTED: skin secretory protein xP2 isoform X2 [Drosophila
>gi|1037001563|ref|XP_017149542.1| PREDICTED: skin secretory protein xP2 isoform X1 [Drosophila
>gi|1037001628|ref|XP_017149546.1| PREDICTED: retinitis pigmentosa 1-like 1 protein [Drosophila
>gi|1037001662|ref|XP_017149548.1| PREDICTED: uncharacterized protein LOC108160213 isoform X1
>gi|1037001644|ref|XP_017149547.1| PREDICTED: uncharacterized protein LOC108160212 [Drosophila
>gi|1036998610|ref|XP_017149369.1| PREDICTED: nephrin isoform X1 [Drosophila miranda]
>gi|1036998628|ref|XP_017149370.1| PREDICTED: nephrin isoform X2 [Drosophila miranda]
>gi|1036986340|ref|XP_017148621.1| PREDICTED: protein charlatan isoform X1 [Drosophila miranda]
>gi|1036986374|ref|XP_017148622.1| PREDICTED: protein charlatan isoform X1 [Drosophila miranda]
>gi|1036986400|ref|XP_017148624.1| PREDICTED: protein charlatan isoform X2 [Drosophila miranda]
>gi|1036986416|ref|XP_017148625.1| PREDICTED: protein charlatan isoform X3 [Drosophila miranda]
>gi|1036940909|ref|XP_017145879.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu
>gi|1036978730|ref|XP_017148145.1| PREDICTED: protein vestigial-like [Drosophila miranda]
>gi|1036978694|ref|XP_017148143.1| PREDICTED: aspartate--tRNA ligase, cytoplasmic-like isoform
>gi|1036978712|ref|XP_017148144.1| PREDICTED: aspartate--tRNA ligase, cytoplasmic-like isoform
>gi|1036978765|ref|XP_017148147.1| PREDICTED: protein lifeguard 1-like isoform X2 [Drosophila r
>gi|1036978748|ref|XP_017148146.1| PREDICTED: protein lifeguard 1-like isoform X1 [Drosophila r
>gi|1036978800|ref|XP_017148149.1| PREDICTED: protein lifeguard 1-like isoform X2 [Drosophila r
>gi|1036978782|ref|XP_017148148.1| PREDICTED: protein lifeguard 1-like isoform X1 [Drosophila r
>gi|1036978818|ref|XP_017148150.1| PREDICTED: protein lifeguard 1-like isoform X3 [Drosophila r
>gi|1036978660|ref|XP_017148140.1| PREDICTED: carcinine transporter-like isoform X1 [Drosophila
>gi|1036978676|ref|XP_017148141.1| PREDICTED: carcinine transporter-like isoform X2 [Drosophila
>gi|1036978836|ref|XP_017148153.1| PREDICTED: transcription factor BTF3 homolog 4-like [Drosoph
>gi|1036937205|ref|XP_017145643.1| PREDICTED: 50S ribosomal protein L3, partial [Drosophila mir
>gi|1036944958|ref|XP_017146117.1| PREDICTED: carbonic anhydrase 4-like, partial [Drosophila mi
>gi|1036937221|ref|XP_017145644.1| PREDICTED: focadhesin-like [Drosophila miranda]
>gi|1036986438|ref|XP_017148626.1| PREDICTED: uncharacterized protein LOC108159660 [Drosophila
>gi|1036986459|ref|XP_017148627.1| PREDICTED: focadhesin-like [Drosophila miranda]
>gi|1037014266|ref|XP_017150308.1| PREDICTED: uncharacterized protein LOC108160673 isoform X1
>gi|1037014284|ref|XP_017150309.1| PREDICTED: uncharacterized protein LOC108160673 isoform X2
>gi|1037014300|ref|XP_017150310.1| PREDICTED: uncharacterized protein LOC108160675 [Drosophila
>gi|1037017353|ref|XP_017150494.1| PREDICTED: uncharacterized protein LOC108160794 isoform X2
>gi|1037017370|ref|XP_017150495.1| PREDICTED: uncharacterized protein LOC108160794 isoform X3
>gi|1037017336|ref|XP_017150493.1| PREDICTED: uncharacterized protein LOC108160794 isoform X1
>gi|1037017384|ref|XP_017150496.1| PREDICTED: uncharacterized protein LOC108160794 isoform X4
>gi|1036937254|ref|XP_017145645.1| PREDICTED: uncharacterized protein LOC108158014 [Drosophila
>gi|1037017319|ref|XP_017150492.1| PREDICTED: odorant receptor 59a [Drosophila miranda]
>gi|1037016628|ref|XP_017150447.1| PREDICTED: uncharacterized protein LOC108160758 [Drosophila
>gi|1037016664|ref|XP_017150449.1| PREDICTED: S-phase kinase-associated protein 1 [Drosophila r
>gi|1037016648|ref|XP_017150448.1| PREDICTED: odorant receptor 59b [Drosophila miranda]
>gi|1036936019|ref|XP_017145566.1| PREDICTED: putative odorant receptor 59c [Drosophila miranda
>gi|1037016682|ref|XP_017150450.1| PREDICTED: uncharacterized protein LOC108160761 [Drosophila
>gi|1036943862|ref|XP_017146057.1| PREDICTED: uncharacterized protein LOC108158327 [Drosophila
>gi|1037021021|ref|XP_017150732.1| PREDICTED: uncharacterized protein LOC108160952 [Drosophila
>gi|1037011453|ref|XP_017150135.1| PREDICTED: vacuolar protein sorting-associated protein 21 is
>gi|1037011433|ref|XP_017150134.1| PREDICTED: ras-related protein RHN1 isoform X1 [Drosophila r
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>gi|1037011471|ref|XP_017150136.1| PREDICTED: ras-related protein Rab-21 isoform X3 [Drosophila
>gi|1037011489|ref|XP_017150137.1| PREDICTED: ras-related protein Rab-21 isoform X3 [Drosophila
>gi|1036973216|ref|XP_017147805.1| PREDICTED: uncharacterized protein LOC108159219 isoform X1
>gi|1036973239|ref|XP_017147806.1| PREDICTED: uncharacterized protein LOC108159219 isoform X1
>gi|1036973255|ref|XP_017147808.1| PREDICTED: uncharacterized protein LOC108159219 isoform X2
>gi|1036973293|ref|XP_017147810.1| PREDICTED: protein brown isoform X1 [Drosophila miranda]
>gi|1036973326|ref|XP_017147811.1| PREDICTED: protein brown isoform X2 [Drosophila miranda]
>gi|1036973393|ref|XP_017147816.1| PREDICTED: pickpocket protein 19 [Drosophila miranda]
>gi|1036973343|ref|XP_017147812.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036973358|ref|XP_017147813.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036973200|ref|XP_017147804.1| PREDICTED: 5-oxoprolinase [Drosophila miranda]
>gi|1036973375|ref|XP_017147814.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
>gi|1036973275|ref|XP_017147809.1| PREDICTED: sodium/potassium-transporting ATPase subunit alph
>gi|1037021794|ref|XP_017150779.1| PREDICTED: uncharacterized protein LOC108160986 [Drosophila
>gi|1036991842|ref|XP_017148945.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1036991860|ref|XP_017148946.1| PREDICTED: probable palmitoyltransferase ZDHHC24 [Drosophila
>gi|1036991878|ref|XP_017148947.1| PREDICTED: uncharacterized protein LOC108159859 [Drosophila
>gi|1036991828|ref|XP_017148944.1| PREDICTED: DNA repair protein RAD50 [Drosophila miranda]
>gi|1037002216|ref|XP_017149585.1| PREDICTED: 28S ribosomal protein S29, mitochondrial [Drosoph
>gi|1037002234|ref|XP_017149587.1| PREDICTED: KH domain-containing, RNA-binding, signal transdu
>gi|1036987619|ref|XP_017148698.1| PREDICTED: KH domain-containing, RNA-binding, signal transdu
>gi|1036987707|ref|XP_017148703.1| PREDICTED: DNA polymerase epsilon subunit 4 [Drosophila mira
>gi|1036987637|ref|XP_017148699.1| PREDICTED: KH domain-containing, RNA-binding, signal transdu
>gi|1036987654|ref|XP_017148700.1| PREDICTED: KH domain-containing, RNA-binding, signal transdu
>gi|1036987689|ref|XP_017148702.1| PREDICTED: ventrally expressed gene D protein [Drosophila mi
>gi|1036987671|ref|XP_017148701.1| PREDICTED: uncharacterized protein LOC108159699 [Drosophila
>gi|1036999369|ref|XP_017149413.1| PREDICTED: uncharacterized protein LOC108160129 [Drosophila
>gi|1036999317|ref|XP_017149410.1| PREDICTED: protein amalgam [Drosophila miranda]
>gi|1036999352|ref|XP_017149412.1| PREDICTED: immunoglobulin superfamily member 10 [Drosophila
>gi|1037002463|ref|XP_017149600.1| PREDICTED: RNA polymerase-associated protein Rtf1 [Drosophil
>gi|1036967941|ref|XP_017147486.1| PREDICTED: uncharacterized protein LOC108159058 isoform X1
>gi|1036967954|ref|XP_017147487.1| PREDICTED: uncharacterized protein LOC108159058 isoform X2
>gi|1036967755|ref|XP_017147475.1| PREDICTED: kazrin isoform X1 [Drosophila miranda]
>gi|1036967770|ref|XP_017147477.1| PREDICTED: kazrin isoform X1 [Drosophila miranda]
>gi|1036967785|ref|XP_017147478.1| PREDICTED: kazrin isoform X1 [Drosophila miranda]
>gi|1036967801|ref|XP_017147479.1| PREDICTED: kazrin isoform X1 [Drosophila miranda]
>gi|1036967817|ref|XP_017147480.1| PREDICTED: kazrin isoform X1 [Drosophila miranda]
>gi|1036967833|ref|XP_017147481.1| PREDICTED: kazrin isoform X1 [Drosophila miranda]
>gi|1036967849|ref|XP_017147482.1| PREDICTED: kazrin-A isoform X2 [Drosophila miranda]
>gi|1036967867|ref|XP_017147483.1| PREDICTED: kazrin-A isoform X3 [Drosophila miranda]
>gi|1036967984|ref|XP_017147490.1| PREDICTED: uncharacterized protein LOC108159060 [Drosophila
>gi|1036967968|ref|XP_017147488.1| PREDICTED: uncharacterized protein LOC108159059 [Drosophila
>gi|1036967927|ref|XP_017147485.1| PREDICTED: probable hydroxyacid-oxoacid transhydrogenase, m
>gi|1036967907|ref|XP_017147484.1| PREDICTED: uncharacterized protein LOC108159055 [Drosophila
>gi|1037010078|ref|XP_017150055.1| PREDICTED: protein windpipe [Drosophila miranda]
>gi|1037010094|ref|XP_017150056.1| PREDICTED: protein windpipe [Drosophila miranda]
>gi|1037010112|ref|XP_017150057.1| PREDICTED: protein windpipe [Drosophila miranda]
>gi|1037010136|ref|XP_017150058.1| PREDICTED: protein windpipe [Drosophila miranda]
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>gi|1037018044|ref|XP_017150537.1| PREDICTED: uncharacterized protein LOC108160820 [Drosophila
>gi|1037018062|ref|XP_017150538.1| PREDICTED: uncharacterized protein LOC108160820 [Drosophila
>gi|1037006040|ref|XP_017149811.1| PREDICTED: uncharacterized protein LOC108160355 [Drosophila
>gi|1037011735|ref|XP_017150151.1| PREDICTED: solute carrier organic anion transporter family r
>gi|1037020683|ref|XP_017150709.1| PREDICTED: solute carrier organic anion transporter family r
>gi|1036996743|ref|XP_017149252.1| PREDICTED: solute carrier organic anion transporter family r
>gi|1036996829|ref|XP_017149257.1| PREDICTED: uncharacterized protein LOC108160029 [Drosophila
>gi|1036996760|ref|XP_017149253.1| PREDICTED: protein ariadne-2 isoform X1 [Drosophila miranda]
>gi|1036996776|ref|XP_017149254.1| PREDICTED: protein ariadne-2 isoform X2 [Drosophila miranda]
>gi|1036996812|ref|XP_017149256.1| PREDICTED: pyridoxal phosphate phosphatase [Drosophila miranda
>gi|1036996794|ref|XP_017149255.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila r
>gi|1037021601|ref|XP_017150765.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila r
>gi|1036994123|ref|XP_017149088.1| PREDICTED: tubulointerstitial nephritis antigen-like [Drosop
>gi|1036994109|ref|XP_017149087.1| PREDICTED: vacuolar protein sorting-associated protein 35 is
>gi|1036994095|ref|XP_017149085.1| PREDICTED: vacuolar protein sorting-associated protein 35 is
>gi|1036997830|ref|XP_017149319.1| PREDICTED: netrin receptor unc-5 isoform X1 [Drosophila mira
>gi|1036997848|ref|XP_017149320.1| PREDICTED: netrin receptor unc-5 isoform X2 [Drosophila mira
>gi|1036997866|ref|XP_017149321.1| PREDICTED: netrin receptor unc-5 isoform X3 [Drosophila mira
>gi|1036989703|ref|XP_017148823.1| PREDICTED: protein apterous [Drosophila miranda]
>gi|1036989687|ref|XP_017148822.1| PREDICTED: LOW QUALITY PROTEIN: disks large-associated prote
>gi|1037010829|ref|XP_017150096.1| PREDICTED: U3 small nucleolar RNA-interacting protein 2 [Dro
>gi|1036976982|ref|XP_017148038.1| PREDICTED: uncharacterized protein LOC108159343 isoform X7
>gi|1036976964|ref|XP_017148036.1| PREDICTED: uncharacterized protein LOC108159343 isoform X6
>gi|1036976946|ref|XP_017148035.1| PREDICTED: uncharacterized protein LOC108159343 isoform X5
>gi|1036976930|ref|XP_017148034.1| PREDICTED: uncharacterized protein LOC108159343 isoform X4
>gi|1036976902|ref|XP_017148032.1| PREDICTED: uncharacterized protein LOC108159343 isoform X2
>gi|1036976916|ref|XP_017148033.1| PREDICTED: uncharacterized protein LOC108159343 isoform X3
>gi|1036976892|ref|XP_017148031.1| PREDICTED: uncharacterized protein LOC108159343 isoform X1
>gi|1036996549|ref|XP_017149240.1| PREDICTED: cytochrome c oxidase assembly protein COX19 [Dros
>gi|1036996501|ref|XP_017149236.1| PREDICTED: phosphopantothenoylcysteine decarboxylase [Drosop
>gi|1036996469|ref|XP_017149234.1| PREDICTED: cyclic nucleotide-gated cation channel beta-1 [D
>gi|1036996485|ref|XP_017149235.1| PREDICTED: cyclic nucleotide-gated cation channel beta-1 [D
>gi|1036996515|ref|XP_017149237.1| PREDICTED: high mobility group protein Z [Drosophila miranda
>gi|1036996533|ref|XP_017149238.1| PREDICTED: high mobility group protein Z [Drosophila miranda
>gi|1037020021|ref|XP_017150668.1| PREDICTED: uncharacterized protein LOC108160906 [Drosophila
>gi|1036993190|ref|XP_017149030.1| PREDICTED: high mobility group protein D [Drosophila miranda
>gi|1036993134|ref|XP_017149027.1| PREDICTED: transport and Golgi organization protein 11 [Dros
>gi|1036993152|ref|XP_017149028.1| PREDICTED: transport and Golgi organization protein 11 [Dros
>gi|1036993168|ref|XP_017149029.1| PREDICTED: MIT domain-containing protein 1 [Drosophila mirar
>gi|1036989502|ref|XP_017148811.1| PREDICTED: lamin-B receptor [Drosophila miranda]
>gi|1036989522|ref|XP_017148812.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1036989661|ref|XP_017148820.1| PREDICTED: glutaredoxin-C4 [Drosophila miranda]
>gi|1036937271|ref|XP_017145647.1| PREDICTED: uncharacterized protein LOC108158015 [Drosophila
>gi|1036989574|ref|XP_017148815.1| PREDICTED: antichymotrypsin-2-like [Drosophila miranda]
>gi|1036989540|ref|XP_017148813.1| PREDICTED: lysyl oxidase homolog 2B [Drosophila miranda]
>gi|1036989671|ref|XP_017148821.1| PREDICTED: uncharacterized protein LOC108159773 [Drosophila
>gi|1036989592|ref|XP_017148816.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036989556|ref|XP_017148814.1| PREDICTED: trypsin-1 [Drosophila miranda]
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>gi|1036989610|ref|XP_017148817.1| PREDICTED: serine/threonine-protein phosphatase PP1-like [D:
>gi|1036989649|ref|XP_017148818.1| PREDICTED: serine/threonine-protein phosphatase PP1 [Drosop]
>gi|1037019893|ref|XP_017150659.1| PREDICTED: serine/threonine-protein phosphatase alpha-3 iso:
>gi|1037019913|ref|XP_017150660.1| PREDICTED: serine/threonine-protein phosphatase alpha-2 iso:
>gi|1037010258|ref|XP_017150066.1| PREDICTED: uncharacterized protein LOC108160521 [Drosophila
>gi|1036942119|ref|XP_017145949.1| PREDICTED: mucin-2 [Drosophila miranda]
>gi|1036942756|ref|XP_017145990.1| PREDICTED: uncharacterized protein LOC108158264 [Drosophila
>gi|1036942327|ref|XP_017145962.1| PREDICTED: uncharacterized protein LOC108158240 [Drosophila
>gi|1036942052|ref|XP_017145944.1| PREDICTED: L-asparaginase-like protein GA18140 [Drosophila n
>gi|1036937288|ref|XP_017145648.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1037018293|ref|XP_017150554.1| PREDICTED: probable serine/threonine-protein kinase kinX [D:
>gi|1036984716|ref|XP_017148519.1| PREDICTED: serine/threonine-protein phosphatase alpha-3 iso:
>gi|1036984653|ref|XP_017148515.1| PREDICTED: nyctalopin [Drosophila miranda]
>gi|1036984671|ref|XP_017148516.1| PREDICTED: nyctalopin [Drosophila miranda]
>gi|1036984732|ref|XP_017148520.1| PREDICTED: uncharacterized protein LOC108159582 [Drosophila
>gi|1036984698|ref|XP_017148518.1| PREDICTED: leukocyte elastase inhibitor C-like [Drosophila n
>gi|1036984749|ref|XP_017148521.1| PREDICTED: uncharacterized protein LOC108159584 [Drosophila
>gi|1036937305|ref|XP_017145649.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036984687|ref|XP_017148517.1| PREDICTED: uncharacterized protein LOC108159579 [Drosophila
>gi|1037020875|ref|XP_017150722.1| PREDICTED: putative gustatory receptor 58c [Drosophila mirar
>gi|1037020891|ref|XP_017150724.1| PREDICTED: putative gustatory receptor 58b [Drosophila mirar
>gi|1037020923|ref|XP_017150725.1| PREDICTED: putative gustatory receptor 58b [Drosophila mirar
>gi|1036937321|ref|XP_017145650.1| PREDICTED: putative gustatory receptor 58a [Drosophila mirar
>gi|1037014230|ref|XP_017150306.1| PREDICTED: uncharacterized protein LOC108160671 [Drosophila
>gi|1037014144|ref|XP_017150301.1| PREDICTED: uncharacterized protein LOC108160669 [Drosophila
>gi|1037014162|ref|XP_017150302.1| PREDICTED: uncharacterized protein LOC108160669 [Drosophila
>gi|1037014180|ref|XP_017150303.1| PREDICTED: uncharacterized protein LOC108160669 [Drosophila
>gi|1037014198|ref|XP_017150304.1| PREDICTED: uncharacterized protein LOC108160669 [Drosophila
>gi|1037014248|ref|XP_017150307.1| PREDICTED: transmembrane emp24 domain-containing protein 2
>gi|1037014214|ref|XP_017150305.1| PREDICTED: serine/threonine-protein phosphatase alpha-2 iso:
>gi|1036937339|ref|XP_017145651.1| PREDICTED: uncharacterized protein LOC108158019 [Drosophila
>gi|1036985658|ref|XP_017148579.1| PREDICTED: uncharacterized protein LOC108159632 [Drosophila
>gi|1036985712|ref|XP_017148582.1| PREDICTED: ERI1 exoribonuclease 2 isoform X2 [Drosophila mi]
>gi|1036985676|ref|XP_017148580.1| PREDICTED: ERI1 exoribonuclease 2 isoform X1 [Drosophila mi]
>gi|1036985694|ref|XP_017148581.1| PREDICTED: ERI1 exoribonuclease 2 isoform X1 [Drosophila mi]
>gi|1036985622|ref|XP_017148576.1| PREDICTED: protein-glutamate O-methyltransferase [Drosophila
>gi|1036985640|ref|XP_017148577.1| PREDICTED: protein-glutamate O-methyltransferase [Drosophila
>gi|1036985604|ref|XP_017148575.1| PREDICTED: protein-glutamate O-methyltransferase [Drosophila
>gi|1037009000|ref|XP_017149994.1| PREDICTED: ceramide glucosyltransferase [Drosophila miranda]
>gi|1036957636|ref|XP_017146887.1| PREDICTED: synaptojanin-1 [Drosophila miranda]
>gi|1036957825|ref|XP_017146897.1| PREDICTED: LOW QUALITY PROTEIN: tetratricopeptide repeat pro
>gi|1036957807|ref|XP_017146895.1| PREDICTED: syndecan isoform X6 [Drosophila miranda]
>gi|1036957771|ref|XP_017146893.1| PREDICTED: syndecan isoform X4 [Drosophila miranda]
>gi|1036957755|ref|XP_017146892.1| PREDICTED: syndecan isoform X3 [Drosophila miranda]
>gi|1036957737|ref|XP_017146891.1| PREDICTED: syndecan isoform X2 [Drosophila miranda]
>gi|1036957789|ref|XP_017146894.1| PREDICTED: syndecan isoform X5 [Drosophila miranda]
>gi|1036957654|ref|XP_017146888.1| PREDICTED: syndecan isoform X1 [Drosophila miranda]
>gi|1036957672|ref|XP_017146889.1| PREDICTED: syndecan isoform X1 [Drosophila miranda]
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>gi|1036957690|ref|XP_017146890.1| PREDICTED: syndecan isoform X1 [Drosophila miranda]
>gi|1037003624|ref|XP_017149668.1| PREDICTED: zinc finger FYVE domain-containing protein 9 [Dro
>gi|1036988949|ref|XP_017148776.1| PREDICTED: FK506-binding protein 2 isoform X2 [Drosophila mi
>gi|1036988930|ref|XP_017148775.1| PREDICTED: FK506-binding protein 2 isoform X1 [Drosophila mi
>gi|1036937355|ref|XP_017145652.1| PREDICTED: uncharacterized protein LOC108158020 [Drosophila
>gi|1036989183|ref|XP_017148790.1| PREDICTED: protein NDRG3 isoform X1 [Drosophila miranda]
>gi|1036989203|ref|XP_017148791.1| PREDICTED: protein NDRG3 isoform X2 [Drosophila miranda]
>gi|1036989219|ref|XP_017148792.1| PREDICTED: protein NDRG3 isoform X3 [Drosophila miranda]
>gi|1036989265|ref|XP_017148796.1| PREDICTED: protein NDRG3 isoform X6 [Drosophila miranda]
>gi|1036989301|ref|XP_017148798.1| PREDICTED: protein NDRG3 isoform X8 [Drosophila miranda]
>gi|1036989327|ref|XP_017148799.1| PREDICTED: protein NDRG3 isoform X9 [Drosophila miranda]
>gi|1036989339|ref|XP_017148800.1| PREDICTED: protein NDRG3 isoform X10 [Drosophila miranda]
>gi|1036989283|ref|XP_017148797.1| PREDICTED: protein NDRG3 isoform X7 [Drosophila miranda]
>gi|1036989237|ref|XP_017148793.1| PREDICTED: protein NDRG3 isoform X4 [Drosophila miranda]
>gi|1036989249|ref|XP_017148794.1| PREDICTED: protein NDRG3 isoform X5 [Drosophila miranda]
>gi|1036989353|ref|XP_017148801.1| PREDICTED: uncharacterized protein LOC108159757 [Drosophila
>gi|1036979903|ref|XP_017148221.1| PREDICTED: epidermal growth factor receptor isoform X2 [Dros
>gi|1036936037|ref|XP_017145568.1| PREDICTED: transmembrane protease serine 9-like [Drosophila
>gi|1036979889|ref|XP_017148220.1| PREDICTED: epidermal growth factor receptor isoform X1 [Dros
>gi|1037017400|ref|XP_017150497.1| PREDICTED: BTB/POZ domain-containing protein kctd15-like iso
>gi|1037017418|ref|XP_017150499.1| PREDICTED: BTB/POZ domain-containing protein kctd15-like iso
>gi|1037017111|ref|XP_017150477.1| PREDICTED: uncharacterized protein LOC108160784 isoform X1
>gi|1037017129|ref|XP_017150478.1| PREDICTED: uncharacterized protein LOC108160784 isoform X2
>gi|1037017097|ref|XP_017150476.1| PREDICTED: uncharacterized protein LOC108160783 [Drosophila
>gi|1037011216|ref|XP_017150122.1| PREDICTED: uncharacterized protein LOC108160559 [Drosophila
>gi|1037015416|ref|XP_017150379.1| PREDICTED: solute carrier family 25 member 35-like [Drosophi
>gi|1037015400|ref|XP_017150378.1| PREDICTED: solute carrier family 25 member 35-like [Drosophi
>gi|1037012558|ref|XP_017150203.1| PREDICTED: uncharacterized protein LOC108160606 isoform X1
>gi|1037012576|ref|XP_017150204.1| PREDICTED: uncharacterized protein LOC108160606 isoform X2
>gi|1037012540|ref|XP_017150202.1| PREDICTED: peptidyl-prolyl cis-trans isomerase 6 [Drosophila
>gi|1037010398|ref|XP_017150074.1| PREDICTED: solute carrier family 35 member F5 [Drosophila mi
>gi|1037017436|ref|XP_017150500.1| PREDICTED: uncharacterized protein LOC108160796 [Drosophila
>gi|1037017454|ref|XP_017150501.1| PREDICTED: uncharacterized protein LOC108160796 [Drosophila
>gi|1037017471|ref|XP_017150502.1| PREDICTED: uncharacterized protein LOC108160796 [Drosophila
>gi|1036941819|ref|XP_017145927.1| PREDICTED: beta-galactoside alpha-2,6-sialyltransferase 2 [
>gi|1036999547|ref|XP_017149424.1| PREDICTED: stromelysin-3 isoform X1 [Drosophila miranda]
>gi|1036999565|ref|XP_017149425.1| PREDICTED: stromelysin-3 isoform X1 [Drosophila miranda]
>gi|1036999583|ref|XP_017149426.1| PREDICTED: stromelysin-3 isoform X1 [Drosophila miranda]
>gi|1036999600|ref|XP_017149427.1| PREDICTED: stromelysin-3 isoform X1 [Drosophila miranda]
>gi|1036999618|ref|XP_017149428.1| PREDICTED: stromelysin-3 isoform X2 [Drosophila miranda]
>gi|1036999636|ref|XP_017149429.1| PREDICTED: matrix metalloproteinase-14 isoform X3 [Drosophil
>gi|1036999654|ref|XP_017149430.1| PREDICTED: matrix metalloproteinase-14 isoform X4 [Drosophil
>gi|1036999670|ref|XP_017149431.1| PREDICTED: matrix metalloproteinase-14 isoform X5 [Drosophil
>gi|1036999720|ref|XP_017149432.1| PREDICTED: matrix metalloproteinase-14 isoform X6 [Drosophil
>gi|1036999758|ref|XP_017149435.1| PREDICTED: protein painting of fourth isoform X2 [Drosophila
>gi|1036999740|ref|XP_017149434.1| PREDICTED: protein painting of fourth isoform X1 [Drosophila
>gi|1036999776|ref|XP_017149436.1| PREDICTED: uncharacterized protein LOC108160139 [Drosophila
>gi|1037014318|ref|XP_017150311.1| PREDICTED: uncharacterized protein LOC108160676 [Drosophila
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>gi|1036946529|ref|XP_017146214.1| PREDICTED: odorant receptor 45b [Drosophila miranda]
>gi|1037021187|ref|XP_017150741.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila n
>gi|1036966793|ref|XP_017147424.1| PREDICTED: CCR4-NOT transcription complex subunit 1 isoform
>gi|1036966773|ref|XP_017147423.1| PREDICTED: CCR4-NOT transcription complex subunit 1 isoform
>gi|1036966831|ref|XP_017147426.1| PREDICTED: CCR4-NOT transcription complex subunit 1 isoform
>gi|1036966813|ref|XP_017147425.1| PREDICTED: CCR4-NOT transcription complex subunit 1 isoform
>gi|1036966849|ref|XP_017147427.1| PREDICTED: 5'-nucleotidase domain-containing protein 3 isofo
>gi|1036966865|ref|XP_017147428.1| PREDICTED: 5'-nucleotidase domain-containing protein 3 isofo
>gi|1036964317|ref|XP_017147277.1| PREDICTED: electron transfer flavoprotein-ubiquinone oxidore
>gi|1036964333|ref|XP_017147278.1| PREDICTED: electron transfer flavoprotein-ubiquinone oxidore
>gi|1036964391|ref|XP_017147279.1| PREDICTED: electron transfer flavoprotein-ubiquinone oxidore
>gi|1036964409|ref|XP_017147280.1| PREDICTED: myocyte-specific enhancer factor 2 [Drosophila mi
>gi|1036964535|ref|XP_017147288.1| PREDICTED: uncharacterized protein LOC108158964 [Drosophila
>gi|1036964464|ref|XP_017147283.1| PREDICTED: peptidyl-alpha-hydroxyglycine alpha-amidating lya
>gi|1036964427|ref|XP_017147281.1| PREDICTED: peptidyl-alpha-hydroxyglycine alpha-amidating lya
>gi|1036964449|ref|XP_017147282.1| PREDICTED: peptidyl-alpha-hydroxyglycine alpha-amidating lya
>gi|1036964482|ref|XP_017147284.1| PREDICTED: prenylated Rab acceptor protein 1 [Drosophila mir
>gi|1036964495|ref|XP_017147285.1| PREDICTED: prenylated Rab acceptor protein 1 [Drosophila mir
>gi|1036964515|ref|XP_017147287.1| PREDICTED: prenylated Rab acceptor protein 1 [Drosophila mir
>gi|1037006026|ref|XP_017149810.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037005986|ref|XP_017149808.1| PREDICTED: WD repeat-containing protein 89 isoform X1 [Droso
>gi|1037006008|ref|XP_017149809.1| PREDICTED: WD repeat-containing protein 89 isoform X2 [Droso
>gi|1036940823|ref|XP_017145875.1| PREDICTED: segmentation protein even-skipped [Drosophila mir
>gi|1036993278|ref|XP_017149037.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 31 isoform
>gi|1036993296|ref|XP_017149038.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 31 isoform
>gi|1037022037|ref|XP_017150794.1| PREDICTED: uncharacterized protein LOC108160997 [Drosophila
>gi|1036978439|ref|XP_017148129.1| PREDICTED: suppressor APC domain-containing protein 2 isofor
>gi|1036978455|ref|XP_017148130.1| PREDICTED: suppressor APC domain-containing protein 2 isofor
>gi|1036978513|ref|XP_017148134.1| PREDICTED: coiled-coil-helix-coiled-coil-helix domain-contai
>gi|1036978473|ref|XP_017148132.1| PREDICTED: neuralized-like protein 2 [Drosophila miranda]
>gi|1036978495|ref|XP_017148133.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1036978397|ref|XP_017148127.1| PREDICTED: epidermal growth factor receptor substrate 15-lik
>gi|1036978383|ref|XP_017148126.1| PREDICTED: epidermal growth factor receptor substrate 15-lik
>gi|1036978415|ref|XP_017148128.1| PREDICTED: epidermal growth factor receptor substrate 15-lik
>gi|1036970587|ref|XP_017147643.1| PREDICTED: histone-lysine N-methyltransferase eggless [Droso
>gi|1036970817|ref|XP_017147656.1| PREDICTED: larval cuticle protein 9 [Drosophila miranda]
>gi|1036970799|ref|XP_017147655.1| PREDICTED: uncharacterized RNA-binding protein C365.04c [Dro
>gi|1036970661|ref|XP_017147647.1| PREDICTED: tyrosine--tRNA ligase, mitochondrial [Drosophila
>gi|1036970606|ref|XP_017147644.1| PREDICTED: uncharacterized protein LOC108159157 isoform X1
>gi|1036970623|ref|XP_017147645.1| PREDICTED: uncharacterized protein LOC108159157 isoform X2
>gi|1036970641|ref|XP_017147646.1| PREDICTED: serine protease MT3772 isoform X3 [Drosophila mir
>gi|1036970785|ref|XP_017147654.1| PREDICTED: NF-kappa-B essential modulator isoform X2 [Drosop
>gi|1036970677|ref|XP_017147649.1| PREDICTED: NF-kappa-B essential modulator isoform X1 [Drosop
>gi|1036970695|ref|XP_017147650.1| PREDICTED: NF-kappa-B essential modulator isoform X1 [Drosop
>gi|1036970715|ref|XP_017147651.1| PREDICTED: NF-kappa-B essential modulator isoform X1 [Drosop
>gi|1036970749|ref|XP_017147652.1| PREDICTED: NF-kappa-B essential modulator isoform X1 [Drosop
>gi|1036970769|ref|XP_017147653.1| PREDICTED: NF-kappa-B essential modulator isoform X1 [Drosop
>gi|1037005149|ref|XP_017149751.1| PREDICTED: rhythmically expressed gene 5 protein [Drosophila
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>gi|1037005167|ref|XP_017149752.1| PREDICTED: uncharacterized protein LOC108160332 [Drosophila
>gi|1037005131|ref|XP_017149750.1| PREDICTED: origin recognition complex subunit 4 isoform X2
>gi|1037005113|ref|XP_017149749.1| PREDICTED: origin recognition complex subunit 4 isoform X1
>gi|1037021471|ref|XP_017150756.1| PREDICTED: uncharacterized protein LOC108160969 [Drosophila
>gi|1036937370|ref|XP_017145654.1| PREDICTED: uncharacterized protein LOC108158021 [Drosophila
>gi|1036943990|ref|XP_017146064.1| PREDICTED: uncharacterized protein LOC108158335 [Drosophila
>gi|1037007999|ref|XP_017149934.1| PREDICTED: homeotic protein distal-less [Drosophila miranda]
>gi|1036937388|ref|XP_017145655.1| PREDICTED: trypsin alpha-3 [Drosophila miranda]
>gi|1036999996|ref|XP_017149448.1| PREDICTED: cyclic AMP-dependent transcription factor ATF-2
>gi|1037000014|ref|XP_017149449.1| PREDICTED: cyclic AMP-dependent transcription factor ATF-2
>gi|1037000051|ref|XP_017149451.1| PREDICTED: uncharacterized protein LOC108160147 isoform X4
>gi|1037000034|ref|XP_017149450.1| PREDICTED: cyclic AMP-dependent transcription factor ATF-2
>gi|1036999982|ref|XP_017149447.1| PREDICTED: uncharacterized protein LOC108160146 [Drosophila
>gi|1037016908|ref|XP_017150463.1| PREDICTED: leptin receptor gene-related protein [Drosophila
>gi|1036989830|ref|XP_017148830.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1036989848|ref|XP_017148832.1| PREDICTED: protein PET100 homolog, mitochondrial [Drosophila
>gi|1036989810|ref|XP_017148829.1| PREDICTED: uncharacterized protein LOC108159780 [Drosophila
>gi|1036985586|ref|XP_017148574.1| PREDICTED: methylosome subunit pICln [Drosophila miranda]
>gi|1036985558|ref|XP_017148572.1| PREDICTED: uncharacterized protein LOC108159626 [Drosophila
>gi|1036985534|ref|XP_017148569.1| PREDICTED: E3 ubiquitin-protein ligase Smurf1 [Drosophila mi
>gi|1036985544|ref|XP_017148571.1| PREDICTED: NADH dehydrogenase [ubiquinone] flavoprotein 1, m
>gi|1036985572|ref|XP_017148573.1| PREDICTED: uncharacterized protein LOC108159627 [Drosophila
>gi|1036987258|ref|XP_017148678.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036987191|ref|XP_017148673.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036987209|ref|XP_017148674.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036987226|ref|XP_017148675.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036987238|ref|XP_017148677.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036987355|ref|XP_017148683.1| PREDICTED: heterochromatin protein 1-like isoform X2 [Drosop
>gi|1036987337|ref|XP_017148682.1| PREDICTED: chromo domain-containing protein cec-1-like isofo
>gi|1036987309|ref|XP_017148680.1| PREDICTED: neuropeptide CCHamide-1 receptor isoform X2 [Dros
>gi|1036987276|ref|XP_017148679.1| PREDICTED: neuropeptide CCHamide-1 receptor isoform X1 [Dros
>gi|1036987323|ref|XP_017148681.1| PREDICTED: neuropeptide CCHamide-1 receptor isoform X3 [Dros
>gi|1036990375|ref|XP_017148866.1| PREDICTED: LOW QUALITY PROTEIN: eukaryotic translation initi
>gi|1036990357|ref|XP_017148865.1| PREDICTED: TP53-regulated inhibitor of apoptosis 1-like [Dro
>gi|1036990339|ref|XP_017148863.1| PREDICTED: complement component 1 Q subcomponent-binding pro
>gi|1036989485|ref|XP_017148810.1| PREDICTED: semaphorin-1A [Drosophila miranda]
>gi|1036990660|ref|XP_017148880.1| PREDICTED: uncharacterized protein LOC108159820 isoform X2
>gi|1036990728|ref|XP_017148884.1| PREDICTED: uncharacterized protein LOC108159824 isoform X1
>gi|1036990748|ref|XP_017148885.1| PREDICTED: uncharacterized protein LOC108159824 isoform X2
>gi|1036990766|ref|XP_017148886.1| PREDICTED: uncharacterized protein LOC108159824 isoform X2
>gi|1036990787|ref|XP_017148887.1| PREDICTED: uncharacterized protein LOC108159825 [Drosophila
>gi|1036990642|ref|XP_017148879.1| PREDICTED: uncharacterized protein LOC108159820 isoform X1
>gi|1036990676|ref|XP_017148881.1| PREDICTED: protein singed wings 2 isoform X1 [Drosophila mi
>gi|1036990694|ref|XP_017148882.1| PREDICTED: protein singed wings 2 isoform X2 [Drosophila mi
>gi|1036990712|ref|XP_017148883.1| PREDICTED: cytoplasmic phosphatidylinositol transfer protei
>gi|1036991271|ref|XP_017148918.1| PREDICTED: protein FAM13A isoform X1 [Drosophila miranda]
>gi|1036991287|ref|XP_017148919.1| PREDICTED: protein FAM13A isoform X2 [Drosophila miranda]
>gi|1036991303|ref|XP_017148920.1| PREDICTED: uncharacterized protein LOC108159840 [Drosophila
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>gi|1036987829|ref|XP_017148710.1| PREDICTED: intersectin-1 [Drosophila miranda]
>gi|1036987809|ref|XP_017148709.1| PREDICTED: uncharacterized protein LOC108159705 [Drosophila
>gi|1036987793|ref|XP_017148708.1| PREDICTED: ataxin-10 [Drosophila miranda]
>gi|1036987725|ref|XP_017148704.1| PREDICTED: uncharacterized protein LOC108159703 isoform X1
>gi|1036987741|ref|XP_017148705.1| PREDICTED: uncharacterized protein LOC108159703 isoform X1
>gi|1036987758|ref|XP_017148706.1| PREDICTED: uncharacterized protein LOC108159703 isoform X1
>gi|1036987775|ref|XP_017148707.1| PREDICTED: uncharacterized protein LOC108159703 isoform X2
>gi|1036988963|ref|XP_017148777.1| PREDICTED: heterogeneous nuclear ribonucleoprotein U-like pr
>gi|1036988997|ref|XP_017148779.1| PREDICTED: heterogeneous nuclear ribonucleoprotein U-like pr
>gi|1036989013|ref|XP_017148780.1| PREDICTED: heterogeneous nuclear ribonucleoprotein U isoform
>gi|1036988987|ref|XP_017148778.1| PREDICTED: heterogeneous nuclear ribonucleoprotein U isoform
>gi|1036989029|ref|XP_017148781.1| PREDICTED: heterogeneous nuclear ribonucleoprotein U isoform
>gi|1036989047|ref|XP_017148783.1| PREDICTED: heterogeneous nuclear ribonucleoprotein U-like pr
>gi|1036991647|ref|XP_017148934.1| PREDICTED: serine palmitoyltransferase 1 [Drosophila miranda
>gi|1036991665|ref|XP_017148935.1| PREDICTED: serine palmitoyltransferase 1 [Drosophila miranda
>gi|1036991625|ref|XP_017148933.1| PREDICTED: 1,4-alpha-glucan-branching enzyme [Drosophila mir
>gi|1036991683|ref|XP_017148936.1| PREDICTED: uncharacterized protein LOC108159854 isoform X1
>gi|1036991699|ref|XP_017148937.1| PREDICTED: uncharacterized protein LOC108159854 isoform X1
>gi|1036991723|ref|XP_017148938.1| PREDICTED: uncharacterized protein LOC108159854 isoform X1
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>gi|1036991773|ref|XP_017148941.1| PREDICTED: uncharacterized protein LOC108159854 isoform X1
>gi|1036991793|ref|XP_017148942.1| PREDICTED: uncharacterized protein LOC108159854 isoform X2
>gi|1036945960|ref|XP_017146178.1| PREDICTED: uncharacterized protein LOC108158417 isoform X3
>gi|1036945942|ref|XP_017146177.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X2
>gi|1036945890|ref|XP_017146174.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X1
>gi|1036945908|ref|XP_017146175.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X1
>gi|1036945926|ref|XP_017146176.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X1
>gi|1036945978|ref|XP_017146179.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X4
>gi|1036945994|ref|XP_017146180.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X5
>gi|1036946012|ref|XP_017146181.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X5
>gi|1036946030|ref|XP_017146182.1| PREDICTED: probable lysine-specific demethylase 4B isoform X
>gi|1036946101|ref|XP_017146186.1| PREDICTED: valine--tRNA ligase [Drosophila miranda]
>gi|1036946135|ref|XP_017146189.1| PREDICTED: origin recognition complex subunit 3 [Drosophila
>gi|1036946256|ref|XP_017146197.1| PREDICTED: uncharacterized protein LOC108158426 isoform X2
>gi|1036946240|ref|XP_017146195.1| PREDICTED: uncharacterized protein LOC108158426 isoform X1
>gi|1036946048|ref|XP_017146183.1| PREDICTED: uncharacterized protein LOC108158418 isoform X1
>gi|1036946084|ref|XP_017146185.1| PREDICTED: serine/arginine repetitive matrix protein 5 isofo
>gi|1036946066|ref|XP_017146184.1| PREDICTED: uncharacterized protein LOC108158418 isoform X2
>gi|1036946119|ref|XP_017146188.1| PREDICTED: WD repeat-containing protein 60 [Drosophila mira
>gi|1036946205|ref|XP_017146193.1| PREDICTED: transferrin [Drosophila miranda]
>gi|1036946153|ref|XP_017146190.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036946170|ref|XP_017146191.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036946188|ref|XP_017146192.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036946291|ref|XP_017146199.1| PREDICTED: lysozyme P [Drosophila miranda]
>gi|1036946308|ref|XP_017146200.1| PREDICTED: uncharacterized protein LOC108158429 isoform X1
>gi|1036946326|ref|XP_017146201.1| PREDICTED: uncharacterized protein LOC108158429 isoform X2
>gi|1036946222|ref|XP_017146194.1| PREDICTED: trypsin 3A1 [Drosophila miranda]
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>gi|1036946274|ref|XP_017146198.1| PREDICTED: cuticle protein 16.8 [Drosophila miranda]
>gi|1036937406|ref|XP_017145656.1| PREDICTED: general odorant-binding protein 57e-like [Drosopl
>gi|1036942630|ref|XP_017145981.1| PREDICTED: gram-negative bacteria-binding protein 3 [Drosopl
>gi|1037011979|ref|XP_017150166.1| PREDICTED: actin cytoskeleton-regulatory complex protein pal
>gi|1037011991|ref|XP_017150167.1| PREDICTED: actin cytoskeleton-regulatory complex protein pal
>gi|1037012007|ref|XP_017150168.1| PREDICTED: uncharacterized protein LOC108160585 [Drosophila
>gi|1037011957|ref|XP_017150165.1| PREDICTED: bleomycin hydrolase [Drosophila miranda]
>gi|1037019494|ref|XP_017150633.1| PREDICTED: uncharacterized protein LOC108160877 isoform X2
>gi|1037019478|ref|XP_017150632.1| PREDICTED: uncharacterized protein LOC108160877 isoform X1
>gi|1037019460|ref|XP_017150631.1| PREDICTED: uncharacterized protein LOC108160876 [Drosophila
>gi|1036942939|ref|XP_017146000.1| PREDICTED: uncharacterized protein LOC108158275 [Drosophila
>gi|1036943202|ref|XP_017146016.1| PREDICTED: keratin, type I cytoskeletal 9 [Drosophila mirand
>gi|1036942788|ref|XP_017145992.1| PREDICTED: gram-negative bacteria-binding protein 3 [Drosopl
>gi|1036944173|ref|XP_017146075.1| PREDICTED: protein transport protein SSS1 [Drosophila mirand
>gi|1036945024|ref|XP_017146121.1| PREDICTED: general odorant-binding protein 57c [Drosophila r
>gi|1036943458|ref|XP_017146032.1| PREDICTED: uncharacterized protein LOC108158303 [Drosophila
>gi|1036943350|ref|XP_017146026.1| PREDICTED: uncharacterized protein LOC108158296 [Drosophila
>gi|1037018687|ref|XP_017150581.1| PREDICTED: uncharacterized protein LOC108160840 isoform X6
>gi|1037018654|ref|XP_017150579.1| PREDICTED: uncharacterized protein LOC108160840 isoform X4
>gi|1037018640|ref|XP_017150578.1| PREDICTED: uncharacterized protein LOC108160840 isoform X3
>gi|1037018671|ref|XP_017150580.1| PREDICTED: uncharacterized protein LOC108160840 isoform X5
>gi|1037018623|ref|XP_017150577.1| PREDICTED: uncharacterized protein LOC108160840 isoform X2
>gi|1037018703|ref|XP_017150582.1| PREDICTED: uncharacterized protein LOC108160840 isoform X7
>gi|1037018569|ref|XP_017150574.1| PREDICTED: uncharacterized protein LOC108160840 isoform X1
>gi|1037018587|ref|XP_017150575.1| PREDICTED: uncharacterized protein LOC108160840 isoform X1
>gi|1037018605|ref|XP_017150576.1| PREDICTED: uncharacterized protein LOC108160840 isoform X1
>gi|1036937422|ref|XP_017145657.1| PREDICTED: trypsin beta [Drosophila miranda]
>gi|1036943602|ref|XP_017146041.1| PREDICTED: uncharacterized protein LOC108158312 [Drosophila
>gi|1037020811|ref|XP_017150718.1| PREDICTED: uncharacterized protein LOC108160941 [Drosophila
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>gi|1036944105|ref|XP_017146071.1| PREDICTED: uncharacterized protein LOC108158343 [Drosophila
>gi|1036941786|ref|XP_017145925.1| PREDICTED: uncharacterized protein LOC108158213 [Drosophila
>gi|1036979660|ref|XP_017148205.1| PREDICTED: basic salivary proline-rich protein 1 isoform X2
>gi|1036979644|ref|XP_017148204.1| PREDICTED: basic salivary proline-rich protein 1 isoform X1
>gi|1036979678|ref|XP_017148206.1| PREDICTED: basic salivary proline-rich protein 1 isoform X3
>gi|1036979695|ref|XP_017148207.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036979586|ref|XP_017148200.1| PREDICTED: tudor domain-containing protein 7B isoform X1 [Di
>gi|1036979602|ref|XP_017148201.1| PREDICTED: tudor domain-containing protein 7B isoform X2 [Di
>gi|1036979618|ref|XP_017148202.1| PREDICTED: uncharacterized protein LOC108159436 isoform X3
>gi|1036979632|ref|XP_017148203.1| PREDICTED: uncharacterized protein LOC108159436 isoform X4
>gi|1037017157|ref|XP_017150480.1| PREDICTED: leukocyte antigen CD37 isoform X2 [Drosophila mil
>gi|1037017143|ref|XP_017150479.1| PREDICTED: tetraspanin-8 isoform X1 [Drosophila miranda]
>gi|1037007981|ref|XP_017149933.1| PREDICTED: protein krueppel [Drosophila miranda]
>gi|1036980013|ref|XP_017148230.1| PREDICTED: MORN repeat-containing protein 3 [Drosophila mira
>gi|1036980033|ref|XP_017148231.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1036980002|ref|XP_017148229.1| PREDICTED: Fanconi anemia group I protein [Drosophila mirand
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>gi|1036980049|ref|XP_017148232.1| PREDICTED: uncharacterized protein LOC108159452 [Drosophila
>gi|1036980070|ref|XP_017148233.1| PREDICTED: uncharacterized protein LOC108159453 [Drosophila
>gi|1037011398|ref|XP_017150132.1| PREDICTED: uncharacterized protein LOC108160564 isoform X2
>gi|1037011381|ref|XP_017150130.1| PREDICTED: uncharacterized protein LOC108160564 isoform X1
>gi|1037011415|ref|XP_017150133.1| PREDICTED: uncharacterized protein LOC108160564 isoform X3
>gi|1037021923|ref|XP_017150786.1| PREDICTED: ejaculatory bulb-specific protein 1 [Drosophila r
>gi|1036944088|ref|XP_017146069.1| PREDICTED: keratin, type I cytoskeletal 14-like [Drosophila
>gi|1036942615|ref|XP_017145980.1| PREDICTED: uncharacterized protein LOC108158254 [Drosophila
>gi|1036943908|ref|XP_017146059.1| PREDICTED: glutamic acid-rich protein [Drosophila miranda]
>gi|1036991483|ref|XP_017148924.1| PREDICTED: protein jim lovell [Drosophila miranda]
>gi|1036991497|ref|XP_017148925.1| PREDICTED: protein goliath [Drosophila miranda]
>gi|1036991517|ref|XP_017148926.1| PREDICTED: protein goliath [Drosophila miranda]
>gi|1037015968|ref|XP_017150406.1| PREDICTED: protein gooseberry [Drosophila miranda]
>gi|1037014868|ref|XP_017150345.1| PREDICTED: protein gooseberry-neuro [Drosophila miranda]
>gi|1037019410|ref|XP_017150628.1| PREDICTED: neuropeptide-like 1 isoform X1 [Drosophila mirand
>gi|1037019444|ref|XP_017150630.1| PREDICTED: neuropeptide-like 1 isoform X3 [Drosophila mirand
>gi|1037019426|ref|XP_017150629.1| PREDICTED: neuropeptide-like 1 isoform X2 [Drosophila mirand
>gi|1036988873|ref|XP_017148771.1| PREDICTED: protein unzipped [Drosophila miranda]
>gi|1036988888|ref|XP_017148772.1| PREDICTED: protein unzipped [Drosophila miranda]
>gi|1036988904|ref|XP_017148773.1| PREDICTED: protein lin-37 homolog [Drosophila miranda]
>gi|1036988917|ref|XP_017148774.1| PREDICTED: 12 kDa FK506-binding protein [Drosophila miranda]
>gi|1036992061|ref|XP_017148959.1| PREDICTED: zinc finger protein 33A isoform X1 [Drosophila mi
>gi|1036992079|ref|XP_017148960.1| PREDICTED: zinc finger protein 33A isoform X2 [Drosophila mi
>gi|1036992043|ref|XP_017148958.1| PREDICTED: longitudinals lacking protein, isoforms J/P/Q/S/Z
>gi|1037018171|ref|XP_017150545.1| PREDICTED: protein windbeutel [Drosophila miranda]
>gi|1036989914|ref|XP_017148836.1| PREDICTED: protein 4.1 homolog isoform X3 [Drosophila mirand
>gi|1036989896|ref|XP_017148835.1| PREDICTED: protein 4.1 homolog isoform X2 [Drosophila mirand
>gi|1036989932|ref|XP_017148837.1| PREDICTED: protein 4.1 homolog isoform X4 [Drosophila mirand
>gi|1036989866|ref|XP_017148833.1| PREDICTED: protein 4.1 homolog isoform X1 [Drosophila mirand
>gi|1036989884|ref|XP_017148834.1| PREDICTED: protein 4.1 homolog isoform X1 [Drosophila mirand
>gi|1036989949|ref|XP_017148838.1| PREDICTED: ribosomal RNA-processing protein 8 [Drosophila mi
>gi|1036957989|ref|XP_017146908.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957961|ref|XP_017146905.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957927|ref|XP_017146903.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957975|ref|XP_017146907.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957943|ref|XP_017146904.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957843|ref|XP_017146898.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957857|ref|XP_017146899.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957875|ref|XP_017146900.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957895|ref|XP_017146901.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036957913|ref|XP_017146902.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036958153|ref|XP_017146915.1| PREDICTED: uncharacterized protein LOC108158777 isoform X6
>gi|1036958165|ref|XP_017146917.1| PREDICTED: uncharacterized protein LOC108158777 isoform X7
>gi|1036958326|ref|XP_017146927.1| PREDICTED: uncharacterized protein LOC108158777 isoform X16
>gi|1036958223|ref|XP_017146921.1| PREDICTED: uncharacterized protein LOC108158777 isoform X11
>gi|1036958120|ref|XP_017146913.1| PREDICTED: uncharacterized protein LOC108158777 isoform X4
>gi|1036958381|ref|XP_017146930.1| PREDICTED: uncharacterized protein LOC108158777 isoform X19
>gi|1036958090|ref|XP_017146912.1| PREDICTED: uncharacterized protein LOC108158777 isoform X3
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>gi|1036958306|ref|XP_017146925.1| PREDICTED: uncharacterized protein LOC108158777 isoform X15
>gi|1036958360|ref|XP_017146929.1| PREDICTED: uncharacterized protein LOC108158777 isoform X18
>gi|1036958289|ref|XP_017146924.1| PREDICTED: uncharacterized protein LOC108158777 isoform X14
>gi|1036958256|ref|XP_017146923.1| PREDICTED: uncharacterized protein LOC108158777 isoform X13
>gi|1036958193|ref|XP_017146919.1| PREDICTED: uncharacterized protein LOC108158777 isoform X9
>gi|1036958177|ref|XP_017146918.1| PREDICTED: uncharacterized protein LOC108158777 isoform X8
>gi|1036958004|ref|XP_017146909.1| PREDICTED: uncharacterized protein LOC108158777 isoform X1
>gi|1036958042|ref|XP_017146911.1| PREDICTED: uncharacterized protein LOC108158777 isoform X1
>gi|1036958211|ref|XP_017146920.1| PREDICTED: uncharacterized protein LOC108158777 isoform X10
>gi|1036958343|ref|XP_017146928.1| PREDICTED: uncharacterized protein LOC108158777 isoform X17
>gi|1036958240|ref|XP_017146922.1| PREDICTED: uncharacterized protein LOC108158777 isoform X12
>gi|1036958140|ref|XP_017146914.1| PREDICTED: uncharacterized protein LOC108158777 isoform X5
>gi|1036958435|ref|XP_017146931.1| PREDICTED: uncharacterized protein LOC108158777 isoform X20
>gi|1036958024|ref|XP_017146910.1| PREDICTED: uncharacterized protein LOC108158777 isoform X2
>gi|1036958584|ref|XP_017146939.1| PREDICTED: protein enabled isoform X7 [Drosophila miranda]
>gi|1036958596|ref|XP_017146940.1| PREDICTED: protein enabled isoform X7 [Drosophila miranda]
>gi|1036958626|ref|XP_017146941.1| PREDICTED: protein enabled isoform X8 [Drosophila miranda]
>gi|1036958566|ref|XP_017146938.1| PREDICTED: protein enabled isoform X6 [Drosophila miranda]
>gi|1036958534|ref|XP_017146935.1| PREDICTED: protein enabled isoform X4 [Drosophila miranda]
>gi|1036958518|ref|XP_017146934.1| PREDICTED: protein enabled isoform X3 [Drosophila miranda]
>gi|1036958496|ref|XP_017146933.1| PREDICTED: protein enabled isoform X2 [Drosophila miranda]
>gi|1036958550|ref|XP_017146936.1| PREDICTED: protein enabled isoform X5 [Drosophila miranda]
>gi|1036958478|ref|XP_017146932.1| PREDICTED: protein enabled isoform X1 [Drosophila miranda]
>gi|1037022445|ref|XP_017150818.1| PREDICTED: uncharacterized protein LOC108161017 [Drosophila
>gi|1037004784|ref|XP_017149729.1| PREDICTED: uncharacterized protein LOC108160316, partial [D
>gi|1037012405|ref|XP_017150192.1| PREDICTED: dehydrogenase/reductase SDR family member on chr
>gi|1037012421|ref|XP_017150193.1| PREDICTED: dehydrogenase/reductase SDR family member on chr
>gi|1037012439|ref|XP_017150195.1| PREDICTED: dehydrogenase/reductase SDR family member on chr
>gi|1037012455|ref|XP_017150196.1| PREDICTED: dehydrogenase/reductase SDR family member on chr
>gi|1037018137|ref|XP_017150543.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1037018155|ref|XP_017150544.1| PREDICTED: casein kinase II subunit beta' [Drosophila miran
>gi|1036945472|ref|XP_017146148.1| PREDICTED: uncharacterized protein LOC108158402 [Drosophila
>gi|1036992498|ref|XP_017148987.1| PREDICTED: VWFA and cache domain-containing protein CG16868
>gi|1036997140|ref|XP_017149277.1| PREDICTED: phosphate carrier protein, mitochondrial [Drosop
>gi|1036997107|ref|XP_017149275.1| PREDICTED: ATP-dependent RNA helicase DDX24 [Drosophila mir
>gi|1036997123|ref|XP_017149276.1| PREDICTED: sister chromatid cohesion protein DCC1 [Drosophil
>gi|1037021091|ref|XP_017150737.1| PREDICTED: homeobox protein unplugged [Drosophila miranda]
>gi|1037010978|ref|XP_017150105.1| PREDICTED: N-acetylglucosamine-1-phosphotransferase subunits
>gi|1036958797|ref|XP_017146951.1| PREDICTED: helicase domino isoform X2 [Drosophila miranda]
>gi|1036958824|ref|XP_017146952.1| PREDICTED: helicase domino isoform X3 [Drosophila miranda]
>gi|1036958779|ref|XP_017146950.1| PREDICTED: helicase domino isoform X1 [Drosophila miranda]
>gi|1037001857|ref|XP_017149562.1| PREDICTED: tubulin beta-1 chain [Drosophila miranda]
>gi|1037008223|ref|XP_017149949.1| PREDICTED: transcription elongation factor B polypeptide 1
>gi|1037008187|ref|XP_017149947.1| PREDICTED: LOW QUALITY PROTEIN: ADP-ribosylation factor-like
>gi|1037008205|ref|XP_017149948.1| PREDICTED: 60S ribosomal protein L11 [Drosophila miranda]
>gi|1036958764|ref|XP_017146949.1| PREDICTED: NF-kappa-B inhibitor-interacting Ras-like protei
>gi|1036958679|ref|XP_017146943.1| PREDICTED: far upstream element-binding protein 3 [Drosophil
>gi|1036958712|ref|XP_017146944.1| PREDICTED: protein DEK isoform X1 [Drosophila miranda]
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>gi|1036958728|ref|XP_017146947.1| PREDICTED: protein DEK isoform X2 [Drosophila miranda]
>gi|1036958746|ref|XP_017146948.1| PREDICTED: flap endonuclease 1 [Drosophila miranda]
>gi|1036958642|ref|XP_017146942.1| PREDICTED: integrator complex subunit 8 [Drosophila miranda]
>gi|1036945094|ref|XP_017146126.1| PREDICTED: uncharacterized protein LOC108158383 [Drosophila
>gi|1037018534|ref|XP_017150571.1| PREDICTED: uncharacterized protein LOC108160838 isoform X2
>gi|1037018516|ref|XP_017150570.1| PREDICTED: uncharacterized protein LOC108160838 isoform X1
>gi|1037021372|ref|XP_017150753.1| PREDICTED: uncharacterized protein LOC108160966 isoform X1
>gi|1037021390|ref|XP_017150754.1| PREDICTED: uncharacterized protein LOC108160966 isoform X2
>gi|1036995674|ref|XP_017149185.1| PREDICTED: uncharacterized protein LOC108159988 [Drosophila
>gi|1036995692|ref|XP_017149186.1| PREDICTED: uncharacterized protein LOC108159989 [Drosophila
>gi|1036995613|ref|XP_017149181.1| PREDICTED: ras-specific guanine nucleotide-releasing factor
>gi|1036995631|ref|XP_017149183.1| PREDICTED: ras-specific guanine nucleotide-releasing factor
>gi|1036995656|ref|XP_017149184.1| PREDICTED: ras-specific guanine nucleotide-releasing factor
>gi|1036971413|ref|XP_017147690.1| PREDICTED: probable ascorbate-specific transmembrane electro
>gi|1036971433|ref|XP_017147691.1| PREDICTED: probable ascorbate-specific transmembrane electro
>gi|1036971451|ref|XP_017147692.1| PREDICTED: probable ascorbate-specific transmembrane electro
>gi|1036971469|ref|XP_017147693.1| PREDICTED: probable ascorbate-specific transmembrane electro
>gi|1036971325|ref|XP_017147685.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971289|ref|XP_017147683.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971361|ref|XP_017147687.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971397|ref|XP_017147689.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971309|ref|XP_017147684.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971267|ref|XP_017147682.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971379|ref|XP_017147688.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971345|ref|XP_017147686.1| PREDICTED: glutaminase kidney isoform, mitochondrial isoform
>gi|1036971251|ref|XP_017147681.1| PREDICTED: nucleoporin NUP188 homolog isoform X2 [Drosophila
>gi|1036971233|ref|XP_017147680.1| PREDICTED: nucleoporin NUP188 homolog isoform X1 [Drosophila
>gi|1036941097|ref|XP_017145885.1| PREDICTED: putative uncharacterized protein DDB_G0294196 [D
>gi|1037012819|ref|XP_017150220.1| PREDICTED: uncharacterized protein LOC108160615 [Drosophila
>gi|1037012851|ref|XP_017150222.1| PREDICTED: general odorant-binding protein 67 [Drosophila m
>gi|1037012837|ref|XP_017150221.1| PREDICTED: epimerase family protein SDR39U1 [Drosophila mira
>gi|1036937440|ref|XP_017145658.1| PREDICTED: uncharacterized protein LOC108158026 [Drosophila
>gi|1036962932|ref|XP_017147196.1| PREDICTED: uncharacterized protein LOC108158912 isoform X1
>gi|1036962950|ref|XP_017147198.1| PREDICTED: uncharacterized protein LOC108158912 isoform X2
>gi|1036962965|ref|XP_017147199.1| PREDICTED: uncharacterized protein LOC108158912 isoform X3
>gi|1036963006|ref|XP_017147201.1| PREDICTED: LOW QUALITY PROTEIN: proto-oncogene serine/threor
>gi|1036963024|ref|XP_017147202.1| PREDICTED: tafazzin homolog [Drosophila miranda]
>gi|1036963058|ref|XP_017147204.1| PREDICTED: 39S ribosomal protein L18, mitochondrial [Drosopl
>gi|1036963212|ref|XP_017147215.1| PREDICTED: cytochrome b-c1 complex subunit 9 [Drosophila mi
>gi|1036963158|ref|XP_017147212.1| PREDICTED: uncharacterized protein LOC108158920 isoform X3
>gi|1036963108|ref|XP_017147208.1| PREDICTED: uncharacterized protein LOC108158920 isoform X1
>gi|1036963126|ref|XP_017147209.1| PREDICTED: uncharacterized protein LOC108158920 isoform X1
>gi|1036963144|ref|XP_017147211.1| PREDICTED: uncharacterized protein LOC108158920 isoform X2
>gi|1036963176|ref|XP_017147213.1| PREDICTED: uncharacterized protein LOC108158921 [Drosophila
>gi|1036962990|ref|XP_017147200.1| PREDICTED: NADPH:adrenodoxin oxidoreductase, mitochondrial
>gi|1036963194|ref|XP_017147214.1| PREDICTED: uncharacterized protein LOC108158922 [Drosophila
>gi|1036963042|ref|XP_017147203.1| PREDICTED: UPF0605 protein GA14893 [Drosophila miranda]
>gi|1036962895|ref|XP_017147194.1| PREDICTED: F-box/LRR-repeat protein 6 [Drosophila miranda]
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>gi|1036962913|ref|XP_017147195.1| PREDICTED: F-box/LRR-repeat protein 6 [Drosophila miranda]
>gi|1036963074|ref|XP_017147206.1| PREDICTED: uncharacterized protein LOC108158919 isoform X1
>gi|1036963090|ref|XP_017147207.1| PREDICTED: uncharacterized protein LOC108158919 isoform X2
>gi|1036990987|ref|XP_017148900.1| PREDICTED: WD repeat-containing protein 48 homolog [Drosoph:
>gi|1036991001|ref|XP_017148901.1| PREDICTED: WD repeat-containing protein 48 homolog [Drosoph:
>gi|1036991197|ref|XP_017148914.1| PREDICTED: protein Vhl [Drosophila miranda]
>gi|1036991235|ref|XP_017148916.1| PREDICTED: trafficking protein particle complex subunit 2-l:
>gi|1036991253|ref|XP_017148917.1| PREDICTED: cuticular protein 47Eg [Drosophila miranda]
>gi|1036991217|ref|XP_017148915.1| PREDICTED: troponin C [Drosophila miranda]
>gi|1036991014|ref|XP_017148902.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1036991040|ref|XP_017148904.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1036991026|ref|XP_017148903.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1036991068|ref|XP_017148906.1| PREDICTED: glycine-rich cell wall structural protein 1 isofc
>gi|1036991114|ref|XP_017148909.1| PREDICTED: pupal cuticle protein 36 isoform X8 [Drosophila r
>gi|1036991054|ref|XP_017148905.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1036991146|ref|XP_017148911.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1036991098|ref|XP_017148908.1| PREDICTED: glycine-rich cell wall structural protein 1 isofc
>gi|1036991082|ref|XP_017148907.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1036991128|ref|XP_017148910.1| PREDICTED: pupal cuticle protein 36 isoform X9 [Drosophila r
>gi|1036991158|ref|XP_017148912.1| PREDICTED: pupal cuticle protein 20 isoform X11 [Drosophila
>gi|1036991177|ref|XP_017148913.1| PREDICTED: histone-lysine N-methyltransferase 2D [Drosophila
>gi|1036944123|ref|XP_017146072.1| PREDICTED: larval cuticle protein LCP-17 [Drosophila miranda
>gi|1036943737|ref|XP_017146049.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036945042|ref|XP_017146123.1| PREDICTED: sodium/potassium/calcium exchanger 6, mitochondr:
>gi|1036942986|ref|XP_017146003.1| PREDICTED: uncharacterized protein LOC108158279 [Drosophila
>gi|1037018765|ref|XP_017150587.1| PREDICTED: cuticle protein CP14.6 [Drosophila miranda]
>gi|1037018747|ref|XP_017150586.1| PREDICTED: odorant receptor 47a [Drosophila miranda]
>gi|1037018783|ref|XP_017150588.1| PREDICTED: neuropeptide-like protein 32 [Drosophila miranda]
>gi|1037001342|ref|XP_017149527.1| PREDICTED: glycine-rich RNA-binding protein 2-like [Drosoph:
>gi|1037001320|ref|XP_017149526.1| PREDICTED: glycine-rich protein 3 [Drosophila miranda]
>gi|1037001156|ref|XP_017149515.1| PREDICTED: prisilkin-39 [Drosophila miranda]
>gi|1037001208|ref|XP_017149518.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1037001222|ref|XP_017149519.1| PREDICTED: glycine-rich cell wall structural protein isoforr
>gi|1037001174|ref|XP_017149516.1| PREDICTED: uncharacterized protein LOC108160193 [Drosophila
>gi|1037001138|ref|XP_017149514.1| PREDICTED: annexin A7 [Drosophila miranda]
>gi|1037001190|ref|XP_017149517.1| PREDICTED: uncharacterized protein LOC108160194 [Drosophila
>gi|1037001292|ref|XP_017149524.1| PREDICTED: glycine-rich cell wall structural protein 1 [Dros
>gi|1037001310|ref|XP_017149525.1| PREDICTED: uncharacterized protein LOC108160201 [Drosophila
>gi|1037001272|ref|XP_017149522.1| PREDICTED: uncharacterized protein F12A10.7 [Drosophila mira
>gi|1037001236|ref|XP_017149520.1| PREDICTED: shematrin-like protein 2 [Drosophila miranda]
>gi|1037001254|ref|XP_017149521.1| PREDICTED: glycine-rich protein 5 [Drosophila miranda]
>gi|1037000121|ref|XP_017149456.1| PREDICTED: protein transport protein Sec24A isoform X2 [Dros
>gi|1037000104|ref|XP_017149455.1| PREDICTED: protein transport protein Sec24A isoform X1 [Dros
>gi|1036997742|ref|XP_017149313.1| PREDICTED: oxysterol-binding protein-related protein 9 isofc
>gi|1036997760|ref|XP_017149314.1| PREDICTED: oxysterol-binding protein-related protein 9 isofc
>gi|1036997778|ref|XP_017149315.1| PREDICTED: uncharacterized protein LOC108160072 isoform X1
>gi|1036997796|ref|XP_017149316.1| PREDICTED: uncharacterized protein LOC108160072 isoform X2
>gi|1036997812|ref|XP_017149317.1| PREDICTED: elongation of very long chain fatty acids protein
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>gi|1037016126|ref|XP_017150416.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037016143|ref|XP_017150417.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037019598|ref|XP_017150639.1| PREDICTED: FMRFamide-related peptides [Drosophila miranda]
>gi|1036999277|ref|XP_017149408.1| PREDICTED: cathepsin L [Drosophila miranda]
>gi|1036999301|ref|XP_017149409.1| PREDICTED: uncharacterized protein LOC108160126 [Drosophila
>gi|1036981133|ref|XP_017148300.1| PREDICTED: protein LTV1 homolog [Drosophila miranda]
>gi|1036981151|ref|XP_017148301.1| PREDICTED: glycine receptor subunit alpha-2 isoform X1 [Dros
>gi|1036981167|ref|XP_017148302.1| PREDICTED: glycine receptor subunit alpha-2 isoform X2 [Dros
>gi|1036981098|ref|XP_017148298.1| PREDICTED: periodic tryptophan protein 2 homolog [Drosophila
>gi|1036981185|ref|XP_017148303.1| PREDICTED: pre-mRNA-splicing factor Syf2 [Drosophila miranda
>gi|1037012073|ref|XP_017150173.1| PREDICTED: U7 snRNA-associated Sm-like protein LSm10 [Drosop
>gi|1037012058|ref|XP_017150172.1| PREDICTED: proteasome subunit beta type-5 [Drosophila mirand
>gi|1036994604|ref|XP_017149119.1| PREDICTED: uncharacterized protein LOC108159960 [Drosophila
>gi|1036994621|ref|XP_017149121.1| PREDICTED: desumoylating isopeptidase 2 [Drosophila miranda]
>gi|1037002368|ref|XP_017149595.1| PREDICTED: transmembrane protein 170A [Drosophila miranda]
>gi|1037002360|ref|XP_017149594.1| PREDICTED: crossover junction endonuclease EME1 [Drosophila
>gi|1036986602|ref|XP_017148636.1| PREDICTED: H/ACA ribonucleoprotein complex subunit 3 [Drosop
>gi|1036986584|ref|XP_017148635.1| PREDICTED: transmembrane protein 223 [Drosophila miranda]
>gi|1036986546|ref|XP_017148633.1| PREDICTED: activating transcription factor 7-interacting pro
>gi|1036986564|ref|XP_017148634.1| PREDICTED: activating transcription factor 7-interacting pro
>gi|1036988101|ref|XP_017148725.1| PREDICTED: la-related protein CG11505 [Drosophila miranda]
>gi|1036985448|ref|XP_017148564.1| PREDICTED: CDK-activating kinase assembly factor MAT1 [Droso
>gi|1036985520|ref|XP_017148568.1| PREDICTED: ubiquitin-conjugating enzyme E2 W isoform X2 [Dro
>gi|1036985488|ref|XP_017148566.1| PREDICTED: ubiquitin-conjugating enzyme E2 W isoform X1 [Dro
>gi|1036985504|ref|XP_017148567.1| PREDICTED: ubiquitin-conjugating enzyme E2 W isoform X1 [Dro
>gi|1036985430|ref|XP_017148563.1| PREDICTED: D-aspartate oxidase [Drosophila miranda]
>gi|1036985412|ref|XP_017148562.1| PREDICTED: uncharacterized protein LOC108159618 [Drosophila
>gi|1036985464|ref|XP_017148565.1| PREDICTED: ras-related protein Rab-3 [Drosophila miranda]
>gi|1036994288|ref|XP_017149099.1| PREDICTED: protocadherin-like wing polarity protein stan iso
>gi|1036994326|ref|XP_017149101.1| PREDICTED: protocadherin-like wing polarity protein stan iso
>gi|1036994310|ref|XP_017149100.1| PREDICTED: protocadherin-like wing polarity protein stan iso
>gi|1037008170|ref|XP_017149946.1| PREDICTED: uncharacterized protein LOC108160446 [Drosophila
>gi|1037008150|ref|XP_017149944.1| PREDICTED: probable elongator complex protein 2 [Drosophila
>gi|1036939704|ref|XP_017145803.1| PREDICTED: ARF GTPase-activating protein GIT2 isoform X1 [Dr
>gi|1036939723|ref|XP_017145804.1| PREDICTED: ARF GTPase-activating protein GIT2 isoform X2 [Dr
>gi|1036939652|ref|XP_017145800.1| PREDICTED: mannosylglucosyl-3-phosphoglycerate phosphatase
>gi|1036939668|ref|XP_017145801.1| PREDICTED: mannosylglucosyl-3-phosphoglycerate phosphatase
>gi|1036939686|ref|XP_017145802.1| PREDICTED: mannosylglucosyl-3-phosphoglycerate phosphatase
>gi|1036939741|ref|XP_017145805.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1036939860|ref|XP_017145812.1| PREDICTED: longitudinals lacking protein isoform X3 [Drosopl
>gi|1036940006|ref|XP_017145823.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939938|ref|XP_017145818.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939954|ref|XP_017145819.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939906|ref|XP_017145816.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939842|ref|XP_017145811.1| PREDICTED: longitudinals lacking protein, isoforms J/P/Q/S/Z
>gi|1036940146|ref|XP_017145832.1| PREDICTED: longitudinals lacking protein, isoforms J/P/Q/S/Z
>gi|1036939876|ref|XP_017145813.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036939770|ref|XP_017145806.1| PREDICTED: longitudinals lacking protein, isoforms F/I/K/T
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>gi|1036939788|ref|XP_017145807.1| PREDICTED: longitudinals lacking protein, isoforms F/I/K/T
>gi|1036939806|ref|XP_017145809.1| PREDICTED: longitudinals lacking protein, isoforms F/I/K/T
>gi|1036939824|ref|XP_017145810.1| PREDICTED: longitudinals lacking protein, isoforms F/I/K/T
>gi|1036940095|ref|XP_017145828.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036940059|ref|XP_017145826.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036940077|ref|XP_017145827.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036940112|ref|XP_017145829.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036940042|ref|XP_017145825.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939888|ref|XP_017145815.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939970|ref|XP_017145821.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939924|ref|XP_017145817.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036940024|ref|XP_017145824.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036940129|ref|XP_017145830.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036939988|ref|XP_017145822.1| PREDICTED: longitudinals lacking protein, isoforms H/M/V iso
>gi|1036940307|ref|XP_017145842.1| PREDICTED: longitudinals lacking protein, isoforms N/O/W/X/
>gi|1036940274|ref|XP_017145839.1| PREDICTED: peflin [Drosophila miranda]
>gi|1036940256|ref|XP_017145838.1| PREDICTED: translin [Drosophila miranda]
>gi|1036940164|ref|XP_017145833.1| PREDICTED: carnitine O-palmitoyltransferase 1, liver isoform
>gi|1036940184|ref|XP_017145834.1| PREDICTED: carnitine O-palmitoyltransferase 1, liver isoform
>gi|1036940202|ref|XP_017145835.1| PREDICTED: carnitine O-palmitoyltransferase 1, liver isoform
>gi|1036940238|ref|XP_017145837.1| PREDICTED: carnitine O-palmitoyltransferase 1, liver isoform
>gi|1036940220|ref|XP_017145836.1| PREDICTED: carnitine O-palmitoyltransferase 1, liver isoform
>gi|1036940289|ref|XP_017145841.1| PREDICTED: peptidyl-prolyl cis-trans isomerase-like 3 [Dros
>gi|1036949033|ref|XP_017146369.1| PREDICTED: chromatin assembly factor 1 subunit B [Drosophila
>gi|1036949134|ref|XP_017146376.1| PREDICTED: G protein alpha o subunit isoform X1 [Drosophila
>gi|1036949152|ref|XP_017146377.1| PREDICTED: G protein alpha o subunit isoform X2 [Drosophila
>gi|1036949087|ref|XP_017146373.1| PREDICTED: probable cytochrome P450 49a1 isoform X1 [Drosoph
>gi|1036949103|ref|XP_017146374.1| PREDICTED: probable cytochrome P450 49a1 isoform X1 [Drosoph
>gi|1036949118|ref|XP_017146375.1| PREDICTED: probable cytochrome P450 49a1 isoform X2 [Drosoph
>gi|1036949214|ref|XP_017146381.1| PREDICTED: succinate dehydrogenase assembly factor 2-A, mit
>gi|1036949267|ref|XP_017146385.1| PREDICTED: putative protein TPRXL [Drosophila miranda]
>gi|1036949072|ref|XP_017146372.1| PREDICTED: ran-binding proteins 9/10 homolog isoform X2 [Dro
>gi|1036949051|ref|XP_017146370.1| PREDICTED: ran-binding proteins 9/10 homolog isoform X1 [Dro
>gi|1036949231|ref|XP_017146382.1| PREDICTED: uncharacterized protein LOC108158516 isoform X1
>gi|1036949249|ref|XP_017146383.1| PREDICTED: uncharacterized protein LOC108158516 isoform X2
>gi|1036949165|ref|XP_017146378.1| PREDICTED: receptor expression-enhancing protein 5 isoform
>gi|1036949181|ref|XP_017146379.1| PREDICTED: receptor expression-enhancing protein 5 isoform
>gi|1036949199|ref|XP_017146380.1| PREDICTED: receptor expression-enhancing protein 5 isoform
>gi|1036949285|ref|XP_017146386.1| PREDICTED: probable 28S ribosomal protein S16, mitochondrial
>gi|1036948729|ref|XP_017146350.1| PREDICTED: zinc finger protein 189 isoform X1 [Drosophila mi
>gi|1036948763|ref|XP_017146352.1| PREDICTED: zinc finger protein 189 isoform X3 [Drosophila mi
>gi|1036948797|ref|XP_017146354.1| PREDICTED: zinc finger protein 189 isoform X5 [Drosophila mi
>gi|1036948747|ref|XP_017146351.1| PREDICTED: zinc finger protein 135 isoform X2 [Drosophila mi
>gi|1036948814|ref|XP_017146355.1| PREDICTED: zinc finger protein 189 isoform X6 [Drosophila mi
>gi|1036948779|ref|XP_017146353.1| PREDICTED: zinc finger protein 189 isoform X4 [Drosophila mi
>gi|1036948829|ref|XP_017146357.1| PREDICTED: zinc finger protein 300 isoform X7 [Drosophila mi
>gi|1036948849|ref|XP_017146358.1| PREDICTED: zinc finger protein 189 isoform X8 [Drosophila mi
>gi|1036948966|ref|XP_017146365.1| PREDICTED: zinc finger protein 665 isoform X15 [Drosophila m
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>gi|1036948983|ref|XP_017146366.1| PREDICTED: zinc finger protein 189 isoform X16 [Drosophila r
>gi|1036949017|ref|XP_017146368.1| PREDICTED: zinc finger protein 567 isoform X18 [Drosophila r
>gi|1036948867|ref|XP_017146359.1| PREDICTED: zinc finger protein 189 isoform X9 [Drosophila mi
>gi|1036948883|ref|XP_017146360.1| PREDICTED: zinc finger protein 189 isoform X10 [Drosophila r
>gi|1036948899|ref|XP_017146361.1| PREDICTED: zinc finger protein 189 isoform X11 [Drosophila r
>gi|1036948917|ref|XP_017146362.1| PREDICTED: zinc finger protein 135 isoform X12 [Drosophila r
>gi|1036948933|ref|XP_017146363.1| PREDICTED: zinc finger protein 189 isoform X13 [Drosophila r
>gi|1036948949|ref|XP_017146364.1| PREDICTED: zinc finger protein 189 isoform X14 [Drosophila r
>gi|1036949001|ref|XP_017146367.1| PREDICTED: zinc finger protein 189 isoform X17 [Drosophila r
>gi|1037004800|ref|XP_017149730.1| PREDICTED: uncharacterized protein LOC108160317 [Drosophila
>gi|1037004852|ref|XP_017149734.1| PREDICTED: vesicular inhibitory amino acid transporter [Dros
>gi|1037004818|ref|XP_017149731.1| PREDICTED: putative transcription factor SOX-15 isoform X1
>gi|1037004834|ref|XP_017149732.1| PREDICTED: putative transcription factor SOX-15 isoform X2
>gi|1036975311|ref|XP_017147933.1| PREDICTED: LOW QUALITY PROTEIN: nucleoprotein TPR [Drosophil
>gi|1036975329|ref|XP_017147934.1| PREDICTED: 40S ribosomal protein S23 [Drosophila miranda]
>gi|1036998556|ref|XP_017149366.1| PREDICTED: putative ferric-chelate reductase 1 homolog isofo
>gi|1036998592|ref|XP_017149368.1| PREDICTED: putative ferric-chelate reductase 1 homolog isofo
>gi|1036998574|ref|XP_017149367.1| PREDICTED: putative ferric-chelate reductase 1 homolog isofo
>gi|1036998540|ref|XP_017149365.1| PREDICTED: FAS-associated factor 1 isoform X2 [Drosophila mi
>gi|1036998525|ref|XP_017149364.1| PREDICTED: FAS-associated factor 1 isoform X1 [Drosophila mi
>gi|1036943110|ref|XP_017146011.1| PREDICTED: uncharacterized protein LOC108158285 [Drosophila
>gi|1037015660|ref|XP_017150391.1| PREDICTED: pikachurin isoform X1 [Drosophila miranda]
>gi|1037015676|ref|XP_017150392.1| PREDICTED: pikachurin isoform X1 [Drosophila miranda]
>gi|1037015690|ref|XP_017150393.1| PREDICTED: pikachurin isoform X2 [Drosophila miranda]
>gi|1036998403|ref|XP_017149356.1| PREDICTED: uncharacterized protein LOC108160090 [Drosophila
>gi|1036967528|ref|XP_017147460.1| PREDICTED: dystroglycan isoform X4 [Drosophila miranda]
>gi|1036967511|ref|XP_017147459.1| PREDICTED: dystroglycan isoform X3 [Drosophila miranda]
>gi|1036967564|ref|XP_017147462.1| PREDICTED: uncharacterized protein LOC108159044 isoform X6
>gi|1036967546|ref|XP_017147461.1| PREDICTED: uncharacterized protein LOC108159044 isoform X5
>gi|1036967491|ref|XP_017147458.1| PREDICTED: dystroglycan isoform X2 [Drosophila miranda]
>gi|1036967473|ref|XP_017147457.1| PREDICTED: dystroglycan isoform X1 [Drosophila miranda]
>gi|1036967619|ref|XP_017147465.1| PREDICTED: 39S ribosomal protein L34, mitochondrial [Drosoph
>gi|1036967584|ref|XP_017147463.1| PREDICTED: polynucleotide 5'-hydroxyl-kinase NOL9 [Drosophil
>gi|1036967601|ref|XP_017147464.1| PREDICTED: polynucleotide 5'-hydroxyl-kinase NOL9 [Drosophil
>gi|1037006145|ref|XP_017149818.1| PREDICTED: uncharacterized protein LOC108160359 [Drosophila
>gi|1037006109|ref|XP_017149816.1| PREDICTED: serine protease snake isoform X1 [Drosophila mira
>gi|1037006127|ref|XP_017149817.1| PREDICTED: serine protease snake isoform X2 [Drosophila mira
>gi|1037001379|ref|XP_017149530.1| PREDICTED: methyltransferase-like protein 6 [Drosophila mira
>gi|1037001365|ref|XP_017149529.1| PREDICTED: hyccin isoform X2 [Drosophila miranda]
>gi|1037001350|ref|XP_017149528.1| PREDICTED: hyccin isoform X1 [Drosophila miranda]
>gi|1036985282|ref|XP_017148553.1| PREDICTED: stomatin-4 [Drosophila miranda]
>gi|1036985266|ref|XP_017148552.1| PREDICTED: phosphatidylinositol N-acetylglucosaminyltransfer
>gi|1036985219|ref|XP_017148549.1| PREDICTED: syndetin [Drosophila miranda]
>gi|1036985248|ref|XP_017148551.1| PREDICTED: uncharacterized protein LOC108159608 [Drosophila
>gi|1036985232|ref|XP_017148550.1| PREDICTED: larval serum protein 2 [Drosophila miranda]
>gi|1036937458|ref|XP_017145659.1| PREDICTED: WD repeat-containing protein 5 [Drosophila mirand
>gi|1036984586|ref|XP_017148510.1| PREDICTED: sodium-independent sulfate anion transporter isof
>gi|1036984572|ref|XP_017148509.1| PREDICTED: sodium-independent sulfate anion transporter isof
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>gi|1036984602|ref|XP_017148511.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1036984620|ref|XP_017148512.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1036984499|ref|XP_017148505.1| PREDICTED: sarcosine dehydrogenase, mitochondrial [Drosophil
>gi|1036984636|ref|XP_017148514.1| PREDICTED: neurogenic differentiation factor 2 [Drosophila r
>gi|1036984517|ref|XP_017148506.1| PREDICTED: probable peroxisomal acyl-coenzyme A oxidase 1 [
>gi|1036984554|ref|XP_017148508.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1037021003|ref|XP_017150731.1| PREDICTED: uncharacterized protein LOC108160951 [Drosophila
>gi|1037020987|ref|XP_017150730.1| PREDICTED: uncharacterized protein LOC108160950 [Drosophila
>gi|1037018092|ref|XP_017150540.1| PREDICTED: uncharacterized protein LOC108160821 isoform X2
>gi|1037018078|ref|XP_017150539.1| PREDICTED: uncharacterized protein LOC108160821 isoform X1
>gi|1036990591|ref|XP_017148877.1| PREDICTED: putative 1-phosphatidylinositol 3-phosphate 5-kir
>gi|1036990609|ref|XP_017148878.1| PREDICTED: anaphase-promoting complex subunit 13 [Drosophila
>gi|1036981521|ref|XP_017148324.1| PREDICTED: cap-specific mRNA (nucleoside-2'-O-)-methyltransi
>gi|1036981556|ref|XP_017148327.1| PREDICTED: ubiquitin-conjugating enzyme E2 L3 [Drosophila m:
>gi|1036981507|ref|XP_017148323.1| PREDICTED: ribosome biogenesis protein BOP1 homolog [Drosopi
>gi|1036981538|ref|XP_017148326.1| PREDICTED: regulator of G-protein signaling 17 [Drosophila r
>gi|1036983456|ref|XP_017148440.1| PREDICTED: protein grainyhead isoform X1 [Drosophila miranda
>gi|1036983472|ref|XP_017148441.1| PREDICTED: protein grainyhead isoform X2 [Drosophila miranda
>gi|1036983529|ref|XP_017148444.1| PREDICTED: protein grainyhead isoform X5 [Drosophila miranda
>gi|1036983565|ref|XP_017148446.1| PREDICTED: protein grainyhead isoform X7 [Drosophila miranda
>gi|1036983491|ref|XP_017148442.1| PREDICTED: protein grainyhead isoform X3 [Drosophila miranda
>gi|1036983511|ref|XP_017148443.1| PREDICTED: protein grainyhead isoform X4 [Drosophila miranda
>gi|1036983547|ref|XP_017148445.1| PREDICTED: protein grainyhead isoform X6 [Drosophila miranda
>gi|1036983579|ref|XP_017148447.1| PREDICTED: protein grainyhead isoform X8 [Drosophila miranda
>gi|1036941699|ref|XP_017145919.1| PREDICTED: uncharacterized protein LOC108158208 [Drosophila
>gi|1036968129|ref|XP_017147498.1| PREDICTED: uncharacterized protein LOC108159064 [Drosophila
>gi|1036968073|ref|XP_017147494.1| PREDICTED: calcium release-activated calcium channel protei
>gi|1036968095|ref|XP_017147495.1| PREDICTED: calcium release-activated calcium channel protei
>gi|1036968111|ref|XP_017147496.1| PREDICTED: calcium release-activated calcium channel protei
>gi|1036968147|ref|XP_017147499.1| PREDICTED: trypsin I-P1 isoform X1 [Drosophila miranda]
>gi|1036968215|ref|XP_017147500.1| PREDICTED: trypsin I-P1 isoform X2 [Drosophila miranda]
>gi|1036968053|ref|XP_017147493.1| PREDICTED: calcium release-activated calcium channel protei
>gi|1036968233|ref|XP_017147501.1| PREDICTED: MORN repeat-containing protein 3 [Drosophila mira
>gi|1036968035|ref|XP_017147492.1| PREDICTED: protein three rows isoform X2 [Drosophila miranda
>gi|1036968015|ref|XP_017147491.1| PREDICTED: protein three rows isoform X1 [Drosophila miranda
>gi|1037000161|ref|XP_017149458.1| PREDICTED: acidic leucine-rich nuclear phosphoprotein 32 far
>gi|1037000141|ref|XP_017149457.1| PREDICTED: acidic leucine-rich nuclear phosphoprotein 32 far
>gi|1037006058|ref|XP_017149812.1| PREDICTED: enoyl-CoA hydratase domain-containing protein 3,
>gi|1037006074|ref|XP_017149814.1| PREDICTED: glutathione S-transferase S1 [Drosophila miranda]
>gi|1037006091|ref|XP_017149815.1| PREDICTED: glutathione S-transferase S1 [Drosophila miranda]
>gi|1037014644|ref|XP_017150331.1| PREDICTED: triple functional domain protein [Drosophila mira
>gi|1037014660|ref|XP_017150332.1| PREDICTED: uncharacterized protein LOC108160691 [Drosophila
>gi|1037019362|ref|XP_017150625.1| PREDICTED: kalirin isoform X1 [Drosophila miranda]
>gi|1037019378|ref|XP_017150627.1| PREDICTED: kalirin isoform X2 [Drosophila miranda]
>gi|1036943845|ref|XP_017146056.1| PREDICTED: uncharacterized protein LOC108158326 [Drosophila
>gi|1036943094|ref|XP_017146010.1| PREDICTED: eIF-2-alpha kinase GCN2-like [Drosophila miranda]
>gi|1036942403|ref|XP_017145966.1| PREDICTED: uncharacterized protein LOC108158244 [Drosophila
>gi|1036942922|ref|XP_017145999.1| PREDICTED: uncharacterized protein LOC108158274 [Drosophila
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>gi|1036942345|ref|XP_017145963.1| PREDICTED: uncharacterized protein LOC108158241 [Drosophila
>gi|1036942509|ref|XP_017145973.1| PREDICTED: uncharacterized protein LOC108158248 [Drosophila
>gi|1037016233|ref|XP_017150423.1| PREDICTED: glutenin, high molecular weight subunit DX5 isof
>gi|1037016249|ref|XP_017150424.1| PREDICTED: glutenin, high molecular weight subunit DX5 isof
>gi|1036940789|ref|XP_017145873.1| PREDICTED: uncharacterized protein LOC108158161 [Drosophila
>gi|1036940806|ref|XP_017145874.1| PREDICTED: uncharacterized protein LOC108158162 [Drosophila
>gi|1036946344|ref|XP_017146202.1| PREDICTED: protein TonB [Drosophila miranda]
>gi|1036937476|ref|XP_017145660.1| PREDICTED: probable palmitoyltransferase ZDHHC20 [Drosophila
>gi|1036946560|ref|XP_017146216.1| PREDICTED: leucine-rich repeat extensin-like protein 3 [Dros
>gi|1036946362|ref|XP_017146203.1| PREDICTED: skin secretory protein xP2 [Drosophila miranda]
>gi|1036946444|ref|XP_017146209.1| PREDICTED: skin secretory protein xP2 [Drosophila miranda]
>gi|1036976876|ref|XP_017148030.1| PREDICTED: uncharacterized protein LOC108159342 [Drosophila
>gi|1036976822|ref|XP_017148027.1| PREDICTED: uncharacterized protein LOC108159340 [Drosophila
>gi|1036976804|ref|XP_017148026.1| PREDICTED: antifreeze protein Maxi-like [Drosophila miranda]
>gi|1036976840|ref|XP_017148028.1| PREDICTED: uncharacterized protein LOC108159341 isoform X1
>gi|1036976858|ref|XP_017148029.1| PREDICTED: uncharacterized protein LOC108159341 isoform X2
>gi|1036937494|ref|XP_017145662.1| PREDICTED: polyglutamine-repeat protein pqn-41 [Drosophila r
>gi|1037002200|ref|XP_017149584.1| PREDICTED: Na(+)/H(+) exchange regulatory cofactor NHE-RF1
>gi|1037002182|ref|XP_017149583.1| PREDICTED: uncharacterized protein LOC108160237 [Drosophila
>gi|1036991607|ref|XP_017148932.1| PREDICTED: uncharacterized protein LOC108159851 [Drosophila
>gi|1036991591|ref|XP_017148931.1| PREDICTED: LOW QUALITY PROTEIN: bcl-2-related ovarian killer
>gi|1036991573|ref|XP_017148930.1| PREDICTED: electron transfer flavoprotein subunit alpha, mit
>gi|1036991553|ref|XP_017148928.1| PREDICTED: electron transfer flavoprotein subunit alpha, mit
>gi|1036991535|ref|XP_017148927.1| PREDICTED: RNA/RNP complex-1-interacting phosphatase [Droso
>gi|1036937512|ref|XP_017145663.1| PREDICTED: uncharacterized protein LOC108158030 [Drosophila
>gi|1036990847|ref|XP_017148891.1| PREDICTED: uncharacterized protein LOC108159827 isoform X1
>gi|1036990865|ref|XP_017148892.1| PREDICTED: uncharacterized protein LOC108159827 isoform X2
>gi|1036990885|ref|XP_017148893.1| PREDICTED: uncharacterized protein LOC108159827 isoform X2
>gi|1036990901|ref|XP_017148894.1| PREDICTED: uncharacterized protein LOC108159827 isoform X2
>gi|1036936055|ref|XP_017145569.1| PREDICTED: integrator complex subunit 5-like protein [Droso
>gi|1036990805|ref|XP_017148888.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036990919|ref|XP_017148895.1| PREDICTED: uncharacterized protein LOC108159828 isoform X1
>gi|1036990935|ref|XP_017148896.1| PREDICTED: uncharacterized protein LOC108159828 isoform X2
>gi|1036990835|ref|XP_017148890.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036990971|ref|XP_017148898.1| PREDICTED: uncharacterized protein LOC108159830 isoform X2
>gi|1036990953|ref|XP_017148897.1| PREDICTED: uncharacterized protein LOC108159830 isoform X1
>gi|1036936073|ref|XP_017145570.1| PREDICTED: uncharacterized protein LOC108157937 [Drosophila
>gi|1036990821|ref|XP_017148889.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036994450|ref|XP_017149109.1| PREDICTED: guanine nucleotide-binding protein subunit beta-1
>gi|1036994344|ref|XP_017149102.1| PREDICTED: sine oculis-binding protein homolog isoform X1 [D
>gi|1036994366|ref|XP_017149103.1| PREDICTED: sine oculis-binding protein homolog isoform X1 [D
>gi|1036994384|ref|XP_017149104.1| PREDICTED: sine oculis-binding protein homolog isoform X1 [D
>gi|1036994401|ref|XP_017149105.1| PREDICTED: sine oculis-binding protein homolog isoform X2 [D
>gi|1036994418|ref|XP_017149106.1| PREDICTED: sine oculis-binding protein homolog isoform X3 [D
>gi|1036994468|ref|XP_017149110.1| PREDICTED: LYR motif-containing protein 2 [Drosophila mirand
>gi|1036994435|ref|XP_017149107.1| PREDICTED: membrane-bound transcription factor site-2 protea
>gi|1036942740|ref|XP_017145988.1| PREDICTED: uncharacterized protein LOC108158263 [Drosophila
>gi|1037018257|ref|XP_017150552.1| PREDICTED: uncharacterized protein LOC108160830 isoform X1
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>gi|1037018275|ref|XP_017150553.1| PREDICTED: uncharacterized protein LOC108160830 isoform X2
>gi|1036946378|ref|XP_017146204.1| PREDICTED: uncharacterized protein LOC108158433 [Drosophila
>gi|1036941629|ref|XP_017145915.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036954865|ref|XP_017146729.1| PREDICTED: uncharacterized protein LOC108158702 [Drosophila
>gi|1036954768|ref|XP_017146725.1| PREDICTED: uncharacterized protein LOC108158701 isoform X1
>gi|1036954788|ref|XP_017146726.1| PREDICTED: uncharacterized protein LOC108158701 isoform X2
>gi|1036954837|ref|XP_017146727.1| PREDICTED: uncharacterized protein LOC108158701 isoform X3
>gi|1036954742|ref|XP_017146724.1| PREDICTED: caspase-3 [Drosophila miranda]
>gi|1036954662|ref|XP_017146720.1| PREDICTED: LOW QUALITY PROTEIN: midasin [Drosophila miranda]
>gi|1036954702|ref|XP_017146722.1| PREDICTED: protein cutoff isoform X1 [Drosophila miranda]
>gi|1036954724|ref|XP_017146723.1| PREDICTED: protein cutoff isoform X2 [Drosophila miranda]
>gi|1036954684|ref|XP_017146721.1| PREDICTED: protein disulfide-isomerase A3 [Drosophila miran
>gi|1036966016|ref|XP_017147376.1| PREDICTED: elongation factor 1-alpha 1 [Drosophila miranda]
>gi|1036965982|ref|XP_017147374.1| PREDICTED: uncharacterized protein LOC108159012 isoform X1
>gi|1036965999|ref|XP_017147375.1| PREDICTED: uncharacterized protein LOC108159012 isoform X2
>gi|1036966052|ref|XP_017147379.1| PREDICTED: zinc transporter ZIP11 [Drosophila miranda]
>gi|1036966034|ref|XP_017147378.1| PREDICTED: uncharacterized protein LOC108159014 [Drosophila
>gi|1037001906|ref|XP_017149566.1| PREDICTED: uncharacterized protein LOC108160226 isoform X1
>gi|1037001923|ref|XP_017149567.1| PREDICTED: lateral signaling target protein 2 homolog isofo
>gi|1037001941|ref|XP_017149568.1| PREDICTED: proteasome inhibitor PI31 subunit [Drosophila mi
>gi|1036958943|ref|XP_017146962.1| PREDICTED: uncharacterized protein LOC108158791 isoform X1
>gi|1036958956|ref|XP_017146963.1| PREDICTED: uncharacterized protein LOC108158791 isoform X2
>gi|1036958969|ref|XP_017146964.1| PREDICTED: uncharacterized protein LOC108158791 isoform X2
>gi|1036958985|ref|XP_017146965.1| PREDICTED: uncharacterized protein LOC108158791 isoform X2
>gi|1036959000|ref|XP_017146966.1| PREDICTED: uncharacterized protein LOC108158791 isoform X2
>gi|1036959032|ref|XP_017146968.1| PREDICTED: origin recognition complex subunit 6 [Drosophila
>gi|1036959017|ref|XP_017146967.1| PREDICTED: proteasome subunit alpha type-3 [Drosophila mira
>gi|1036959078|ref|XP_017146971.1| PREDICTED: transducin beta-like protein 3 [Drosophila miran
>gi|1036959064|ref|XP_017146970.1| PREDICTED: pyruvate carboxylase, mitochondrial isoform X2 [
>gi|1036959047|ref|XP_017146969.1| PREDICTED: pyruvate carboxylase, mitochondrial isoform X1 [
>gi|1036959094|ref|XP_017146973.1| PREDICTED: probable aspartate aminotransferase, cytoplasmic
>gi|1036959144|ref|XP_017146976.1| PREDICTED: DNA-binding protein SMUBP-2 [Drosophila miranda]
>gi|1036959129|ref|XP_017146975.1| PREDICTED: UPF0598 protein CG30010 [Drosophila miranda]
>gi|1036959111|ref|XP_017146974.1| PREDICTED: protein crossbronx [Drosophila miranda]
>gi|1037016328|ref|XP_017150429.1| PREDICTED: leucine-rich repeat neuronal protein 2 [Drosophil
>gi|1037016365|ref|XP_017150431.1| PREDICTED: venom allergen 5.01 [Drosophila miranda]
>gi|1036937530|ref|XP_017145664.1| PREDICTED: uncharacterized protein LOC108158031, partial [Dr
>gi|1037016345|ref|XP_017150430.1| PREDICTED: venom allergen 5-like [Drosophila miranda]
>gi|1036969387|ref|XP_017147569.1| PREDICTED: cytochrome P450 9c1 [Drosophila miranda]
>gi|1036969534|ref|XP_017147577.1| PREDICTED: uncharacterized protein LOC108159110 [Drosophila
>gi|1036969551|ref|XP_017147579.1| PREDICTED: uncharacterized protein LOC108159110 [Drosophila
>gi|1036969569|ref|XP_017147580.1| PREDICTED: uncharacterized protein LOC108159110 [Drosophila
>gi|1036969691|ref|XP_017147587.1| PREDICTED: uncharacterized protein LOC108159115 isoform X2
>gi|1036969677|ref|XP_017147586.1| PREDICTED: uncharacterized protein LOC108159115 isoform X1
>gi|1036969585|ref|XP_017147581.1| PREDICTED: uncharacterized aarF domain-containing protein ki
>gi|1036969603|ref|XP_017147582.1| PREDICTED: uncharacterized aarF domain-containing protein ki
>gi|1036969477|ref|XP_017147574.1| PREDICTED: probable ribonuclease ZC3H12C isoform X4 [Drosoph
>gi|1036969443|ref|XP_017147572.1| PREDICTED: probable ribonuclease ZC3H12C isoform X2 [Drosoph
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>gi|1036969459|ref|XP_017147573.1| PREDICTED: probable ribonuclease ZC3H12C isoform X3 [Drosop]
>gi|1036969403|ref|XP_017147570.1| PREDICTED: probable ribonuclease ZC3H12C isoform X1 [Drosop]
>gi|1036969425|ref|XP_017147571.1| PREDICTED: probable ribonuclease ZC3H12C isoform X1 [Drosop]
>gi|1036969496|ref|XP_017147575.1| PREDICTED: probable ribonuclease ZC3H12C isoform X5 [Drosop]
>gi|1036969514|ref|XP_017147576.1| PREDICTED: uncharacterized protein LOC108159108 isoform X6
>gi|1036969370|ref|XP_017147568.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036969640|ref|XP_017147584.1| PREDICTED: probable methyltransferase BTM2 homolog [Drosoph]
>gi|1036969658|ref|XP_017147585.1| PREDICTED: uncharacterized protein LOC108159114 [Drosophila
>gi|1036969623|ref|XP_017147583.1| PREDICTED: uncharacterized protein LOC108159112 [Drosophila
>gi|1036982812|ref|XP_017148402.1| PREDICTED: putative ATP synthase subunit f, mitochondrial []
>gi|1036982794|ref|XP_017148401.1| PREDICTED: uncharacterized protein LOC108159536 [Drosophila
>gi|1036982760|ref|XP_017148399.1| PREDICTED: uncharacterized protein LOC108159534 isoform X2
>gi|1036982742|ref|XP_017148398.1| PREDICTED: uncharacterized protein LOC108159534 isoform X1
>gi|1036982776|ref|XP_017148400.1| PREDICTED: testis-expressed sequence 9 protein [Drosophila n
>gi|1036982830|ref|XP_017148403.1| PREDICTED: neuropeptide SIFamide [Drosophila miranda]
>gi|1036982705|ref|XP_017148396.1| PREDICTED: stAR-related lipid transfer protein 3 [Drosophila
>gi|1036982723|ref|XP_017148397.1| PREDICTED: stAR-related lipid transfer protein 3 [Drosophila
>gi|1036968656|ref|XP_017147526.1| PREDICTED: peptidylprolyl isomerase domain and WD repeat-con
>gi|1036968639|ref|XP_017147525.1| PREDICTED: peptidylprolyl isomerase domain and WD repeat-con
>gi|1036968676|ref|XP_017147527.1| PREDICTED: peptidylprolyl isomerase domain and WD repeat-con
>gi|1036968592|ref|XP_017147522.1| PREDICTED: RNA polymerase II subunit A C-terminal domain pho
>gi|1036968624|ref|XP_017147524.1| PREDICTED: myosin-3 [Drosophila miranda]
>gi|1036968610|ref|XP_017147523.1| PREDICTED: zinc finger protein 598 [Drosophila miranda]
>gi|1036997912|ref|XP_017149324.1| PREDICTED: inorganic pyrophosphatase [Drosophila miranda]
>gi|1036997928|ref|XP_017149325.1| PREDICTED: ion transport peptide isoform X1 [Drosophila mira
>gi|1036997984|ref|XP_017149330.1| PREDICTED: ion transport peptide-like isoform X4 [Drosophila
>gi|1036997950|ref|XP_017149326.1| PREDICTED: ion transport peptide-like isoform X2 [Drosophila
>gi|1036997968|ref|XP_017149327.1| PREDICTED: ion transport peptide-like isoform X3 [Drosophila
>gi|1036998001|ref|XP_017149331.1| PREDICTED: ion transport peptide-like isoform X5 [Drosophila
>gi|1036998018|ref|XP_017149332.1| PREDICTED: ion transport peptide-like isoform X5 [Drosophila
>gi|1036998036|ref|XP_017149333.1| PREDICTED: ion transport peptide isoform X6 [Drosophila mira
>gi|1036937544|ref|XP_017145665.1| PREDICTED: chorion protein S38 [Drosophila miranda]
>gi|1037006220|ref|XP_017149823.1| PREDICTED: zinc finger protein 227 [Drosophila miranda]
>gi|1037006238|ref|XP_017149824.1| PREDICTED: geminin [Drosophila miranda]
>gi|1036993368|ref|XP_017149043.1| PREDICTED: uncharacterized protein LOC108159918 [Drosophila
>gi|1036993384|ref|XP_017149044.1| PREDICTED: tetratricopeptide repeat protein 4 [Drosophila mi
>gi|1036984912|ref|XP_017148531.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036984930|ref|XP_017148532.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036984896|ref|XP_017148530.1| PREDICTED: diacylglycerol kinase epsilon [Drosophila miranda
>gi|1036984860|ref|XP_017148528.1| PREDICTED: 25S rRNA (cytosine-C(5))-methyltransferase nop2
>gi|1036984878|ref|XP_017148529.1| PREDICTED: endothelin-converting enzyme 1 [Drosophila mirand
>gi|1036973631|ref|XP_017147831.1| PREDICTED: diuretic hormone receptor isoform X1 [Drosophila
>gi|1036973645|ref|XP_017147833.1| PREDICTED: diuretic hormone receptor isoform X1 [Drosophila
>gi|1036973658|ref|XP_017147834.1| PREDICTED: diuretic hormone receptor isoform X2 [Drosophila
>gi|1036973670|ref|XP_017147835.1| PREDICTED: diuretic hormone receptor isoform X3 [Drosophila
>gi|1036973686|ref|XP_017147836.1| PREDICTED: diuretic hormone receptor isoform X4 [Drosophila
>gi|1036973706|ref|XP_017147837.1| PREDICTED: diuretic hormone receptor isoform X5 [Drosophila
>gi|1036973533|ref|XP_017147824.1| PREDICTED: probable beta-hexosaminidase fdl isoform X2 [Dros
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>gi|1036973517|ref|XP_017147823.1| PREDICTED: probable beta-hexosaminidase fdl isoform X1 [Dros
>gi|1036973549|ref|XP_017147825.1| PREDICTED: probable beta-hexosaminidase fdl isoform X1 [Dros
>gi|1036973753|ref|XP_017147840.1| PREDICTED: uncharacterized protein DDB_G0271670-like [Drosop
>gi|1036973719|ref|XP_017147838.1| PREDICTED: uncharacterized protein LOC108159236 isoform X1
>gi|1036973736|ref|XP_017147839.1| PREDICTED: uncharacterized protein LOC108159236 isoform X2
>gi|1036973602|ref|XP_017147829.1| PREDICTED: homeobox protein PKNOX2 isoform X2 [Drosophila mi
>gi|1036973574|ref|XP_017147827.1| PREDICTED: uncharacterized protein LOC108159232 isoform X1
>gi|1036973588|ref|XP_017147828.1| PREDICTED: uncharacterized protein LOC108159232 isoform X1
>gi|1036973561|ref|XP_017147826.1| PREDICTED: uncharacterized protein LOC108159231 [Drosophila
>gi|1036973616|ref|XP_017147830.1| PREDICTED: uncharacterized protein LOC108159234 [Drosophila
>gi|1036951017|ref|XP_017146497.1| PREDICTED: transmembrane protein 131 homolog [Drosophila mi
>gi|1036951303|ref|XP_017146514.1| PREDICTED: ubiquitin-fold modifier-conjugating enzyme 1 [Dro
>gi|1036951177|ref|XP_017146506.1| PREDICTED: transcription factor grauzone isoform X2 [Drosop
>gi|1036951159|ref|XP_017146505.1| PREDICTED: transcription factor grauzone isoform X1 [Drosop
>gi|1036951192|ref|XP_017146507.1| PREDICTED: monocarboxylate transporter 13 [Drosophila miran
>gi|1036951210|ref|XP_017146509.1| PREDICTED: monocarboxylate transporter 13 [Drosophila miran
>gi|1036951275|ref|XP_017146513.1| PREDICTED: proteasome subunit beta type-6 [Drosophila miran
>gi|1036951226|ref|XP_017146510.1| PREDICTED: exosome complex exonuclease RRP42 [Drosophila mi
>gi|1036951245|ref|XP_017146511.1| PREDICTED: ER membrane protein complex subunit 7 homolog [Dr
>gi|1036951262|ref|XP_017146512.1| PREDICTED: uncharacterized protein LOC108158580 [Drosophila
>gi|1036951107|ref|XP_017146502.1| PREDICTED: monocarboxylate transporter 3 [Drosophila miranda
>gi|1036951125|ref|XP_017146503.1| PREDICTED: monocarboxylate transporter 3 [Drosophila miranda
>gi|1036951143|ref|XP_017146504.1| PREDICTED: monocarboxylate transporter 3 [Drosophila miranda
>gi|1036951089|ref|XP_017146501.1| PREDICTED: dynamin-like 120 kDa protein, mitochondrial isofo
>gi|1036951071|ref|XP_017146500.1| PREDICTED: dynamin-like 120 kDa protein, mitochondrial isofo
>gi|1036951053|ref|XP_017146499.1| PREDICTED: dynamin-like 120 kDa protein, mitochondrial isofo
>gi|1036951035|ref|XP_017146498.1| PREDICTED: dynamin-like 120 kDa protein, mitochondrial isofo
>gi|1036951492|ref|XP_017146528.1| PREDICTED: SNF-related serine/threonine-protein kinase [Dros
>gi|1036951480|ref|XP_017146527.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 20 [Drosop
>gi|1036951560|ref|XP_017146532.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036951466|ref|XP_017146526.1| PREDICTED: multidrug resistance protein 1A [Drosophila mira
>gi|1036951510|ref|XP_017146529.1| PREDICTED: heat shock 70 kDa protein cognate 5 [Drosophila
>gi|1036951582|ref|XP_017146533.1| PREDICTED: dnaJ homolog subfamily C member 11 [Drosophila m
>gi|1036951599|ref|XP_017146534.1| PREDICTED: dnaJ homolog subfamily C member 11 [Drosophila m
>gi|1036951619|ref|XP_017146535.1| PREDICTED: beta-1,4-N-acetylgalactosaminyltransferase bre-4
>gi|1036951544|ref|XP_017146531.1| PREDICTED: putative GPI-anchored protein PB15E9.01c isoform
>gi|1036951526|ref|XP_017146530.1| PREDICTED: putative GPI-anchored protein PB15E9.01c isoform
>gi|1036951637|ref|XP_017146536.1| PREDICTED: selenide, water dikinase [Drosophila miranda]
>gi|1036972875|ref|XP_017147783.1| PREDICTED: chaoptin isoform X1 [Drosophila miranda]
>gi|1036972889|ref|XP_017147784.1| PREDICTED: chaoptin isoform X2 [Drosophila miranda]
>gi|1036972907|ref|XP_017147785.1| PREDICTED: uncharacterized protein LOC108159216 isoform X1
>gi|1036972925|ref|XP_017147787.1| PREDICTED: uncharacterized protein LOC108159216 isoform X2
>gi|1036973014|ref|XP_017147792.1| PREDICTED: uncharacterized protein LOC108159216 isoform X7
>gi|1036973032|ref|XP_017147793.1| PREDICTED: uncharacterized protein LOC108159216 isoform X8
>gi|1036972963|ref|XP_017147789.1| PREDICTED: uncharacterized protein LOC108159216 isoform X4
>gi|1036973050|ref|XP_017147794.1| PREDICTED: uncharacterized protein LOC108159216 isoform X9
>gi|1036973084|ref|XP_017147796.1| PREDICTED: uncharacterized protein LOC108159216 isoform X11
>gi|1036972945|ref|XP_017147788.1| PREDICTED: uncharacterized protein LOC108159216 isoform X3
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>gi|1036972980|ref|XP_017147790.1| PREDICTED: uncharacterized protein LOC108159216 isoform X5
>gi|1036973066|ref|XP_017147795.1| PREDICTED: uncharacterized protein LOC108159216 isoform X10
>gi|1036972998|ref|XP_017147791.1| PREDICTED: uncharacterized protein LOC108159216 isoform X6
>gi|1036973101|ref|XP_017147798.1| PREDICTED: uncharacterized protein LOC108159216 isoform X12
>gi|1036973119|ref|XP_017147799.1| PREDICTED: uncharacterized protein LOC108159216 isoform X13
>gi|1036973136|ref|XP_017147800.1| PREDICTED: potassium/sodium hyperpolarization-activated cycl
>gi|1036973182|ref|XP_017147803.1| PREDICTED: uncharacterized protein LOC108159217 isoform X2
>gi|1036973152|ref|XP_017147801.1| PREDICTED: uncharacterized protein LOC108159217 isoform X1
>gi|1036973168|ref|XP_017147802.1| PREDICTED: uncharacterized protein LOC108159217 isoform X1
>gi|1037013532|ref|XP_017150263.1| PREDICTED: 60S ribosomal protein L23 [Drosophila miranda]
>gi|1037013514|ref|XP_017150262.1| PREDICTED: inactivation-no-after-potential D protein [Drosop
>gi|1037021856|ref|XP_017150782.1| PREDICTED: fork head domain-containing protein FD3 [Drosophi
>gi|1037021872|ref|XP_017150783.1| PREDICTED: fork head domain-containing protein FD3 [Drosophi
>gi|1037022167|ref|XP_017150800.1| PREDICTED: uncharacterized protein LOC108161005 isoform X1
>gi|1037022185|ref|XP_017150802.1| PREDICTED: uncharacterized protein LOC108161005 isoform X2
>gi|1036944460|ref|XP_017146088.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944498|ref|XP_017146090.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944372|ref|XP_017146083.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944404|ref|XP_017146085.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944442|ref|XP_017146087.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944386|ref|XP_017146084.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944426|ref|XP_017146086.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944480|ref|XP_017146089.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944516|ref|XP_017146091.1| PREDICTED: phosphatidylinositol 4-phosphate 5-kinase type-1
>gi|1036944767|ref|XP_017146107.1| PREDICTED: ER membrane protein complex subunit 8/9 homolog
>gi|1036944585|ref|XP_017146096.1| PREDICTED: ATP-dependent zinc metalloprotease YME1 homolog
>gi|1036944567|ref|XP_017146095.1| PREDICTED: ATP-dependent zinc metalloprotease YME1 homolog
>gi|1036944550|ref|XP_017146094.1| PREDICTED: ATP-dependent zinc metalloprotease YME1 homolog
>gi|1036944534|ref|XP_017146093.1| PREDICTED: ATP-dependent zinc metalloprotease YME1 homolog
>gi|1036944749|ref|XP_017146106.1| PREDICTED: OCIA domain-containing protein 1 [Drosophila mira
>gi|1036944683|ref|XP_017146103.1| PREDICTED: probable GDP-L-fucose synthase [Drosophila mirand
>gi|1036944330|ref|XP_017146081.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036944665|ref|XP_017146102.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1036944651|ref|XP_017146100.1| PREDICTED: chymotrypsin-like protease CTRL-1 [Drosophila mi
>gi|1036944785|ref|XP_017146108.1| PREDICTED: uncharacterized protein LOC108158365 [Drosophila
>gi|1036944803|ref|XP_017146109.1| PREDICTED: uncharacterized protein LOC108158367 [Drosophila
>gi|1036944603|ref|XP_017146097.1| PREDICTED: dynein intermediate chain 2, ciliary [Drosophila
>gi|1036944636|ref|XP_017146099.1| PREDICTED: fukutin-related protein [Drosophila miranda]
>gi|1036944819|ref|XP_017146110.1| PREDICTED: atrial natriuretic peptide receptor 1 [Drosophila
>gi|1036944618|ref|XP_017146098.1| PREDICTED: uncharacterized protein LOC108158356 [Drosophila
>gi|1036944731|ref|XP_017146105.1| PREDICTED: proteasome subunit alpha type-4 [Drosophila miran
>gi|1036944700|ref|XP_017146104.1| PREDICTED: uncharacterized protein LOC108158361 [Drosophila
>gi|1036944222|ref|XP_017146078.1| PREDICTED: uncharacterized protein LOC108158351 isoform X1
>gi|1036944312|ref|XP_017146080.1| PREDICTED: uncharacterized protein LOC108158351 isoform X3
>gi|1036944294|ref|XP_017146079.1| PREDICTED: uncharacterized protein LOC108158351 isoform X2
>gi|1036976630|ref|XP_017148015.1| PREDICTED: CDK5RAP3-like protein [Drosophila miranda]
>gi|1036976614|ref|XP_017148014.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036976596|ref|XP_017148012.1| PREDICTED: U2 snRNP-associated SURP motif-containing protein
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>gi|1036976651|ref|XP_017148016.1| PREDICTED: protein RMD5 homolog A [Drosophila miranda]
>gi|1037014904|ref|XP_017150347.1| PREDICTED: uncharacterized protein LOC108160704 isoform X1
>gi|1037014921|ref|XP_017150349.1| PREDICTED: uncharacterized protein LOC108160704 isoform X1
>gi|1037014953|ref|XP_017150351.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037014939|ref|XP_017150350.1| PREDICTED: uncharacterized protein LOC108160704 isoform X2
>gi|1037014967|ref|XP_017150352.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037015015|ref|XP_017150355.1| PREDICTED: uncharacterized protein LOC108160704 isoform X7
>gi|1037015064|ref|XP_017150358.1| PREDICTED: uncharacterized protein LOC108160704 isoform X10
>gi|1037014983|ref|XP_017150353.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037015033|ref|XP_017150356.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037014999|ref|XP_017150354.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037015049|ref|XP_017150357.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037015082|ref|XP_017150360.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037015100|ref|XP_017150361.1| PREDICTED: resistance to inhibitors of cholinesterase prote
>gi|1037015158|ref|XP_017150363.1| PREDICTED: uncharacterized protein LOC108160704 isoform X14
>gi|1037015140|ref|XP_017150362.1| PREDICTED: uncharacterized protein LOC108160704 isoform X13
>gi|1037017080|ref|XP_017150475.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcompl
>gi|1036961036|ref|XP_017147077.1| PREDICTED: DE-cadherin [Drosophila miranda]
>gi|1036961129|ref|XP_017147083.1| PREDICTED: F-actin-capping protein subunit alpha [Drosophil
>gi|1036961138|ref|XP_017147085.1| PREDICTED: uncharacterized protein LOC108158863 [Drosophila
>gi|1036961151|ref|XP_017147086.1| PREDICTED: 39S ribosomal protein L54, mitochondrial [Drosop
>gi|1036961052|ref|XP_017147078.1| PREDICTED: chitinase-3-like protein 1 [Drosophila miranda]
>gi|1036961079|ref|XP_017147079.1| PREDICTED: chitinase-like protein 3 [Drosophila miranda]
>gi|1036961097|ref|XP_017147081.1| PREDICTED: chitinase-like protein 3 isoform X1 [Drosophila
>gi|1036961113|ref|XP_017147082.1| PREDICTED: chitinase-like protein 3 isoform X2 [Drosophila
>gi|1036961089|ref|XP_017147080.1| PREDICTED: acidic mammalian chitinase [Drosophila miranda]
>gi|1037021569|ref|XP_017150763.1| PREDICTED: acidic mammalian chitinase [Drosophila miranda]
>gi|1036976580|ref|XP_017148011.1| PREDICTED: C3 and PZP-like alpha-2-macroglobulin domain-con
>gi|1036976562|ref|XP_017148010.1| PREDICTED: trehalase isoform X2 [Drosophila miranda]
>gi|1036976528|ref|XP_017148009.1| PREDICTED: trehalase isoform X1 [Drosophila miranda]
>gi|1036976510|ref|XP_017148008.1| PREDICTED: rab GTPase-binding effector protein 1 [Drosophil
>gi|1036937562|ref|XP_017145666.1| PREDICTED: uncharacterized protein LOC108158033 [Drosophila
>gi|1036976457|ref|XP_017148005.1| PREDICTED: uncharacterized protein LOC108159324 [Drosophila
>gi|1036976474|ref|XP_017148006.1| PREDICTED: uncharacterized protein LOC108159324 [Drosophila
>gi|1036976492|ref|XP_017148007.1| PREDICTED: uncharacterized protein LOC108159324 [Drosophila
>gi|1036985025|ref|XP_017148538.1| PREDICTED: protein king tubby [Drosophila miranda]
>gi|1036985008|ref|XP_017148536.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [
>gi|1036984946|ref|XP_017148533.1| PREDICTED: transmembrane and TPR repeat-containing protein
>gi|1036984972|ref|XP_017148534.1| PREDICTED: transmembrane and TPR repeat-containing protein
>gi|1036984990|ref|XP_017148535.1| PREDICTED: transmembrane and TPR repeat-containing protein
>gi|1036985059|ref|XP_017148539.1| PREDICTED: protein mago nashi [Drosophila miranda]
>gi|1036989794|ref|XP_017148828.1| PREDICTED: uncharacterized protein LOC108159779 [Drosophila
>gi|1036989719|ref|XP_017148824.1| PREDICTED: membrane-associated guanylate kinase, WW and PDZ
>gi|1036989735|ref|XP_017148825.1| PREDICTED: membrane-associated guanylate kinase, WW and PDZ
>gi|1036989764|ref|XP_017148826.1| PREDICTED: EF-hand calcium-binding domain-containing protei
>gi|1036989776|ref|XP_017148827.1| PREDICTED: EF-hand calcium-binding domain-containing protei
>gi|1037010276|ref|XP_017150067.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036987602|ref|XP_017148697.1| PREDICTED: SWI/SNF-related matrix-associated actin-dependent
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>gi|1036987584|ref|XP_017148696.1| PREDICTED: macoilin-2 isoform X2 [Drosophila miranda]
>gi|1036987546|ref|XP_017148693.1| PREDICTED: macoilin isoform X1 [Drosophila miranda]
>gi|1036987562|ref|XP_017148694.1| PREDICTED: macoilin isoform X1 [Drosophila miranda]
>gi|1036990285|ref|XP_017148860.1| PREDICTED: splicing factor, arginine/serine-rich 15 isoform
>gi|1036990267|ref|XP_017148859.1| PREDICTED: splicing factor, arginine/serine-rich 15 isoform
>gi|1036990303|ref|XP_017148861.1| PREDICTED: splicing factor, arginine/serine-rich 15 isoform
>gi|1036990321|ref|XP_017148862.1| PREDICTED: uncharacterized protein LOC108159804 isoform X4
>gi|1036997898|ref|XP_017149323.1| PREDICTED: nuclear pore complex protein Nup133-like isoform
>gi|1036997882|ref|XP_017149322.1| PREDICTED: nuclear pore complex protein Nup133-like isoform
>gi|1037018410|ref|XP_017150562.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1036992163|ref|XP_017148966.1| PREDICTED: uncharacterized protein LOC108159873 [Drosophila
>gi|1036992199|ref|XP_017148968.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm1 [Drosoph
>gi|1036992093|ref|XP_017148961.1| PREDICTED: TFIIH basal transcription factor complex helicase
>gi|1036992145|ref|XP_017148965.1| PREDICTED: GTP cyclohydrolase 1 isoform X3 [Drosophila miran
>gi|1036992181|ref|XP_017148967.1| PREDICTED: uncharacterized protein LOC108159874 [Drosophila
>gi|1036992109|ref|XP_017148963.1| PREDICTED: GTP cyclohydrolase 1 isoform X1 [Drosophila miran
>gi|1036992127|ref|XP_017148964.1| PREDICTED: GTP cyclohydrolase 1 isoform X2 [Drosophila miran
>gi|1037008880|ref|XP_017149986.1| PREDICTED: uncharacterized protein LOC108160477 [Drosophila
>gi|1037003435|ref|XP_017149654.1| PREDICTED: locomotion-related protein Hikaru genki isoform
>gi|1037003417|ref|XP_017149653.1| PREDICTED: locomotion-related protein Hikaru genki isoform
>gi|1037003451|ref|XP_017149656.1| PREDICTED: locomotion-related protein Hikaru genki isoform
>gi|1037003504|ref|XP_017149659.1| PREDICTED: probable cytochrome P450 4p2 isoform X3 [Drosoph
>gi|1037003486|ref|XP_017149658.1| PREDICTED: probable cytochrome P450 4p2 isoform X2 [Drosoph
>gi|1037003468|ref|XP_017149657.1| PREDICTED: probable cytochrome P450 4p2 isoform X1 [Drosoph
>gi|1037003540|ref|XP_017149662.1| PREDICTED: cytochrome P450 4p1-like [Drosophila miranda]
>gi|1037003522|ref|XP_017149661.1| PREDICTED: cytochrome P450 4p1-like [Drosophila miranda]
>gi|1036960568|ref|XP_017147048.1| PREDICTED: uncharacterized protein LOC108158838 [Drosophila
>gi|1036960212|ref|XP_017147030.1| PREDICTED: serine/threonine-protein kinase N isoform X6 [Dro
>gi|1036960234|ref|XP_017147031.1| PREDICTED: serine/threonine-protein kinase N isoform X6 [Dro
>gi|1036960252|ref|XP_017147032.1| PREDICTED: serine/threonine-protein kinase N isoform X6 [Dro
>gi|1036960268|ref|XP_017147033.1| PREDICTED: serine/threonine-protein kinase N isoform X6 [Dro
>gi|1036960310|ref|XP_017147036.1| PREDICTED: serine/threonine-protein kinase N isoform X9 [Dro
>gi|1036960282|ref|XP_017147034.1| PREDICTED: serine/threonine-protein kinase N isoform X7 [Dro
>gi|1036960413|ref|XP_017147041.1| PREDICTED: serine/threonine-protein kinase N isoform X13 [Dr
>gi|1036960194|ref|XP_017147029.1| PREDICTED: serine/threonine-protein kinase N isoform X5 [Dro
>gi|1036960296|ref|XP_017147035.1| PREDICTED: serine/threonine-protein kinase N isoform X8 [Dro
>gi|1036960176|ref|XP_017147028.1| PREDICTED: serine/threonine-protein kinase N isoform X4 [Dro
>gi|1036960159|ref|XP_017147026.1| PREDICTED: serine/threonine-protein kinase N isoform X3 [Dro
>gi|1036960141|ref|XP_017147025.1| PREDICTED: serine/threonine-protein kinase N isoform X2 [Dro
>gi|1036960354|ref|XP_017147039.1| PREDICTED: serine/threonine-protein kinase N isoform X11 [Dr
>gi|1036960372|ref|XP_017147040.1| PREDICTED: serine/threonine-protein kinase N isoform X12 [Dr
>gi|1036960096|ref|XP_017147024.1| PREDICTED: serine/threonine-protein kinase N isoform X1 [Dro
>gi|1036960338|ref|XP_017147037.1| PREDICTED: serine/threonine-protein kinase N isoform X10 [Dr
>gi|1036960444|ref|XP_017147042.1| PREDICTED: uncharacterized protein LOC108158834 [Drosophila
>gi|1036960462|ref|XP_017147043.1| PREDICTED: uncharacterized protein LOC108158834 [Drosophila
>gi|1036960512|ref|XP_017147046.1| PREDICTED: SAP30-binding protein [Drosophila miranda]
>gi|1036960496|ref|XP_017147045.1| PREDICTED: conserved oligomeric Golgi complex subunit 6 [Dro
>gi|1036960478|ref|XP_017147044.1| PREDICTED: mitotic spindle assembly checkpoint protein MAD1
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>gi|1037000844|ref|XP_017149501.1| PREDICTED: uncharacterized protein LOC108160184 isoform X2
>gi|1037000821|ref|XP_017149500.1| PREDICTED: uncharacterized protein LOC108160184 isoform X1
>gi|1037009458|ref|XP_017150022.1| PREDICTED: myeloid differentiation primary response protein
>gi|1036975115|ref|XP_017147921.1| PREDICTED: uncharacterized protein LOC108159274 [Drosophila
>gi|1036975137|ref|XP_017147922.1| PREDICTED: uncharacterized protein LOC108159274 [Drosophila
>gi|1036975099|ref|XP_017147920.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036975209|ref|XP_017147927.1| PREDICTED: odorant receptor 45a [Drosophila miranda]
>gi|1036975293|ref|XP_017147932.1| PREDICTED: uncharacterized protein LOC108159283 [Drosophila
>gi|1036975259|ref|XP_017147930.1| PREDICTED: dynein light chain 1, axonemal [Drosophila mirand
>gi|1036975155|ref|XP_017147923.1| PREDICTED: nucleolar GTP-binding protein 1 [Drosophila miran
>gi|1036975245|ref|XP_017147929.1| PREDICTED: protein lethal(2)k10201 [Drosophila miranda]
>gi|1036975276|ref|XP_017147931.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1036975173|ref|XP_017147925.1| PREDICTED: putative phosphatidate phosphatase isoform X1 [Dr
>gi|1036975191|ref|XP_017147926.1| PREDICTED: putative phosphatidate phosphatase isoform X2 [Dr
>gi|1036975227|ref|XP_017147928.1| PREDICTED: putative phosphatidate phosphatase [Drosophila mi
>gi|1036945441|ref|XP_017146147.1| PREDICTED: uncharacterized abhydrolase domain-containing pro
>gi|1036999841|ref|XP_017149438.1| PREDICTED: uncharacterized protein LOC108160142 [Drosophila
>gi|1037002039|ref|XP_017149573.1| PREDICTED: sestrin homolog isoform X2 [Drosophila miranda]
>gi|1037002019|ref|XP_017149572.1| PREDICTED: sestrin homolog isoform X1 [Drosophila miranda]
>gi|1037002057|ref|XP_017149574.1| PREDICTED: purine nucleoside phosphorylase 1 [Drosophila mi
>gi|1036954310|ref|XP_017146698.1| PREDICTED: lethal(2)neighbour of tid protein [Drosophila mi
>gi|1036954285|ref|XP_017146696.1| PREDICTED: protein tumorous imaginal discs, mitochondrial is
>gi|1036954297|ref|XP_017146697.1| PREDICTED: protein tumorous imaginal discs, mitochondrial is
>gi|1036954273|ref|XP_017146695.1| PREDICTED: protein lethal(2)denticleless [Drosophila miranda
>gi|1036954407|ref|XP_017146705.1| PREDICTED: tumor necrosis factor alpha-induced protein 8-lik
>gi|1036954424|ref|XP_017146706.1| PREDICTED: tumor necrosis factor alpha-induced protein 8-lik
>gi|1036954440|ref|XP_017146707.1| PREDICTED: 39S ribosomal protein L43, mitochondrial [Drosopl
>gi|1036954259|ref|XP_017146694.1| PREDICTED: actin-binding protein anillin [Drosophila miranda
>gi|1036954359|ref|XP_017146702.1| PREDICTED: protein angel [Drosophila miranda]
>gi|1036954344|ref|XP_017146700.1| PREDICTED: peptidyl-alpha-hydroxyglycine alpha-amidating lya
>gi|1036954327|ref|XP_017146699.1| PREDICTED: peptidyl-alpha-hydroxyglycine alpha-amidating lya
>gi|1036954464|ref|XP_017146709.1| PREDICTED: uncharacterized protein LOC108158687 [Drosophila
>gi|1036954454|ref|XP_017146708.1| PREDICTED: protein lethal(2)essential for life [Drosophila r
>gi|1036954395|ref|XP_017146704.1| PREDICTED: estrogen sulfotransferase-like [Drosophila mirand
>gi|1036954378|ref|XP_017146703.1| PREDICTED: amine sulfotransferase [Drosophila miranda]
>gi|1036999508|ref|XP_017149422.1| PREDICTED: high affinity cAMP-specific and IBMX-insensitive
>gi|1036999494|ref|XP_017149421.1| PREDICTED: high affinity cAMP-specific and IBMX-insensitive
>gi|1036999524|ref|XP_017149423.1| PREDICTED: high affinity cAMP-specific and IBMX-insensitive
>gi|1036999476|ref|XP_017149420.1| PREDICTED: high affinity cAMP-specific and IBMX-insensitive
>gi|1036999458|ref|XP_017149418.1| PREDICTED: high affinity cAMP-specific and IBMX-insensitive
>gi|1036989394|ref|XP_017148804.1| PREDICTED: protein dead ringer isoform X3 [Drosophila mirand
>gi|1036989380|ref|XP_017148803.1| PREDICTED: protein dead ringer isoform X2 [Drosophila mirand
>gi|1036989364|ref|XP_017148802.1| PREDICTED: protein dead ringer isoform X1 [Drosophila mirand
>gi|1036989410|ref|XP_017148805.1| PREDICTED: protein dead ringer isoform X4 [Drosophila mirand
>gi|1037012225|ref|XP_017150183.1| PREDICTED: aquaporin isoform X1 [Drosophila miranda]
>gi|1037012243|ref|XP_017150184.1| PREDICTED: aquaporin isoform X2 [Drosophila miranda]
>gi|1037012293|ref|XP_017150187.1| PREDICTED: aquaporin AQPcic isoform X2 [Drosophila miranda]
>gi|1037012275|ref|XP_017150186.1| PREDICTED: aquaporin AQPcic isoform X1 [Drosophila miranda]
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>gi|1037012309|ref|XP_017150189.1| PREDICTED: aquaporin AQPcic isoform X3 [Drosophila miranda]
>gi|1037012327|ref|XP_017150190.1| PREDICTED: aquaporin AQPcic isoform X3 [Drosophila miranda]
>gi|1037012345|ref|XP_017150191.1| PREDICTED: aquaporin [Drosophila miranda]
>gi|1037012257|ref|XP_017150185.1| PREDICTED: aquaporin isoform X3 [Drosophila miranda]
>gi|1036955290|ref|XP_017146750.1| PREDICTED: uncharacterized protein LOC108158718 [Drosophila
>gi|1036955338|ref|XP_017146753.1| PREDICTED: immune-induced peptide 18 [Drosophila miranda]
>gi|1036955182|ref|XP_017146746.1| PREDICTED: uncharacterized protein LOC108158714 [Drosophila
>gi|1036937580|ref|XP_017145667.1| PREDICTED: uncharacterized protein LOC108158034, partial [D
>gi|1036955105|ref|XP_017146744.1| PREDICTED: phosphatidylinositol 3-kinase catalytic subunit
>gi|1036955258|ref|XP_017146749.1| PREDICTED: exosome complex component RRP4 [Drosophila miran
>gi|1036955128|ref|XP_017146745.1| PREDICTED: zinc finger protein 629 [Drosophila miranda]
>gi|1036955198|ref|XP_017146747.1| PREDICTED: caspase-1 isoform X1 [Drosophila miranda]
>gi|1036955216|ref|XP_017146748.1| PREDICTED: caspase-1 isoform X2 [Drosophila miranda]
>gi|1036955306|ref|XP_017146752.1| PREDICTED: cytochrome c oxidase subunit 6A, mitochondrial [
>gi|1036999121|ref|XP_017149401.1| PREDICTED: serine-threonine kinase receptor-associated prot
>gi|1036999135|ref|XP_017149402.1| PREDICTED: eukaryotic translation elongation factor 1 epsil
>gi|1036937598|ref|XP_017145668.1| PREDICTED: putative gustatory receptor 59f [Drosophila mira
>gi|1036999107|ref|XP_017149400.1| PREDICTED: putative gustatory receptor 59e [Drosophila mira
>gi|1037008697|ref|XP_017149975.1| PREDICTED: protein eiger isoform X1 [Drosophila miranda]
>gi|1037008714|ref|XP_017149976.1| PREDICTED: protein eiger isoform X2 [Drosophila miranda]
>gi|1037016566|ref|XP_017150442.1| PREDICTED: facilitated trehalose transporter Tret1 [Drosoph
>gi|1037016510|ref|XP_017150438.1| PREDICTED: uncharacterized protein LOC108160752 isoform X1
>gi|1037016528|ref|XP_017150439.1| PREDICTED: uncharacterized protein LOC108160752 isoform X1
>gi|1037016548|ref|XP_017150440.1| PREDICTED: uncharacterized protein LOC108160752 isoform X2
>gi|1037007377|ref|XP_017149897.1| PREDICTED: transcription factor AP-1 [Drosophila miranda]
>gi|1037007397|ref|XP_017149898.1| PREDICTED: mRNA turnover protein 4 homolog [Drosophila mira
>gi|1036981448|ref|XP_017148319.1| PREDICTED: 14-3-3 protein zeta isoform X1 [Drosophila miran
>gi|1036981462|ref|XP_017148320.1| PREDICTED: 14-3-3 protein zeta isoform X1 [Drosophila miran
>gi|1036981480|ref|XP_017148321.1| PREDICTED: 14-3-3 protein zeta isoform X2 [Drosophila miran
>gi|1036981494|ref|XP_017148322.1| PREDICTED: 14-3-3 protein zeta isoform X3 [Drosophila miran
>gi|1036981291|ref|XP_017148309.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X2 [
>gi|1036981321|ref|XP_017148310.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X3 [
>gi|1036981271|ref|XP_017148308.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X1 [
>gi|1036981424|ref|XP_017148317.1| PREDICTED: ATP-dependent 6-phosphofructokinase 1 isoform X7
>gi|1036981390|ref|XP_017148315.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X5 [
>gi|1036981406|ref|XP_017148316.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X6 [
>gi|1036981339|ref|XP_017148312.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X4 [
>gi|1036981356|ref|XP_017148313.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X4 [
>gi|1036981372|ref|XP_017148314.1| PREDICTED: ATP-dependent 6-phosphofructokinase isoform X4 [
>gi|1036981436|ref|XP_017148318.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1037019959|ref|XP_017150663.1| PREDICTED: uncharacterized protein LOC108160902 [Drosophila
>gi|1037019945|ref|XP_017150662.1| PREDICTED: uncharacterized protein LOC108160900 isoform X2
>gi|1037019930|ref|XP_017150661.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1036937616|ref|XP_017145670.1| PREDICTED: angiotensin-converting enzyme, partial [Drosophi
>gi|1037008951|ref|XP_017149990.1| PREDICTED: phenoloxidase 2 [Drosophila miranda]
>gi|1037008986|ref|XP_017149992.1| PREDICTED: putative sodium-coupled neutral amino acid transp
>gi|1037008968|ref|XP_017149991.1| PREDICTED: putative sodium-coupled neutral amino acid transp
>gi|1037021073|ref|XP_017150736.1| PREDICTED: leucine-rich repeat-containing protein 34 isoform
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>gi|1037021057|ref|XP_017150734.1| PREDICTED: leucine-rich repeat-containing protein 34 isofor
>gi|1037010348|ref|XP_017150071.1| PREDICTED: protein anachronism isoform X1 [Drosophila miran
>gi|1037010366|ref|XP_017150072.1| PREDICTED: protein anachronism isoform X2 [Drosophila miran
>gi|1037010384|ref|XP_017150073.1| PREDICTED: protein anachronism isoform X3 [Drosophila miran
>gi|1037010310|ref|XP_017150069.1| PREDICTED: solute carrier family 28 member 3 [Drosophila mi
>gi|1037010328|ref|XP_017150070.1| PREDICTED: solute carrier family 28 member 3 [Drosophila mi
>gi|1037003374|ref|XP_017149650.1| PREDICTED: solute carrier family 28 member 3 isoform X1 [Dro
>gi|1037003390|ref|XP_017149651.1| PREDICTED: solute carrier family 28 member 3 isoform X2 [Dro
>gi|1037003344|ref|XP_017149649.1| PREDICTED: uncharacterized protein LOC108160278 [Drosophila
>gi|1036993081|ref|XP_017149024.1| PREDICTED: probable ATP-dependent RNA helicase Dbp45A isofor
>gi|1036993097|ref|XP_017149025.1| PREDICTED: probable ATP-dependent RNA helicase Dbp45A isofor
>gi|1036993113|ref|XP_017149026.1| PREDICTED: probable ATP-dependent RNA helicase Dbp45A isofor
>gi|1036993049|ref|XP_017149022.1| PREDICTED: NAD kinase 2, mitochondrial-like isoform X4 [Dros
>gi|1036993013|ref|XP_017149020.1| PREDICTED: NAD kinase 2, mitochondrial-like isoform X2 [Dros
>gi|1036993067|ref|XP_017149023.1| PREDICTED: NAD kinase 2, mitochondrial-like isoform X5 [Dros
>gi|1036993031|ref|XP_017149021.1| PREDICTED: NAD kinase 2, mitochondrial-like isoform X3 [Dros
>gi|1036992995|ref|XP_017149019.1| PREDICTED: putative uncharacterized protein DDB_G0282133 iso
>gi|1036987901|ref|XP_017148714.1| PREDICTED: cytoplasmic tRNA 2-thiolation protein 1 [Drosophi
>gi|1036987847|ref|XP_017148711.1| PREDICTED: tetratricopeptide repeat protein 37 isoform X1 [
>gi|1036987865|ref|XP_017148712.1| PREDICTED: tetratricopeptide repeat protein 37 isoform X2 [
>gi|1036987883|ref|XP_017148713.1| PREDICTED: tetratricopeptide repeat protein 37 isoform X3 [
>gi|1036968728|ref|XP_017147531.1| PREDICTED: vang-like protein 1 [Drosophila miranda]
>gi|1036968710|ref|XP_017147529.1| PREDICTED: phosphoglucomutase-2 [Drosophila miranda]
>gi|1036968770|ref|XP_017147533.1| PREDICTED: RNA-binding protein 8A [Drosophila miranda]
>gi|1036968694|ref|XP_017147528.1| PREDICTED: protein SDA1 homolog [Drosophila miranda]
>gi|1036968752|ref|XP_017147532.1| PREDICTED: phosphorylated adapter RNA export protein [Drosop
>gi|1036953721|ref|XP_017146662.1| PREDICTED: E3 SUMO-protein ligase PIAS1 isoform X4 [Drosophi
>gi|1036953671|ref|XP_017146659.1| PREDICTED: E3 SUMO-protein ligase PIAS2 isoform X1 [Drosophi
>gi|1036953702|ref|XP_017146661.1| PREDICTED: E3 SUMO-protein ligase PIAS2 isoform X3 [Drosophi
>gi|1036953687|ref|XP_017146660.1| PREDICTED: E3 SUMO-protein ligase PIAS2 isoform X2 [Drosophi
>gi|1036953826|ref|XP_017146668.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1036953838|ref|XP_017146669.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1036953605|ref|XP_017146655.1| PREDICTED: DENN domain-containing protein 1A isoform X6 [Dro
>gi|1036953556|ref|XP_017146651.1| PREDICTED: uncharacterized protein LOC108158664 isoform X3
>gi|1036953590|ref|XP_017146654.1| PREDICTED: DENN domain-containing protein 1A isoform X5 [Dro
>gi|1036953573|ref|XP_017146652.1| PREDICTED: DENN domain-containing protein 1A isoform X4 [Dro
>gi|1036953537|ref|XP_017146650.1| PREDICTED: uncharacterized protein LOC108158664 isoform X2
>gi|1036953522|ref|XP_017146649.1| PREDICTED: uncharacterized protein LOC108158664 isoform X1
>gi|1036953619|ref|XP_017146656.1| PREDICTED: COPII coat assembly protein sec16 isoform X7 [Dro
>gi|1036953635|ref|XP_017146657.1| PREDICTED: COPII coat assembly protein sec16 isoform X7 [Dro
>gi|1036953653|ref|XP_017146658.1| PREDICTED: DENN domain-containing protein 1A isoform X8 [Dro
>gi|1036953752|ref|XP_017146663.1| PREDICTED: phospholipase ABHD3 isoform X1 [Drosophila miran
>gi|1036953770|ref|XP_017146664.1| PREDICTED: phospholipase ABHD3 isoform X2 [Drosophila miran
>gi|1036953788|ref|XP_017146665.1| PREDICTED: phospholipase ABHD3 isoform X2 [Drosophila miran
>gi|1036953808|ref|XP_017146666.1| PREDICTED: 5'-AMP-activated protein kinase subunit beta-1 [
>gi|1036953851|ref|XP_017146670.1| PREDICTED: uncharacterized protein LOC108158670 [Drosophila
>gi|1036953506|ref|XP_017146648.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036967221|ref|XP_017147444.1| PREDICTED: protein lava lamp [Drosophila miranda]
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>gi|1036967242|ref|XP_017147445.1| PREDICTED: uncharacterized protein LOC108159035 [Drosophila
>gi|1036980782|ref|XP_017148277.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036980788|ref|XP_017148279.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036980804|ref|XP_017148280.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036980818|ref|XP_017148281.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036980830|ref|XP_017148282.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036980844|ref|XP_017148283.1| PREDICTED: acylphosphatase-2 [Drosophila miranda]
>gi|1037017731|ref|XP_017150518.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1037017679|ref|XP_017150514.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1037017696|ref|XP_017150516.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1037017713|ref|XP_017150517.1| PREDICTED: serine proteinase stubble isoform X2 [Drosophila
>gi|1036942136|ref|XP_017145950.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036942188|ref|XP_017145953.1| PREDICTED: uncharacterized protein LOC108158232 [Drosophila
>gi|1036942206|ref|XP_017145954.1| PREDICTED: uncharacterized protein LOC108158232 [Drosophila
>gi|1036994854|ref|XP_017149136.1| PREDICTED: TGF-beta receptor type-1 isoform X3 [Drosophila
>gi|1036994836|ref|XP_017149135.1| PREDICTED: TGF-beta receptor type-1 isoform X2 [Drosophila
>gi|1036994818|ref|XP_017149134.1| PREDICTED: TGF-beta receptor type-1 isoform X1 [Drosophila
>gi|1036994872|ref|XP_017149137.1| PREDICTED: TGF-beta receptor type-1 isoform X4 [Drosophila
>gi|1037010222|ref|XP_017150064.1| PREDICTED: ADP-ribosylation factor 3 [Drosophila miranda]
>gi|1037010240|ref|XP_017150065.1| PREDICTED: ADP-ribosylation factor 3 [Drosophila miranda]
>gi|1037020699|ref|XP_017150710.1| PREDICTED: photoreceptor-specific nuclear receptor [Drosoph
>gi|1036981064|ref|XP_017148296.1| PREDICTED: uncharacterized protein LOC108159479 [Drosophila
>gi|1036981081|ref|XP_017148297.1| PREDICTED: uncharacterized protein LOC108159479 [Drosophila
>gi|1036980992|ref|XP_017148292.1| PREDICTED: uncharacterized protein LOC108159476 [Drosophila
>gi|1036981028|ref|XP_017148294.1| PREDICTED: E3 ubiquitin-protein ligase TRAIP isoform X1 [Dro
>gi|1036981046|ref|XP_017148295.1| PREDICTED: E3 ubiquitin-protein ligase TRAIP isoform X2 [Dro
>gi|1036981010|ref|XP_017148293.1| PREDICTED: armadillo repeat-containing protein 6 homolog [Di
>gi|1036952846|ref|XP_017146606.1| PREDICTED: uncharacterized protein LOC108158635 isoform X2
>gi|1036952828|ref|XP_017146605.1| PREDICTED: uncharacterized protein LOC108158635 isoform X1
>gi|1036952950|ref|XP_017146612.1| PREDICTED: lithostathine-2 [Drosophila miranda]
>gi|1036952932|ref|XP_017146611.1| PREDICTED: uncharacterized protein LOC108158638 [Drosophila
>gi|1036952862|ref|XP_017146607.1| PREDICTED: extracellular serine/threonine protein kinase fou
>gi|1036952898|ref|XP_017146609.1| PREDICTED: extracellular serine/threonine protein kinase fou
>gi|1036952880|ref|XP_017146608.1| PREDICTED: extracellular serine/threonine protein kinase fou
>gi|1036953098|ref|XP_017146622.1| PREDICTED: LOW QUALITY PROTEIN: otefin [Drosophila miranda]
>gi|1036953054|ref|XP_017146619.1| PREDICTED: gamma-secretase subunit pen-2 [Drosophila miranda
>gi|1036953068|ref|XP_017146620.1| PREDICTED: mucin-19 isoform X1 [Drosophila miranda]
>gi|1036953080|ref|XP_017146621.1| PREDICTED: mucin-19 isoform X2 [Drosophila miranda]
>gi|1036952916|ref|XP_017146610.1| PREDICTED: uncharacterized protein LOC108158637 [Drosophila
>gi|1036952968|ref|XP_017146613.1| PREDICTED: uncharacterized protein LOC108158641 [Drosophila
>gi|1036953003|ref|XP_017146616.1| PREDICTED: uncharacterized protein LOC108158643 [Drosophila
>gi|1036953020|ref|XP_017146617.1| PREDICTED: uncharacterized protein LOC108158644 [Drosophila
>gi|1036952986|ref|XP_017146614.1| PREDICTED: uncharacterized protein LOC108158642 [Drosophila
>gi|1036953036|ref|XP_017146618.1| PREDICTED: UPF0279 protein CG14505 [Drosophila miranda]
>gi|1036943584|ref|XP_017146040.1| PREDICTED: uncharacterized protein LOC108158311 [Drosophila
>gi|1037018918|ref|XP_017150597.1| PREDICTED: immune-induced peptide 23 [Drosophila miranda]
>gi|1037018900|ref|XP_017150596.1| PREDICTED: uncharacterized protein LOC108160850 [Drosophila
>gi|1037018934|ref|XP_017150598.1| PREDICTED: immune-induced peptide 2 [Drosophila miranda]
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>gi|1037018950|ref|XP_017150600.1| PREDICTED: immune-induced peptide 3-like [Drosophila miranda
>gi|1036943944|ref|XP_017146062.1| PREDICTED: uncharacterized protein LOC108158332 [Drosophila
>gi|1036943006|ref|XP_017146004.1| PREDICTED: immune-induced peptide 3-like [Drosophila miranda
>gi|1037021107|ref|XP_017150738.1| PREDICTED: chitinase-like protein Idgf5 [Drosophila miranda]
>gi|1036941115|ref|XP_017145887.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1036941165|ref|XP_017145890.1| PREDICTED: glutathione S-transferase 1-like [Drosophila mira
>gi|1037019973|ref|XP_017150664.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1037006394|ref|XP_017149835.1| PREDICTED: glutathione S-transferase 1-like [Drosophila mira
>gi|1037006358|ref|XP_017149832.1| PREDICTED: glutathione S-transferase 1-like [Drosophila mira
>gi|1037006342|ref|XP_017149831.1| PREDICTED: glutathione S-transferase 1-like [Drosophila mira
>gi|1037006376|ref|XP_017149834.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1037006324|ref|XP_017149830.1| PREDICTED: uncharacterized protein LOC108160369 [Drosophila
>gi|1036953867|ref|XP_017146671.1| PREDICTED: vigilin [Drosophila miranda]
>gi|1036953885|ref|XP_017146672.1| PREDICTED: vigilin [Drosophila miranda]
>gi|1036954141|ref|XP_017146686.1| PREDICTED: tumor protein D54 isoform X5 [Drosophila miranda]
>gi|1036954223|ref|XP_017146691.1| PREDICTED: tumor protein D52 isoform X10 [Drosophila miranda
>gi|1036954153|ref|XP_017146687.1| PREDICTED: tumor protein D54 isoform X6 [Drosophila miranda]
>gi|1036954169|ref|XP_017146688.1| PREDICTED: tumor protein D54 isoform X7 [Drosophila miranda]
>gi|1036954185|ref|XP_017146689.1| PREDICTED: tumor protein D52 isoform X8 [Drosophila miranda]
>gi|1036954205|ref|XP_017146690.1| PREDICTED: tumor protein D52 isoform X9 [Drosophila miranda]
>gi|1036954240|ref|XP_017146692.1| PREDICTED: tumor protein D52 isoform X11 [Drosophila miranda
>gi|1036954029|ref|XP_017146681.1| PREDICTED: uncharacterized protein LOC108158675 isoform X1
>gi|1036954045|ref|XP_017146683.1| PREDICTED: tumor protein D54 isoform X2 [Drosophila miranda]
>gi|1036954067|ref|XP_017146684.1| PREDICTED: tumor protein D52 isoform X3 [Drosophila miranda]
>gi|1036954119|ref|XP_017146685.1| PREDICTED: tumor protein D52 isoform X4 [Drosophila miranda]
>gi|1036953901|ref|XP_017146673.1| PREDICTED: GPI ethanolamine phosphate transferase 3 [Drosoph
>gi|1036954015|ref|XP_017146680.1| PREDICTED: TRPL translocation defect protein 14 isoform X4
>gi|1036953981|ref|XP_017146678.1| PREDICTED: TRPL translocation defect protein 14 isoform X2
>gi|1036953995|ref|XP_017146679.1| PREDICTED: TRPL translocation defect protein 14 isoform X3
>gi|1036953919|ref|XP_017146674.1| PREDICTED: TRPL translocation defect protein 14 isoform X1
>gi|1036953935|ref|XP_017146675.1| PREDICTED: TRPL translocation defect protein 14 isoform X1
>gi|1036953965|ref|XP_017146677.1| PREDICTED: TRPL translocation defect protein 14 isoform X1
>gi|1036972736|ref|XP_017147773.1| PREDICTED: kelch domain-containing protein 10 homolog [Droso
>gi|1036972857|ref|XP_017147782.1| PREDICTED: ragulator complex protein LAMTOR2 homolog [Drosop
>gi|1036972750|ref|XP_017147775.1| PREDICTED: pre-mRNA-processing factor 19 [Drosophila miranda
>gi|1036972780|ref|XP_017147777.1| PREDICTED: uncharacterized protein LOC108159211 isoform X1
>gi|1036972797|ref|XP_017147778.1| PREDICTED: uncharacterized protein LOC108159211 isoform X2
>gi|1036972809|ref|XP_017147779.1| PREDICTED: uncharacterized protein LOC108159211 isoform X3
>gi|1036972823|ref|XP_017147780.1| PREDICTED: uncharacterized protein LOC108159211 isoform X4
>gi|1036972842|ref|XP_017147781.1| PREDICTED: mitochondrial ribonuclease P protein 1 homolog [
>gi|1036972762|ref|XP_017147776.1| PREDICTED: dynein assembly factor 3, axonemal homolog [Droso
>gi|1036972722|ref|XP_017147772.1| PREDICTED: NACHT domain- and WD repeat-containing protein 1
>gi|1036972709|ref|XP_017147771.1| PREDICTED: NACHT domain- and WD repeat-containing protein 1
>gi|1036972655|ref|XP_017147768.1| PREDICTED: NACHT domain- and WD repeat-containing protein 1
>gi|1036972673|ref|XP_017147769.1| PREDICTED: NACHT domain- and WD repeat-containing protein 1
>gi|1036972691|ref|XP_017147770.1| PREDICTED: NACHT domain- and WD repeat-containing protein 1
>gi|1037016612|ref|XP_017150445.1| PREDICTED: glutathione S-transferase 1 [Drosophila miranda]
>gi|1037008559|ref|XP_017149966.1| PREDICTED: tubulin-specific chaperone E [Drosophila miranda]
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>gi|1036985912|ref|XP_017148595.1| PREDICTED: transmembrane protein 164 isoform X1 [Drosophila
>gi|1036985930|ref|XP_017148597.1| PREDICTED: transmembrane protein 164 isoform X2 [Drosophila
>gi|1036985952|ref|XP_017148598.1| PREDICTED: transmembrane protein 164 isoform X2 [Drosophila
>gi|1036985807|ref|XP_017148590.1| PREDICTED: sterol regulatory element-binding protein cleavag
>gi|1036985857|ref|XP_017148592.1| PREDICTED: sterol regulatory element-binding protein cleavag
>gi|1036985824|ref|XP_017148591.1| PREDICTED: sterol regulatory element-binding protein cleavag
>gi|1036985869|ref|XP_017148593.1| PREDICTED: caspase-1 [Drosophila miranda]
>gi|1036985895|ref|XP_017148594.1| PREDICTED: peptide-N(4)-(N-acetyl-beta-glucosaminyl)asparag
>gi|1036973483|ref|XP_017147821.1| PREDICTED: actin-5C [Drosophila miranda]
>gi|1036973467|ref|XP_017147820.1| PREDICTED: tyrosine-protein kinase Src42A isoform X3 [Droso
>gi|1036973439|ref|XP_017147818.1| PREDICTED: uncharacterized protein LOC108159227 isoform X2
>gi|1036973411|ref|XP_017147817.1| PREDICTED: uncharacterized protein LOC108159227 isoform X1
>gi|1036973501|ref|XP_017147822.1| PREDICTED: proteasome subunit alpha type-1-like [Drosophila
>gi|1036971788|ref|XP_017147711.1| PREDICTED: dosage compensation regulator isoform X1 [Drosop
>gi|1036971803|ref|XP_017147712.1| PREDICTED: dosage compensation regulator isoform X2 [Drosop
>gi|1036971772|ref|XP_017147710.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036971821|ref|XP_017147714.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1036971861|ref|XP_017147716.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1036971839|ref|XP_017147715.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1037020731|ref|XP_017150713.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1037009912|ref|XP_017150045.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037009928|ref|XP_017150046.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036937632|ref|XP_017145671.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036995025|ref|XP_017149146.1| PREDICTED: hemicentin-1 isoform X2 [Drosophila miranda]
>gi|1036995045|ref|XP_017149148.1| PREDICTED: hemicentin-2 isoform X3 [Drosophila miranda]
>gi|1036994966|ref|XP_017149143.1| PREDICTED: hemicentin-2 isoform X1 [Drosophila miranda]
>gi|1036994996|ref|XP_017149144.1| PREDICTED: hemicentin-2 isoform X1 [Drosophila miranda]
>gi|1036995011|ref|XP_017149145.1| PREDICTED: hemicentin-2 isoform X1 [Drosophila miranda]
>gi|1036994952|ref|XP_017149142.1| PREDICTED: odorant receptor 42b [Drosophila miranda]
>gi|1036937650|ref|XP_017145672.1| PREDICTED: odorant receptor 42a [Drosophila miranda]
>gi|1036994936|ref|XP_017149141.1| PREDICTED: odorant receptor 42a-like [Drosophila miranda]
>gi|1036995080|ref|XP_017149150.1| PREDICTED: uncharacterized protein LOC108159974 [Drosophila
>gi|1036937667|ref|XP_017145673.1| PREDICTED: J domain-containing protein CG6693-like [Drosoph
>gi|1036937685|ref|XP_017145674.1| PREDICTED: alpha-amylase A-like [Drosophila miranda]
>gi|1036994924|ref|XP_017149140.1| PREDICTED: alpha-amylase A [Drosophila miranda]
>gi|1036937707|ref|XP_017145675.1| PREDICTED: alpha-amylase 2-like, partial [Drosophila miranda
>gi|1036995063|ref|XP_017149149.1| PREDICTED: uncharacterized protein LOC108159973 [Drosophila
>gi|1036994890|ref|XP_017149138.1| PREDICTED: zinc transporter 2 [Drosophila miranda]
>gi|1036994908|ref|XP_017149139.1| PREDICTED: zinc transporter 2 [Drosophila miranda]
>gi|1036954903|ref|XP_017146731.1| PREDICTED: nuclear pore membrane glycoprotein 210 isoform X
>gi|1036954883|ref|XP_017146730.1| PREDICTED: nuclear pore membrane glycoprotein 210 isoform X
>gi|1036955031|ref|XP_017146739.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 [Dro
>gi|1036955049|ref|XP_017146741.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 [Dro
>gi|1036955067|ref|XP_017146742.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 [Dro
>gi|1036954935|ref|XP_017146733.1| PREDICTED: mitochondrial intermediate peptidase isoform X1
>gi|1036954953|ref|XP_017146734.1| PREDICTED: mitochondrial intermediate peptidase isoform X2
>gi|1036955085|ref|XP_017146743.1| PREDICTED: uncharacterized protein LOC108158711 [Drosophila
>gi|1036955014|ref|XP_017146738.1| PREDICTED: glutamate-rich WD repeat-containing protein 1 [Dr

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>gi|1036954918|ref|XP_017146732.1| PREDICTED: mismatch repair endonuclease PMS2 [Drosophila mi
>gi|1036954969|ref|XP_017146735.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036954983|ref|XP_017146736.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036954999|ref|XP_017146737.1| PREDICTED: solute carrier family 2, facilitated glucose tran
>gi|1036978569|ref|XP_017148136.1| PREDICTED: DNA replication factor Cdt1 [Drosophila miranda]
>gi|1036978553|ref|XP_017148135.1| PREDICTED: SRSF protein kinase 3 [Drosophila miranda]
>gi|1036978589|ref|XP_017148137.1| PREDICTED: uncharacterized protein LOC108159407 [Drosophila
>gi|1036997307|ref|XP_017149288.1| PREDICTED: metchnikowin [Drosophila miranda]
>gi|1036997323|ref|XP_017149289.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036997261|ref|XP_017149284.1| PREDICTED: uncharacterized protein LOC108160050 isoform X2
>gi|1036997244|ref|XP_017149283.1| PREDICTED: uncharacterized protein LOC108160050 isoform X1
>gi|1036997291|ref|XP_017149287.1| PREDICTED: uncharacterized protein LOC108160051 isoform X2
>gi|1036997275|ref|XP_017149285.1| PREDICTED: uncharacterized protein LOC108160051 isoform X1
>gi|1037008607|ref|XP_017149970.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isofor
>gi|1037008626|ref|XP_017149971.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like isofor
>gi|1037008589|ref|XP_017149969.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like isofor
>gi|1037008575|ref|XP_017149967.1| PREDICTED: uncharacterized protein LOC108160464 [Drosophila
>gi|1037008643|ref|XP_017149972.1| PREDICTED: uncharacterized protein LOC108160466 [Drosophila
>gi|1037008661|ref|XP_017149973.1| PREDICTED: mastermind-like protein 2 [Drosophila miranda]
>gi|1036937725|ref|XP_017145677.1| PREDICTED: uncharacterized protein LOC108158042 [Drosophila
>gi|1036937742|ref|XP_017145678.1| PREDICTED: uncharacterized protein LOC108158043 [Drosophila
>gi|1037004098|ref|XP_017149695.1| PREDICTED: polypeptide N-acetylgalactosaminyltransferase 1
>gi|1036977437|ref|XP_017148066.1| PREDICTED: kinesin-like protein KIF13A isoform X6 [Drosophil
>gi|1036977388|ref|XP_017148062.1| PREDICTED: kinesin-like protein KIF13A isoform X3 [Drosophil
>gi|1036977422|ref|XP_017148065.1| PREDICTED: kinesin-like protein KIF13A isoform X5 [Drosophil
>gi|1036977406|ref|XP_017148063.1| PREDICTED: kinesin-like protein KIF13A isoform X4 [Drosophil
>gi|1036977370|ref|XP_017148061.1| PREDICTED: kinesin-like protein KIF13A isoform X2 [Drosophil
>gi|1036977352|ref|XP_017148060.1| PREDICTED: kinesin-like protein KIF13A isoform X1 [Drosophil
>gi|1037022021|ref|XP_017150793.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036988766|ref|XP_017148764.1| PREDICTED: protein saal1 [Drosophila miranda]
>gi|1036988837|ref|XP_017148768.1| PREDICTED: uncharacterized protein LOC108159742 [Drosophila
>gi|1036988855|ref|XP_017148770.1| PREDICTED: V-type proton ATPase subunit D 1 [Drosophila mir
>gi|1036988783|ref|XP_017148765.1| PREDICTED: translation initiation factor eIF-2B subunit gamm
>gi|1036988819|ref|XP_017148767.1| PREDICTED: GATA zinc finger domain-containing protein 10 is
>gi|1036988801|ref|XP_017148766.1| PREDICTED: GATA zinc finger domain-containing protein 10 is
>gi|1036949711|ref|XP_017146413.1| PREDICTED: uncharacterized protein LOC108158528 isoform X14
>gi|1036949645|ref|XP_017146409.1| PREDICTED: uncharacterized protein LOC108158528 isoform X10
>gi|1036949663|ref|XP_017146410.1| PREDICTED: uncharacterized protein LOC108158528 isoform X11
>gi|1036949488|ref|XP_017146399.1| PREDICTED: uncharacterized protein LOC108158528 isoform X1
>gi|1036949611|ref|XP_017146407.1| PREDICTED: uncharacterized protein LOC108158528 isoform X8
>gi|1036949744|ref|XP_017146414.1| PREDICTED: uncharacterized protein LOC108158528 isoform X15
>gi|1036949693|ref|XP_017146412.1| PREDICTED: uncharacterized protein LOC108158528 isoform X13
>gi|1036949506|ref|XP_017146400.1| PREDICTED: uncharacterized protein LOC108158528 isoform X2
>gi|1036949559|ref|XP_017146403.1| PREDICTED: uncharacterized protein LOC108158528 isoform X5
>gi|1036949575|ref|XP_017146404.1| PREDICTED: uncharacterized protein LOC108158528 isoform X6
>gi|1036949627|ref|XP_017146408.1| PREDICTED: uncharacterized protein LOC108158528 isoform X9
>gi|1036949593|ref|XP_017146405.1| PREDICTED: uncharacterized protein LOC108158528 isoform X7
>gi|1036949541|ref|XP_017146402.1| PREDICTED: uncharacterized protein LOC108158528 isoform X4
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>gi|1036949523|ref|XP_017146401.1| PREDICTED: uncharacterized protein LOC108158528 isoform X3
>gi|1036949677|ref|XP_017146411.1| PREDICTED: uncharacterized protein LOC108158528 isoform X12
>gi|1036949471|ref|XP_017146398.1| PREDICTED: copper chaperone for superoxide dismutase [Droso
>gi|1036949354|ref|XP_017146390.1| PREDICTED: F-box only protein 28 [Drosophila miranda]
>gi|1036949372|ref|XP_017146391.1| PREDICTED: F-box only protein 28 [Drosophila miranda]
>gi|1036949338|ref|XP_017146389.1| PREDICTED: USP6 N-terminal-like protein isoform X2 [Drosophi
>gi|1036949320|ref|XP_017146388.1| PREDICTED: USP6 N-terminal-like protein isoform X1 [Drosophi
>gi|1036949454|ref|XP_017146397.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036949762|ref|XP_017146415.1| PREDICTED: WD repeat and HMG-box DNA-binding protein 1 [Dros
>gi|1036949303|ref|XP_017146387.1| PREDICTED: DNA-binding protein SMUBP-2 [Drosophila miranda]
>gi|1036949437|ref|XP_017146395.1| PREDICTED: uncharacterized protein LOC108158525 [Drosophila
>gi|1036949406|ref|XP_017146393.1| PREDICTED: sulfotransferase 4A1 isoform X1 [Drosophila miran
>gi|1036949420|ref|XP_017146394.1| PREDICTED: sulfotransferase 4A1 isoform X2 [Drosophila miran
>gi|1036949388|ref|XP_017146392.1| PREDICTED: probable phenylalanine--tRNA ligase, mitochondria
>gi|1036986962|ref|XP_017148656.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036986973|ref|XP_017148658.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036986986|ref|XP_017148659.1| PREDICTED: uncharacterized protein LOC108159678 isoform X1
>gi|1036986997|ref|XP_017148660.1| PREDICTED: uncharacterized protein LOC108159678 isoform X2
>gi|1036941149|ref|XP_017145889.1| PREDICTED: 5-hydroxyisourate hydrolase [Drosophila miranda]
>gi|1036998074|ref|XP_017149335.1| PREDICTED: periodic tryptophan protein 1 homolog [Drosophila
>gi|1036998054|ref|XP_017149334.1| PREDICTED: transcription initiation factor TFIID subunit 5
>gi|1036947727|ref|XP_017146289.1| PREDICTED: peroxisome assembly factor 2 isoform X1 [Drosophi
>gi|1036947745|ref|XP_017146290.1| PREDICTED: peroxisome assembly factor 2 isoform X2 [Drosophi
>gi|1036947762|ref|XP_017146291.1| PREDICTED: peroxisomal (S)-2-hydroxy-acid oxidase GLO5 [Dros
>gi|1036947811|ref|XP_017146294.1| PREDICTED: uncharacterized protein LOC108158494 [Drosophila
>gi|1036947833|ref|XP_017146295.1| PREDICTED: U7 snRNA-associated Sm-like protein LSm11 [Drosop
>gi|1036947796|ref|XP_017146293.1| PREDICTED: LOW QUALITY PROTEIN: mitochondrial amidoxime-redu
>gi|1036947777|ref|XP_017146292.1| PREDICTED: mitochondrial amidoxime reducing component 2-like
>gi|1036947851|ref|XP_017146297.1| PREDICTED: vesicle-associated membrane protein 7 [Drosophila
>gi|1036947869|ref|XP_017146298.1| PREDICTED: vesicle-associated membrane protein 7 [Drosophila
>gi|1036947887|ref|XP_017146299.1| PREDICTED: vesicle-associated membrane protein 7 [Drosophila
>gi|1036947709|ref|XP_017146287.1| PREDICTED: uncharacterized protein LOC108158489 isoform X2
>gi|1036947691|ref|XP_017146286.1| PREDICTED: uncharacterized protein LOC108158489 isoform X1
>gi|1036947904|ref|XP_017146300.1| PREDICTED: beta-1,3-galactosyltransferase 1 [Drosophila mira
>gi|1036947921|ref|XP_017146301.1| PREDICTED: ribonucleoside-diphosphate reductase subunit M2
>gi|1036947956|ref|XP_017146303.1| PREDICTED: RNA-binding protein 48 [Drosophila miranda]
>gi|1036947938|ref|XP_017146302.1| PREDICTED: presenilins-associated rhomboid-like protein, mit
>gi|1036939443|ref|XP_017145787.1| PREDICTED: retinol dehydrogenase 12 [Drosophila miranda]
>gi|1036939407|ref|XP_017145785.1| PREDICTED: retinol dehydrogenase 12 isoform X1 [Drosophila m
>gi|1036939425|ref|XP_017145786.1| PREDICTED: retinol dehydrogenase 12 isoform X2 [Drosophila m
>gi|1036939461|ref|XP_017145789.1| PREDICTED: lysM and putative peptidoglycan-binding domain-co
>gi|1036939477|ref|XP_017145790.1| PREDICTED: lysM and putative peptidoglycan-binding domain-co
>gi|1036939318|ref|XP_017145779.1| PREDICTED: 26S protease regulatory subunit 7 [Drosophila mir
>gi|1036938722|ref|XP_017145743.1| PREDICTED: serine-rich adhesin for platelets isoform X10 [Dr
>gi|1036938597|ref|XP_017145735.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036938704|ref|XP_017145741.1| PREDICTED: serine-rich adhesin for platelets isoform X9 [Dro
>gi|1036938686|ref|XP_017145740.1| PREDICTED: serine-rich adhesin for platelets isoform X8 [Dro
>gi|1036938633|ref|XP_017145737.1| PREDICTED: serine-rich adhesin for platelets isoform X5 [Dro
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>gi|1036938561|ref|XP_017145733.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036938738|ref|XP_017145744.1| PREDICTED: serine-rich adhesin for platelets isoform X11 [Di
>gi|1036938577|ref|XP_017145734.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036938615|ref|XP_017145736.1| PREDICTED: serine-rich adhesin for platelets isoform X4 [Dro
>gi|1036938651|ref|XP_017145738.1| PREDICTED: serine-rich adhesin for platelets isoform X6 [Dro
>gi|1036938669|ref|XP_017145739.1| PREDICTED: serine-rich adhesin for platelets isoform X7 [Dro
>gi|1036938756|ref|XP_017145745.1| PREDICTED: uncharacterized protein LOC108158090 isoform X12
>gi|1036938791|ref|XP_017145747.1| PREDICTED: uncharacterized protein LOC108158090 isoform X14
>gi|1036938774|ref|XP_017145746.1| PREDICTED: uncharacterized protein LOC108158090 isoform X13
>gi|1036939599|ref|XP_017145797.1| PREDICTED: uncharacterized protein LOC108158127 [Drosophila
>gi|1036939246|ref|XP_017145775.1| PREDICTED: feline leukemia virus subgroup C receptor-related
>gi|1036939264|ref|XP_017145776.1| PREDICTED: feline leukemia virus subgroup C receptor-related
>gi|1036938896|ref|XP_017145753.1| PREDICTED: actin-binding protein anillin-like [Drosophila mi
>gi|1036939230|ref|XP_017145774.1| PREDICTED: actin-binding protein anillin-like isoform X3 [Di
>gi|1036939212|ref|XP_017145773.1| PREDICTED: actin-binding protein anillin-like isoform X2 [Di
>gi|1036939194|ref|XP_017145772.1| PREDICTED: actin-binding protein anillin-like isoform X1 [Di
>gi|1036939617|ref|XP_017145798.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036938950|ref|XP_017145757.1| PREDICTED: uncharacterized protein LOC108158101 [Drosophila
>gi|1036938964|ref|XP_017145758.1| PREDICTED: uncharacterized protein LOC108158101 [Drosophila
>gi|1036938982|ref|XP_017145759.1| PREDICTED: uncharacterized protein LOC108158101 [Drosophila
>gi|1036938419|ref|XP_017145724.1| PREDICTED: vacuolar protein sorting-associated protein 13C :
>gi|1036938437|ref|XP_017145725.1| PREDICTED: vacuolar protein sorting-associated protein 13C :
>gi|1036938454|ref|XP_017145726.1| PREDICTED: vacuolar protein sorting-associated protein 13C :
>gi|1036939563|ref|XP_017145795.1| PREDICTED: LDLR chaperone boca [Drosophila miranda]
>gi|1036939545|ref|XP_017145794.1| PREDICTED: ribulose-phosphate 3-epimerase [Drosophila mirano
>gi|1036938809|ref|XP_017145748.1| PREDICTED: uncharacterized protein LOC108158092 [Drosophila
>gi|1036939389|ref|XP_017145784.1| PREDICTED: ATPase ASNA1 homolog [Drosophila miranda]
>gi|1036938932|ref|XP_017145756.1| PREDICTED: glycerol-3-phosphate dehydrogenase, mitochondrial
>gi|1036939581|ref|XP_017145796.1| PREDICTED: cytochrome b5 [Drosophila miranda]
>gi|1036939336|ref|XP_017145781.1| PREDICTED: zinc-type alcohol dehydrogenase-like protein C17
>gi|1036939493|ref|XP_017145791.1| PREDICTED: uncharacterized protein LOC108158122 isoform X1
>gi|1036939509|ref|XP_017145792.1| PREDICTED: uncharacterized protein LOC108158122 isoform X2
>gi|1036939527|ref|XP_017145793.1| PREDICTED: uncharacterized protein LOC108158122 isoform X3
>gi|1036938826|ref|XP_017145749.1| PREDICTED: origin recognition complex subunit 1 [Drosophila
>gi|1036938844|ref|XP_017145750.1| PREDICTED: GPI ethanolamine phosphate transferase 2 [Drosopl
>gi|1036938914|ref|XP_017145755.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosopl
>gi|1036938999|ref|XP_017145760.1| PREDICTED: myoneurin [Drosophila miranda]
>gi|1036939019|ref|XP_017145761.1| PREDICTED: uncharacterized protein LOC108158103 [Drosophila
>gi|1036939037|ref|XP_017145762.1| PREDICTED: zinc finger protein 782 [Drosophila miranda]
>gi|1036939282|ref|XP_017145777.1| PREDICTED: elongation factor Tu [Drosophila miranda]
>gi|1036938489|ref|XP_017145729.1| PREDICTED: unconventional myosin-Va isoform X2 [Drosophila r
>gi|1036938472|ref|XP_017145728.1| PREDICTED: unconventional myosin-Vb isoform X1 [Drosophila r
>gi|1036938525|ref|XP_017145731.1| PREDICTED: unconventional myosin-Va isoform X4 [Drosophila r
>gi|1036938507|ref|XP_017145730.1| PREDICTED: unconventional myosin-Va isoform X3 [Drosophila r
>gi|1036938543|ref|XP_017145732.1| PREDICTED: unconventional myosin-Va isoform X5 [Drosophila r
>gi|1036938862|ref|XP_017145751.1| PREDICTED: DNA replication licensing factor MCM4 [Drosophila
>gi|1036939055|ref|XP_017145763.1| PREDICTED: mesoderm induction early response protein 1 [Dros
>gi|1036939373|ref|XP_017145783.1| PREDICTED: uncharacterized protein LOC108158117 [Drosophila
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>gi|1036938880|ref|XP_017145752.1| PREDICTED: protein wech [Drosophila miranda]
>gi|1036939124|ref|XP_017145768.1| PREDICTED: aldehyde dehydrogenase, dimeric NADP-preferring
>gi|1036939142|ref|XP_017145769.1| PREDICTED: aldehyde dehydrogenase, dimeric NADP-preferring
>gi|1036939160|ref|XP_017145770.1| PREDICTED: aldehyde dehydrogenase, dimeric NADP-preferring
>gi|1036939089|ref|XP_017145765.1| PREDICTED: aldehyde dehydrogenase, dimeric NADP-preferring
>gi|1036939071|ref|XP_017145764.1| PREDICTED: aldehyde dehydrogenase, dimeric NADP-preferring
>gi|1036939107|ref|XP_017145766.1| PREDICTED: aldehyde dehydrogenase, dimeric NADP-preferring
>gi|1036939178|ref|XP_017145771.1| PREDICTED: aldehyde dehydrogenase family 3 member B1 isoform
>gi|1036939355|ref|XP_017145782.1| PREDICTED: NSFL1 cofactor p47 [Drosophila miranda]
>gi|1036939300|ref|XP_017145778.1| PREDICTED: uncharacterized protein LOC108158113 [Drosophila
>gi|1036974775|ref|XP_017147899.1| PREDICTED: muskelin isoform X1 [Drosophila miranda]
>gi|1036974793|ref|XP_017147900.1| PREDICTED: muskelin isoform X2 [Drosophila miranda]
>gi|1036974876|ref|XP_017147907.1| PREDICTED: G protein alpha q subunit isoform X4 [Drosophila
>gi|1036974815|ref|XP_017147903.1| PREDICTED: G protein alpha q subunit isoform X1 [Drosophila
>gi|1036974832|ref|XP_017147904.1| PREDICTED: G protein alpha q subunit isoform X1 [Drosophila
>gi|1036974892|ref|XP_017147908.1| PREDICTED: G protein alpha q subunit isoform X1 [Drosophila
>gi|1036974906|ref|XP_017147909.1| PREDICTED: G protein alpha q subunit isoform X5 [Drosophila
>gi|1036974924|ref|XP_017147910.1| PREDICTED: G protein alpha q subunit isoform X6 [Drosophila
>gi|1036974847|ref|XP_017147905.1| PREDICTED: G protein alpha q subunit isoform X2 [Drosophila
>gi|1036974860|ref|XP_017147906.1| PREDICTED: G protein alpha q subunit isoform X3 [Drosophila
>gi|1036974940|ref|XP_017147911.1| PREDICTED: G protein alpha q subunit isoform X7 [Drosophila
>gi|1037006957|ref|XP_017149871.1| PREDICTED: myc box-dependent-interacting protein 1 isoform
>gi|1037006942|ref|XP_017149870.1| PREDICTED: bridging integrator 2 isoform X1 [Drosophila mira
>gi|1036945487|ref|XP_017146149.1| PREDICTED: uncharacterized protein LOC108158403 isoform X1
>gi|1036945520|ref|XP_017146151.1| PREDICTED: uncharacterized protein LOC108158403 isoform X3
>gi|1036945504|ref|XP_017146150.1| PREDICTED: uncharacterized protein LOC108158403 isoform X2
>gi|1036945538|ref|XP_017146153.1| PREDICTED: paired amphipathic helix protein Sin3a isoform X4
>gi|1036945556|ref|XP_017146154.1| PREDICTED: paired amphipathic helix protein Sin3a isoform X5
>gi|1036945645|ref|XP_017146160.1| PREDICTED: H(+)/Cl(-) exchange transporter 7 [Drosophila mir
>gi|1036945699|ref|XP_017146164.1| PREDICTED: protein stand still [Drosophila miranda]
>gi|1036945717|ref|XP_017146165.1| PREDICTED: protein stand still [Drosophila miranda]
>gi|1036945753|ref|XP_017146167.1| PREDICTED: adenylate kinase isoenzyme 6 homolog [Drosophila
>gi|1036945681|ref|XP_017146162.1| PREDICTED: probable cytochrome P450 301a1, mitochondrial iso
>gi|1036945663|ref|XP_017146161.1| PREDICTED: probable cytochrome P450 301a1, mitochondrial iso
>gi|1036945735|ref|XP_017146166.1| PREDICTED: uncharacterized protein LOC108158408 [Drosophila
>gi|1036945628|ref|XP_017146159.1| PREDICTED: neogenin isoform X4 [Drosophila miranda]
>gi|1036945610|ref|XP_017146157.1| PREDICTED: neogenin isoform X3 [Drosophila miranda]
>gi|1036945592|ref|XP_017146156.1| PREDICTED: neogenin isoform X2 [Drosophila miranda]
>gi|1036945574|ref|XP_017146155.1| PREDICTED: neogenin isoform X1 [Drosophila miranda]
>gi|1036945771|ref|XP_017146168.1| PREDICTED: uncharacterized protein LOC108158410 [Drosophila
>gi|1036945804|ref|XP_017146170.1| PREDICTED: uncharacterized protein LOC108158412 [Drosophila
>gi|1036945788|ref|XP_017146169.1| PREDICTED: uncharacterized protein LOC108158411 [Drosophila
>gi|1036937760|ref|XP_017145679.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037019236|ref|XP_017150617.1| PREDICTED: parkin coregulated gene protein homolog [Drosophi
>gi|1037019254|ref|XP_017150619.1| PREDICTED: uncharacterized protein LOC108160868 [Drosophila
>gi|1037012899|ref|XP_017150225.1| PREDICTED: thioredoxin-related transmembrane protein 2 homol
>gi|1036937778|ref|XP_017145680.1| PREDICTED: uncharacterized protein LOC108158046 [Drosophila
>gi|1036937794|ref|XP_017145681.1| PREDICTED: uncharacterized protein LOC108158047 [Drosophila
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>gi|1036942421|ref|XP_017145968.1| PREDICTED: uncharacterized protein LOC108158245 [Drosophila
>gi|1036971127|ref|XP_017147675.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971067|ref|XP_017147672.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971047|ref|XP_017147671.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971011|ref|XP_017147669.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036970995|ref|XP_017147667.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036970887|ref|XP_017147661.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971031|ref|XP_017147670.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036970977|ref|XP_017147666.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
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>gi|1036970941|ref|XP_017147664.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
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>gi|1036970921|ref|XP_017147663.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036970869|ref|XP_017147660.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036970853|ref|XP_017147659.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971109|ref|XP_017147674.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971087|ref|XP_017147673.1| PREDICTED: heterogeneous nuclear ribonucleoprotein L isofor
>gi|1036971187|ref|XP_017147678.1| PREDICTED: pinin [Drosophila miranda]
>gi|1036971169|ref|XP_017147677.1| PREDICTED: uncharacterized protein LOC108159163 isoform X18
>gi|1036971149|ref|XP_017147676.1| PREDICTED: uncharacterized protein LOC108159163 isoform X17
>gi|1036971204|ref|XP_017147679.1| PREDICTED: uncharacterized protein LOC108159165 [Drosophila
>gi|1036970835|ref|XP_017147657.1| PREDICTED: tubulin polyglutamylase ttll6 [Drosophila miranda
>gi|1036937810|ref|XP_017145682.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037022266|ref|XP_017150807.1| PREDICTED: uncharacterized protein LOC108161010 [Drosophila
>gi|1037022286|ref|XP_017150808.1| PREDICTED: uncharacterized protein LOC108161010 [Drosophila
>gi|1037022306|ref|XP_017150809.1| PREDICTED: uncharacterized protein LOC108161010 [Drosophila
>gi|1036941750|ref|XP_017145923.1| PREDICTED: 39 kDa FK506-binding nuclear protein [Drosophila
>gi|1036941768|ref|XP_017145924.1| PREDICTED: uncharacterized protein LOC108158212 [Drosophila
>gi|1036941927|ref|XP_017145936.1| PREDICTED: uncharacterized protein LOC108158219 isoform X2
>gi|1036941912|ref|XP_017145934.1| PREDICTED: uncharacterized protein LOC108158219 isoform X1
>gi|1037018830|ref|XP_017150592.1| PREDICTED: uncharacterized protein LOC108160847 [Drosophila
>gi|1036984463|ref|XP_017148502.1| PREDICTED: LOW QUALITY PROTEIN: poly(A) polymerase beta [Dro
>gi|1036984481|ref|XP_017148504.1| PREDICTED: ubiquitin-associated domain-containing protein 1
>gi|1036944991|ref|XP_017146119.1| PREDICTED: uncharacterized protein LOC108158378 [Drosophila
>gi|1037021776|ref|XP_017150777.1| PREDICTED: uncharacterized protein LOC108160985 [Drosophila
>gi|1037021756|ref|XP_017150776.1| PREDICTED: general odorant-binding protein 56a [Drosophila
>gi|1037021738|ref|XP_017150775.1| PREDICTED: general odorant-binding protein 56a [Drosophila
>gi|1036944923|ref|XP_017146116.1| PREDICTED: uncharacterized protein LOC108158372 [Drosophila
>gi|1036941061|ref|XP_017145883.1| PREDICTED: general odorant-binding protein 56d [Drosophila
>gi|1036937826|ref|XP_017145684.1| PREDICTED: odorant receptor 56a-like [Drosophila miranda]
>gi|1036937841|ref|XP_017145685.1| PREDICTED: odorant receptor 56a [Drosophila miranda]
>gi|1036942309|ref|XP_017145961.1| PREDICTED: uncharacterized protein LOC108158239 [Drosophila
>gi|1036942291|ref|XP_017145960.1| PREDICTED: general odorant-binding protein 56h [Drosophila
>gi|1036942888|ref|XP_017145997.1| PREDICTED: general odorant-binding protein 56h [Drosophila
>gi|1037007317|ref|XP_017149894.1| PREDICTED: protein toll isoform X2 [Drosophila miranda]
>gi|1037007303|ref|XP_017149893.1| PREDICTED: protein toll isoform X1 [Drosophila miranda]
>gi|1037020219|ref|XP_017150679.1| PREDICTED: uncharacterized protein LOC108160913 [Drosophila
>gi|1037016094|ref|XP_017150414.1| PREDICTED: organic cation transporter protein [Drosophila m
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>gi|1037016112|ref|XP_017150415.1| PREDICTED: organic cation transporter protein [Drosophila m:
>gi|1036942036|ref|XP_017145943.1| PREDICTED: uncharacterized protein LOC108158223 [Drosophila
>gi|1037000439|ref|XP_017149473.1| PREDICTED: protein toll isoform X1 [Drosophila miranda]
>gi|1037000449|ref|XP_017149474.1| PREDICTED: slit homolog 3 protein isoform X2 [Drosophila mi:
>gi|1036937857|ref|XP_017145686.1| PREDICTED: EF-hand calcium-binding domain-containing protei:
>gi|1036942224|ref|XP_017145955.1| PREDICTED: protein crossbronx-like [Drosophila miranda]
>gi|1037016008|ref|XP_017150409.1| PREDICTED: probable cytochrome P450 6a14 [Drosophila miranda
>gi|1037003326|ref|XP_017149648.1| PREDICTED: LOW QUALITY PROTEIN: probable cytochrome P450 6a:
>gi|1037003308|ref|XP_017149647.1| PREDICTED: probable cytochrome P450 6a13 [Drosophila miranda
>gi|1037003290|ref|XP_017149646.1| PREDICTED: cytochrome P450 6a22 [Drosophila miranda]
>gi|1037012173|ref|XP_017150180.1| PREDICTED: probable cytochrome P450 6a23 [Drosophila miranda
>gi|1037018008|ref|XP_017150534.1| PREDICTED: probable cytochrome P450 6a20 [Drosophila miranda
>gi|1037010416|ref|XP_017150077.1| PREDICTED: probable cytochrome P450 6a21 [Drosophila miranda
>gi|1037010434|ref|XP_017150078.1| PREDICTED: cytochrome P450 6a8 [Drosophila miranda]
>gi|1036992380|ref|XP_017148979.1| PREDICTED: probable cytochrome P450 317a1 [Drosophila miran:
>gi|1036992234|ref|XP_017148970.1| PREDICTED: uncharacterized protein LOC108159876 isoform X2
>gi|1036992256|ref|XP_017148971.1| PREDICTED: uncharacterized protein LOC108159876 isoform X2
>gi|1036992272|ref|XP_017148972.1| PREDICTED: uncharacterized protein LOC108159876 isoform X2
>gi|1036992290|ref|XP_017148973.1| PREDICTED: uncharacterized protein LOC108159876 isoform X3
>gi|1036992217|ref|XP_017148969.1| PREDICTED: uncharacterized protein LOC108159876 isoform X1
>gi|1036992344|ref|XP_017148977.1| PREDICTED: uncharacterized protein LOC108159876 isoform X6
>gi|1036992326|ref|XP_017148975.1| PREDICTED: uncharacterized protein LOC108159876 isoform X5
>gi|1036992310|ref|XP_017148974.1| PREDICTED: KN motif and ankyrin repeat domain-containing pro
>gi|1036992362|ref|XP_017148978.1| PREDICTED: KN motif and ankyrin repeat domain-containing pro
>gi|1037017560|ref|XP_017150507.1| PREDICTED: alkyldihydroxyacetonephosphate synthase [Drosophi
>gi|1037017578|ref|XP_017150508.1| PREDICTED: alkyldihydroxyacetonephosphate synthase [Drosophi
>gi|1037017596|ref|XP_017150509.1| PREDICTED: alkyldihydroxyacetonephosphate synthase [Drosophi
>gi|1037017612|ref|XP_017150510.1| PREDICTED: alkyldihydroxyacetonephosphate synthase [Drosophi
>gi|1037017630|ref|XP_017150511.1| PREDICTED: uncharacterized protein LOC108160801 [Drosophila
>gi|1036997211|ref|XP_017149281.1| PREDICTED: protein lap1 isoform X1 [Drosophila miranda]
>gi|1036997228|ref|XP_017149282.1| PREDICTED: protein lap1 isoform X2 [Drosophila miranda]
>gi|1036943654|ref|XP_017146045.1| PREDICTED: fas apoptotic inhibitory molecule 1 [Drosophila r
>gi|1036992708|ref|XP_017149001.1| PREDICTED: atypical protein kinase C isoform X2 [Drosophila
>gi|1036992728|ref|XP_017149002.1| PREDICTED: atypical protein kinase C isoform X2 [Drosophila
>gi|1036992690|ref|XP_017149000.1| PREDICTED: atypical protein kinase C isoform X1 [Drosophila
>gi|1036992746|ref|XP_017149003.1| PREDICTED: atypical protein kinase C isoform X3 [Drosophila
>gi|1036992764|ref|XP_017149004.1| PREDICTED: atypical protein kinase C isoform X4 [Drosophila
>gi|1036995388|ref|XP_017149168.1| PREDICTED: uncharacterized protein LOC108159982 isoform X1
>gi|1036995406|ref|XP_017149169.1| PREDICTED: uncharacterized protein LOC108159982 isoform X1
>gi|1036995422|ref|XP_017149170.1| PREDICTED: caskin-2 isoform X2 [Drosophila miranda]
>gi|1036989967|ref|XP_017148839.1| PREDICTED: uncharacterized protein LOC108159786 [Drosophila
>gi|1036989985|ref|XP_017148841.1| PREDICTED: neo-calmodulin [Drosophila miranda]
>gi|1036990003|ref|XP_017148842.1| PREDICTED: uncharacterized protein LOC108159790 [Drosophila
>gi|1036937873|ref|XP_017145687.1| PREDICTED: uncharacterized protein LOC108158052 [Drosophila
>gi|1036944187|ref|XP_017146076.1| PREDICTED: uncharacterized protein LOC108158348 [Drosophila
>gi|1036982919|ref|XP_017148406.1| PREDICTED: LOW QUALITY PROTEIN: mitochondrial-processing pep
>gi|1036982863|ref|XP_017148405.1| PREDICTED: ribonuclease 3 [Drosophila miranda]
>gi|1036982948|ref|XP_017148408.1| PREDICTED: mitochondrial-processing peptidase subunit alpha-
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>gi|1036982934|ref|XP_017148407.1| PREDICTED: ribonuclease 3-like [Drosophila miranda]
>gi|1037014525|ref|XP_017150324.1| PREDICTED: ribonuclease H1 isoform X1 [Drosophila miranda]
>gi|1037014539|ref|XP_017150325.1| PREDICTED: ribonuclease H1 isoform X2 [Drosophila miranda]
>gi|1036993246|ref|XP_017149034.1| PREDICTED: phosphatidylinositol-glycan biosynthesis class X
>gi|1036993228|ref|XP_017149033.1| PREDICTED: putative inorganic phosphate cotransporter [Drosc
>gi|1036993210|ref|XP_017149032.1| PREDICTED: probable lysine-specific demethylase 4A [Drosophi
>gi|1037000087|ref|XP_017149453.1| PREDICTED: odorant receptor 43b [Drosophila miranda]
>gi|1037000069|ref|XP_017149452.1| PREDICTED: cullin homolog 1 [Drosophila miranda]
>gi|1037012751|ref|XP_017150215.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1037007536|ref|XP_017149906.1| PREDICTED: uncharacterized protein LOC108160418 [Drosophila
>gi|1037007468|ref|XP_017149902.1| PREDICTED: uncharacterized protein LOC108160417 isoform X1
>gi|1037007486|ref|XP_017149903.1| PREDICTED: uncharacterized protein LOC108160417 isoform X1
>gi|1037007500|ref|XP_017149904.1| PREDICTED: uncharacterized protein LOC108160417 isoform X1
>gi|1037007518|ref|XP_017149905.1| PREDICTED: uncharacterized protein LOC108160417 isoform X2
>gi|1037004534|ref|XP_017149712.1| PREDICTED: diacylglycerol kinase 1 isoform X1 [Drosophila mi
>gi|1037004552|ref|XP_017149713.1| PREDICTED: diacylglycerol kinase 1 isoform X2 [Drosophila mi
>gi|1037004568|ref|XP_017149714.1| PREDICTED: diacylglycerol kinase 1 isoform X3 [Drosophila mi
>gi|1037004582|ref|XP_017149715.1| PREDICTED: diacylglycerol kinase 1 isoform X4 [Drosophila mi
>gi|1037004598|ref|XP_017149716.1| PREDICTED: diacylglycerol kinase 1 isoform X5 [Drosophila mi
>gi|1037004642|ref|XP_017149720.1| PREDICTED: ubiquitin-like-conjugating enzyme ATG10 [Drosophi
>gi|1037004658|ref|XP_017149721.1| PREDICTED: ubiquitin-like-conjugating enzyme ATG10 [Drosophi
>gi|1037004628|ref|XP_017149719.1| PREDICTED: tRNA (adenine(37)-N6)-methyltransferase isoform X
>gi|1037004616|ref|XP_017149718.1| PREDICTED: tRNA (adenine(37)-N6)-methyltransferase isoform X
>gi|1037008115|ref|XP_017149942.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylgl
>gi|1037016820|ref|XP_017150458.1| PREDICTED: threonine-rich protein [Drosophila miranda]
>gi|1037016798|ref|XP_017150457.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosoph
>gi|1036942970|ref|XP_017146002.1| PREDICTED: uncharacterized protein LOC108158277 [Drosophila
>gi|1037007253|ref|XP_017149890.1| PREDICTED: uncharacterized protein LOC108160408 [Drosophila
>gi|1037007269|ref|XP_017149891.1| PREDICTED: uncharacterized protein LOC108160408 [Drosophila
>gi|1037007287|ref|XP_017149892.1| PREDICTED: persulfide dioxygenase ETHE1, mitochondrial [Dros
>gi|1037007194|ref|XP_017149887.1| PREDICTED: LOW QUALITY PROTEIN: persulfide dioxygenase ETHE
>gi|1037007229|ref|XP_017149889.1| PREDICTED: superoxide dismutase [Cu-Zn], chloroplastic isofc
>gi|1037007212|ref|XP_017149888.1| PREDICTED: superoxide dismutase [Cu-Zn], chloroplastic isofc
>gi|1036942904|ref|XP_017145998.1| PREDICTED: CD63 antigen [Drosophila miranda]
>gi|1036943282|ref|XP_017146022.1| PREDICTED: uncharacterized protein LOC108158294 isoform X1
>gi|1036943298|ref|XP_017146023.1| PREDICTED: uncharacterized protein LOC108158294 isoform X2
>gi|1036943316|ref|XP_017146024.1| PREDICTED: uncharacterized protein LOC108158294 isoform X3
>gi|1037016161|ref|XP_017150418.1| PREDICTED: translocon-associated protein subunit delta [Dros
>gi|1036982184|ref|XP_017148363.1| PREDICTED: uncharacterized protein LOC108159515 [Drosophila
>gi|1036982119|ref|XP_017148359.1| PREDICTED: uncharacterized protein LOC108159511 [Drosophila
>gi|1036982136|ref|XP_017148360.1| PREDICTED: uncharacterized protein LOC108159511 [Drosophila
>gi|1036982200|ref|XP_017148364.1| PREDICTED: LOW QUALITY PROTEIN: putative gustatory receptor
>gi|1036982168|ref|XP_017148362.1| PREDICTED: probable N-acetyltransferase san [Drosophila mira
>gi|1036982150|ref|XP_017148361.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036982087|ref|XP_017148358.1| PREDICTED: uncharacterized protein LOC108159510 [Drosophila
>gi|1036980167|ref|XP_017148239.1| PREDICTED: probable phospholipid-transporting ATPase IA iso
>gi|1036980149|ref|XP_017148238.1| PREDICTED: probable phospholipid-transporting ATPase IA iso
>gi|1036980101|ref|XP_017148236.1| PREDICTED: probable phospholipid-transporting ATPase IA iso
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>gi|1036980086|ref|XP_017148234.1| PREDICTED: probable phospholipid-transporting ATPase IA iso
>gi|1036980201|ref|XP_017148242.1| PREDICTED: trypsin alpha-3 [Drosophila miranda]
>gi|1036980133|ref|XP_017148237.1| PREDICTED: probable phospholipid-transporting ATPase IA iso
>gi|1036980185|ref|XP_017148240.1| PREDICTED: probable phospholipid-transporting ATPase IA iso
>gi|1036980213|ref|XP_017148243.1| PREDICTED: putative odorant-binding protein A5 [Drosophila r
>gi|1036967681|ref|XP_017147471.1| PREDICTED: N-acetylglucosamine-6-sulfatase-like [Drosophila
>gi|1036967731|ref|XP_017147474.1| PREDICTED: immune-induced peptides [Drosophila miranda]
>gi|1036967647|ref|XP_017147469.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036967633|ref|XP_017147466.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036967697|ref|XP_017147472.1| PREDICTED: aminoacyl tRNA synthase complex-interacting mult
>gi|1036967715|ref|XP_017147473.1| PREDICTED: protein FAM8A1 [Drosophila miranda]
>gi|1036967665|ref|XP_017147470.1| PREDICTED: stromal membrane-associated protein 2 [Drosophila
>gi|1036984805|ref|XP_017148525.1| PREDICTED: serendipity locus protein alpha [Drosophila mirar
>gi|1036984825|ref|XP_017148526.1| PREDICTED: uncharacterized protein LOC108159587 isoform X1
>gi|1036984842|ref|XP_017148527.1| PREDICTED: uncharacterized protein LOC108159587 isoform X2
>gi|1036984769|ref|XP_017148522.1| PREDICTED: protein lines [Drosophila miranda]
>gi|1036984787|ref|XP_017148523.1| PREDICTED: protein lines [Drosophila miranda]
>gi|1036964189|ref|XP_017147270.1| PREDICTED: glucose-6-phosphate isomerase [Drosophila miranda
>gi|1036964172|ref|XP_017147269.1| PREDICTED: WD repeat-containing protein 43 [Drosophila mirar
>gi|1036964229|ref|XP_017147272.1| PREDICTED: uncharacterized protein LOC108158956 [Drosophila
>gi|1036964211|ref|XP_017147271.1| PREDICTED: T-complex protein 1 subunit theta [Drosophila mi
>gi|1036964276|ref|XP_017147275.1| PREDICTED: guanine nucleotide-binding protein subunit gamma-
>gi|1036964297|ref|XP_017147276.1| PREDICTED: guanine nucleotide-binding protein subunit gamma-
>gi|1036964258|ref|XP_017147274.1| PREDICTED: MRG/MORF4L-binding protein [Drosophila miranda]
>gi|1037005860|ref|XP_017149798.1| PREDICTED: trichohyalin isoform X14 [Drosophila miranda]
>gi|1037005631|ref|XP_017149781.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1037005684|ref|XP_017149785.1| PREDICTED: trichohyalin isoform X3 [Drosophila miranda]
>gi|1037005744|ref|XP_017149789.1| PREDICTED: myosin-9 isoform X7 [Drosophila miranda]
>gi|1037005697|ref|XP_017149786.1| PREDICTED: plectin isoform X4 [Drosophila miranda]
>gi|1037005712|ref|XP_017149787.1| PREDICTED: plectin isoform X5 [Drosophila miranda]
>gi|1037005760|ref|XP_017149791.1| PREDICTED: myosin-9 isoform X8 [Drosophila miranda]
>gi|1037005726|ref|XP_017149788.1| PREDICTED: myosin-9 isoform X6 [Drosophila miranda]
>gi|1037005774|ref|XP_017149792.1| PREDICTED: ELKS/Rab6-interacting/CAST family member 1 isofor
>gi|1037005807|ref|XP_017149794.1| PREDICTED: rootletin isoform X11 [Drosophila miranda]
>gi|1037005649|ref|XP_017149782.1| PREDICTED: trichohyalin isoform X2 [Drosophila miranda]
>gi|1037005669|ref|XP_017149783.1| PREDICTED: trichohyalin isoform X2 [Drosophila miranda]
>gi|1037005825|ref|XP_017149795.1| PREDICTED: plectin isoform X12 [Drosophila miranda]
>gi|1037005842|ref|XP_017149797.1| PREDICTED: plectin isoform X13 [Drosophila miranda]
>gi|1037005791|ref|XP_017149793.1| PREDICTED: trichohyalin isoform X10 [Drosophila miranda]
>gi|1037008097|ref|XP_017149941.1| PREDICTED: protein PTCD3 homolog, mitochondrial [Drosophila
>gi|1036999074|ref|XP_017149398.1| PREDICTED: nucleoporin GLE1 [Drosophila miranda]
>gi|1036999091|ref|XP_017149399.1| PREDICTED: zinc finger CCCH domain-containing protein 15 hor
>gi|1036989080|ref|XP_017148784.1| PREDICTED: protein patched [Drosophila miranda]
>gi|1036995517|ref|XP_017149176.1| PREDICTED: long-chain-fatty-acid--CoA ligase 4 isoform X1 [[
>gi|1036995535|ref|XP_017149177.1| PREDICTED: long-chain-fatty-acid--CoA ligase 4 isoform X1 [[
>gi|1036995579|ref|XP_017149179.1| PREDICTED: long-chain-fatty-acid--CoA ligase 4 isoform X3 [[
>gi|1036995553|ref|XP_017149178.1| PREDICTED: long-chain-fatty-acid--CoA ligase 4 isoform X2 [[
>gi|1036995597|ref|XP_017149180.1| PREDICTED: uncharacterized protein LOC108159986 [Drosophila
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>gi|1036941044|ref|XP_017145882.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1037020313|ref|XP_017150685.1| PREDICTED: serine protease 29 isoform X2 [Drosophila miranda
>gi|1037020295|ref|XP_017150684.1| PREDICTED: acrosin isoform X1 [Drosophila miranda]
>gi|1037021304|ref|XP_017150748.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1037021322|ref|XP_017150749.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1037019859|ref|XP_017150657.1| PREDICTED: CTD nuclear envelope phosphatase 1 [Drosophila m
>gi|1037019873|ref|XP_017150658.1| PREDICTED: sodium/potassium/calcium exchanger 6, mitochondri
>gi|1036943494|ref|XP_017146034.1| PREDICTED: sodium/potassium/calcium exchanger 6, mitochondri
>gi|1037021266|ref|XP_017150746.1| PREDICTED: peptidoglycan-recognition protein SC1a/b [Drosoph
>gi|1037021233|ref|XP_017150744.1| PREDICTED: sodium/potassium/calcium exchanger 6, mitochondri
>gi|1037021249|ref|XP_017150745.1| PREDICTED: sodium/potassium/calcium exchanger 6, mitochondri
>gi|1036942002|ref|XP_017145941.1| PREDICTED: sodium/potassium/calcium exchanger 6, mitochondri
>gi|1036942172|ref|XP_017145952.1| PREDICTED: peptidoglycan-recognition protein SC1a/b [Drosoph
>gi|1036950048|ref|XP_017146434.1| PREDICTED: peptidoglycan-recognition protein SC2 [Drosophila
>gi|1036949962|ref|XP_017146429.1| PREDICTED: tetraspanning orphan receptor isoform X3 [Drosoph
>gi|1036949946|ref|XP_017146428.1| PREDICTED: tetraspanning orphan receptor isoform X2 [Drosoph
>gi|1036949932|ref|XP_017146427.1| PREDICTED: uncharacterized protein LOC108158535 isoform X1
>gi|1036949994|ref|XP_017146431.1| PREDICTED: kelch-like protein 9 isoform X2 [Drosophila mira
>gi|1036949976|ref|XP_017146430.1| PREDICTED: kelch-like protein 9 isoform X1 [Drosophila mira
>gi|1036950030|ref|XP_017146433.1| PREDICTED: uncharacterized protein LOC108158536 isoform X4
>gi|1036950012|ref|XP_017146432.1| PREDICTED: uncharacterized protein LOC108158536 isoform X3
>gi|1036937891|ref|XP_017145688.1| PREDICTED: uncharacterized protein LOC108158053 [Drosophila
>gi|1036949864|ref|XP_017146423.1| PREDICTED: myotubularin-related protein 6 isoform X2 [Droso
>gi|1036949831|ref|XP_017146419.1| PREDICTED: myotubularin-related protein 6 isoform X1 [Droso
>gi|1036949846|ref|XP_017146422.1| PREDICTED: myotubularin-related protein 6 isoform X1 [Droso
>gi|1036949898|ref|XP_017146425.1| PREDICTED: translation initiation factor eIF-2B subunit del
>gi|1036949880|ref|XP_017146424.1| PREDICTED: translation initiation factor eIF-2B subunit del
>gi|1036949916|ref|XP_017146426.1| PREDICTED: translation initiation factor eIF-2B subunit del
>gi|1036950066|ref|XP_017146436.1| PREDICTED: uncharacterized protein LOC108158539 [Drosophila
>gi|1036949814|ref|XP_017146418.1| PREDICTED: little elongation complex subunit 1 [Drosophila
>gi|1036949778|ref|XP_017146416.1| PREDICTED: golgin subfamily A member 4 isoform X1 [Drosophil
>gi|1036949796|ref|XP_017146417.1| PREDICTED: golgin subfamily A member 4 isoform X2 [Drosophil
>gi|1037010584|ref|XP_017150087.1| PREDICTED: ATPase inhibitor mai-2, mitochondrial [Drosophila
>gi|1037010549|ref|XP_017150085.1| PREDICTED: major royal jelly protein 1 [Drosophila miranda]
>gi|1037010562|ref|XP_017150086.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1037014595|ref|XP_017150329.1| PREDICTED: 40S ribosomal protein S18 [Drosophila miranda]
>gi|1037012594|ref|XP_017150205.1| PREDICTED: DNA replication inhibitor plutonium-like [Drosoph
>gi|1037012630|ref|XP_017150207.1| PREDICTED: uncharacterized protein LOC108160608 isoform X2
>gi|1037012612|ref|XP_017150206.1| PREDICTED: uncharacterized protein LOC108160608 isoform X1
>gi|1036996452|ref|XP_017149233.1| PREDICTED: proliferating cell nuclear antigen [Drosophila m
>gi|1036996432|ref|XP_017149232.1| PREDICTED: hormone-sensitive lipase-like [Drosophila miranda
>gi|1036937909|ref|XP_017145689.1| PREDICTED: probable cytochrome P450 4p3 [Drosophila miranda]
>gi|1036937923|ref|XP_017145692.1| PREDICTED: uncharacterized protein LOC108158055, partial [D
>gi|1036937939|ref|XP_017145693.1| PREDICTED: probable cytochrome P450 6a23 [Drosophila miranda
>gi|1036937955|ref|XP_017145694.1| PREDICTED: probable cytochrome P450 6a23, partial [Drosophil
>gi|1036937970|ref|XP_017145695.1| PREDICTED: kinesin-like protein subito, partial [Drosophila
>gi|1036977133|ref|XP_017148047.1| PREDICTED: protein germ cell-less [Drosophila miranda]
>gi|1036977083|ref|XP_017148044.1| PREDICTED: protein EFR3 homolog cmp44E isoform X1 [Drosophil
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>gi|1036977115|ref|XP_017148046.1| PREDICTED: protein EFR3 homolog cmp44E isoform X3 [Drosophil
>gi|1036977100|ref|XP_017148045.1| PREDICTED: protein EFR3 homolog cmp44E isoform X2 [Drosophil
>gi|1037008545|ref|XP_017149965.1| PREDICTED: uncharacterized protein LOC108160462 [Drosophila
>gi|1036942579|ref|XP_017145977.1| PREDICTED: uncharacterized protein LOC108158252 [Drosophila
>gi|1037002419|ref|XP_017149598.1| PREDICTED: nephrin isoform X3 [Drosophila miranda]
>gi|1037002384|ref|XP_017149596.1| PREDICTED: nephrin isoform X1 [Drosophila miranda]
>gi|1037002401|ref|XP_017149597.1| PREDICTED: nephrin isoform X2 [Drosophila miranda]
>gi|1037002437|ref|XP_017149599.1| PREDICTED: uncharacterized protein LOC108160248 [Drosophila
>gi|1036988300|ref|XP_017148737.1| PREDICTED: ryanodine receptor isoform X3 [Drosophila miranda
>gi|1036988470|ref|XP_017148745.1| PREDICTED: ryanodine receptor isoform X11 [Drosophila mirand
>gi|1036988375|ref|XP_017148741.1| PREDICTED: ryanodine receptor isoform X7 [Drosophila miranda
>gi|1036988341|ref|XP_017148739.1| PREDICTED: ryanodine receptor isoform X5 [Drosophila miranda
>gi|1036988359|ref|XP_017148740.1| PREDICTED: ryanodine receptor isoform X6 [Drosophila miranda
>gi|1036988429|ref|XP_017148744.1| PREDICTED: ryanodine receptor isoform X10 [Drosophila mirand
>gi|1036988392|ref|XP_017148742.1| PREDICTED: ryanodine receptor isoform X8 [Drosophila miranda
>gi|1036988266|ref|XP_017148735.1| PREDICTED: ryanodine receptor isoform X1 [Drosophila miranda
>gi|1036988317|ref|XP_017148738.1| PREDICTED: ryanodine receptor isoform X4 [Drosophila miranda
>gi|1036988280|ref|XP_017148736.1| PREDICTED: ryanodine receptor isoform X2 [Drosophila miranda
>gi|1036988409|ref|XP_017148743.1| PREDICTED: ryanodine receptor isoform X9 [Drosophila miranda
>gi|1036988487|ref|XP_017148746.1| PREDICTED: SCF E3 ubiquitin ligase complex F-box protein po
>gi|1037013255|ref|XP_017150247.1| PREDICTED: dynactin subunit 2 [Drosophila miranda]
>gi|1036999387|ref|XP_017149414.1| PREDICTED: protein transport protein Sec31A [Drosophila mir
>gi|1036999405|ref|XP_017149415.1| PREDICTED: uncharacterized protein LOC108160131 [Drosophila
>gi|1037019777|ref|XP_017150651.1| PREDICTED: uncharacterized protein LOC108160893 [Drosophila
>gi|1037013399|ref|XP_017150255.1| PREDICTED: protein Wnt-2 isoform X1 [Drosophila miranda]
>gi|1037013416|ref|XP_017150256.1| PREDICTED: protein Wnt-2 isoform X2 [Drosophila miranda]
>gi|1036956368|ref|XP_017146812.1| PREDICTED: uncharacterized protein LOC108158726 isoform X4
>gi|1036956309|ref|XP_017146809.1| PREDICTED: uncharacterized protein LOC108158726 isoform X2
>gi|1036956292|ref|XP_017146808.1| PREDICTED: uncharacterized protein LOC108158726 isoform X1
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>gi|1036956401|ref|XP_017146815.1| PREDICTED: uncharacterized protein LOC108158726 isoform X6
>gi|1036956385|ref|XP_017146814.1| PREDICTED: uncharacterized protein LOC108158726 isoform X5
>gi|1036956328|ref|XP_017146810.1| PREDICTED: uncharacterized protein LOC108158726 isoform X3
>gi|1036956350|ref|XP_017146811.1| PREDICTED: uncharacterized protein LOC108158726 isoform X3
>gi|1036956478|ref|XP_017146820.1| PREDICTED: PTB domain-containing adapter protein ced-6 isof
>gi|1036956514|ref|XP_017146822.1| PREDICTED: PTB domain-containing adapter protein ced-6 isof
>gi|1036956496|ref|XP_017146821.1| PREDICTED: PTB domain-containing adapter protein ced-6 isof
>gi|1036956530|ref|XP_017146823.1| PREDICTED: pyruvate dehydrogenase (acetyl-transferring) kin
>gi|1036956542|ref|XP_017146824.1| PREDICTED: pyruvate dehydrogenase (acetyl-transferring) kin
>gi|1036956461|ref|XP_017146819.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 38 [Drosop
>gi|1036940323|ref|XP_017145843.1| PREDICTED: transmembrane protein 18 isoform X1 [Drosophila
>gi|1036940338|ref|XP_017145844.1| PREDICTED: transmembrane protein 18 isoform X2 [Drosophila
>gi|1036937988|ref|XP_017145696.1| PREDICTED: uncharacterized protein LOC108158059 [Drosophila
>gi|1037000567|ref|XP_017149480.1| PREDICTED: dystrobrevin beta isoform X1 [Drosophila miranda]
>gi|1037000584|ref|XP_017149481.1| PREDICTED: dystrobrevin beta isoform X2 [Drosophila miranda]
>gi|1037000601|ref|XP_017149483.1| PREDICTED: dystrobrevin beta isoform X3 [Drosophila miranda]
>gi|1037005878|ref|XP_017149799.1| PREDICTED: lachesin [Drosophila miranda]
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>gi|1036943566|ref|XP_017146038.1| PREDICTED: uncharacterized protein LOC108158310 [Drosophila
>gi|1036988027|ref|XP_017148720.1| PREDICTED: CWF19-like protein 1 homolog isoform X2 [Drosophi
>gi|1036988009|ref|XP_017148719.1| PREDICTED: CWF19-like protein 1 homolog isoform X1 [Drosophi
>gi|1036988059|ref|XP_017148723.1| PREDICTED: farnesyl pyrophosphate synthase [Drosophila mirar
>gi|1036988045|ref|XP_017148721.1| PREDICTED: putative uncharacterized protein DDB_G0286901 [Dr
>gi|1036987991|ref|XP_017148718.1| PREDICTED: uncharacterized protein LOC108159711 isoform X3
>gi|1036987973|ref|XP_017148717.1| PREDICTED: uncharacterized protein LOC108159711 isoform X2
>gi|1036987953|ref|XP_017148716.1| PREDICTED: uncharacterized protein LOC108159711 isoform X1
>gi|1036988083|ref|XP_017148724.1| PREDICTED: uncharacterized protein LOC108159715 [Drosophila
>gi|1036946543|ref|XP_017146215.1| PREDICTED: odorant receptor 47b [Drosophila miranda]
>gi|1037009345|ref|XP_017150014.1| PREDICTED: uncharacterized protein LOC108160490 [Drosophila
>gi|1037009359|ref|XP_017150016.1| PREDICTED: trypsin delta/gamma-like [Drosophila miranda]
>gi|1036938006|ref|XP_017145697.1| PREDICTED: transmembrane protease serine 9-like [Drosophila
>gi|1037021656|ref|XP_017150769.1| PREDICTED: trypsin theta [Drosophila miranda]
>gi|1037019634|ref|XP_017150641.1| PREDICTED: trypsin alpha [Drosophila miranda]
>gi|1037017822|ref|XP_017150524.1| PREDICTED: trypsin epsilon [Drosophila miranda]
>gi|1037017832|ref|XP_017150525.1| PREDICTED: LOW QUALITY PROTEIN: trypsin alpha-like [Drosophi
>gi|1037016926|ref|XP_017150465.1| PREDICTED: trypsin delta/gamma [Drosophila miranda]
>gi|1037016955|ref|XP_017150467.1| PREDICTED: coiled-coil domain-containing protein 103 [Drosop
>gi|1037016940|ref|XP_017150466.1| PREDICTED: trypsin iota-like [Drosophila miranda]
>gi|1036940841|ref|XP_017145876.1| PREDICTED: death-associated protein 1 [Drosophila miranda]
>gi|1036959264|ref|XP_017146982.1| PREDICTED: uroporphyrinogen decarboxylase [Drosophila mirano
>gi|1036959413|ref|XP_017146992.1| PREDICTED: transmembrane protein 234 homolog [Drosophila mi
>gi|1036959450|ref|XP_017146993.1| PREDICTED: 60S ribosomal protein L31 [Drosophila miranda]
>gi|1036959176|ref|XP_017146977.1| PREDICTED: uncharacterized protein LOC108158802 [Drosophila
>gi|1036959222|ref|XP_017146981.1| PREDICTED: putative N(4)-(beta-N-acetylglucosaminyl)-L-aspar
>gi|1036959208|ref|XP_017146980.1| PREDICTED: uncharacterized protein LOC108158804 [Drosophila
>gi|1036959366|ref|XP_017146988.1| PREDICTED: RWD domain-containing protein 2A [Drosophila mira
>gi|1036959193|ref|XP_017146978.1| PREDICTED: neuropeptide Y receptor type 1 [Drosophila mirano
>gi|1036959350|ref|XP_017146987.1| PREDICTED: alpha-tocopherol transfer protein [Drosophila mi
>gi|1036959275|ref|XP_017146983.1| PREDICTED: uncharacterized protein LOC108158807 [Drosophila
>gi|1036959290|ref|XP_017146984.1| PREDICTED: uncharacterized protein LOC108158807 [Drosophila
>gi|1036959396|ref|XP_017146991.1| PREDICTED: glutathione S-transferase theta-1 [Drosophila mi
>gi|1036959382|ref|XP_017146989.1| PREDICTED: glutathione S-transferase theta-1-like [Drosophi
>gi|1036959302|ref|XP_017146985.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophi
>gi|1036959330|ref|XP_017146986.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophi
>gi|1036975381|ref|XP_017147938.1| PREDICTED: matrix metalloproteinase-2 isoform X1 [Drosophila
>gi|1036975419|ref|XP_017147940.1| PREDICTED: matrix metalloproteinase-2 isoform X3 [Drosophila
>gi|1036975397|ref|XP_017147939.1| PREDICTED: matrix metalloproteinase-2 isoform X2 [Drosophila
>gi|1036975433|ref|XP_017147941.1| PREDICTED: uncharacterized protein LOC108159288 [Drosophila
>gi|1036975346|ref|XP_017147936.1| PREDICTED: ubiquitin-like modifier-activating enzyme 1 [Dros
>gi|1036975364|ref|XP_017147937.1| PREDICTED: ubiquitin-like modifier-activating enzyme 1 [Dros
>gi|1036938022|ref|XP_017145699.1| PREDICTED: uncharacterized protein LOC108158061, partial [Dr
>gi|1036992002|ref|XP_017148956.1| PREDICTED: uncharacterized protein LOC108159866 [Drosophila
>gi|1036992016|ref|XP_017148957.1| PREDICTED: uncharacterized protein LOC108159867 [Drosophila
>gi|1036991968|ref|XP_017148954.1| PREDICTED: serine-threonine kinase receptor-associated prote
>gi|1036991986|ref|XP_017148955.1| PREDICTED: cyclin-dependent kinase inhibitor 1 [Drosophila n
>gi|1036991946|ref|XP_017148953.1| PREDICTED: serine/threonine-protein kinase DDB_G0283821 [Dro
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>gi|1036938039|ref|XP_017145700.1| PREDICTED: uncharacterized protein LOC108158062 [Drosophila
>gi|1037018311|ref|XP_017150556.1| PREDICTED: transient-receptor-potential-like protein isofor
>gi|1037018331|ref|XP_017150557.1| PREDICTED: transient-receptor-potential-like protein isofor
>gi|1037018347|ref|XP_017150558.1| PREDICTED: transient-receptor-potential-like protein isofor
>gi|1037018392|ref|XP_017150561.1| PREDICTED: uncharacterized protein LOC108160833 [Drosophila
>gi|1037018361|ref|XP_017150559.1| PREDICTED: transient-receptor-potential-like protein isofor
>gi|1037018379|ref|XP_017150560.1| PREDICTED: transient-receptor-potential-like protein isofor
>gi|1037009529|ref|XP_017150027.1| PREDICTED: uncharacterized protein LOC108160500 isoform X2
>gi|1037009511|ref|XP_017150026.1| PREDICTED: uncharacterized protein LOC108160500 isoform X1
>gi|1037009543|ref|XP_017150028.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036938056|ref|XP_017145701.1| PREDICTED: uncharacterized protein LOC108158063 [Drosophila
>gi|1036938070|ref|XP_017145702.1| PREDICTED: uncharacterized protein LOC108158064, partial [D
>gi|1037007883|ref|XP_017149926.1| PREDICTED: zinc finger protein 573 isoform X1 [Drosophila m
>gi|1037007932|ref|XP_017149930.1| PREDICTED: zinc finger protein 573 isoform X4 [Drosophila m
>gi|1037007914|ref|XP_017149928.1| PREDICTED: zinc finger protein 573 isoform X3 [Drosophila m
>gi|1037007898|ref|XP_017149927.1| PREDICTED: zinc finger protein 573 isoform X2 [Drosophila m
>gi|1037007950|ref|XP_017149931.1| PREDICTED: uncharacterized protein LOC108160434 isoform X5
>gi|1036982236|ref|XP_017148367.1| PREDICTED: uncharacterized protein LOC108159518 [Drosophila
>gi|1036982218|ref|XP_017148365.1| PREDICTED: eukaryotic translation initiation factor 4 gamma
>gi|1036997605|ref|XP_017149304.1| PREDICTED: uncharacterized protein LOC108160066 [Drosophila
>gi|1036997585|ref|XP_017149303.1| PREDICTED: zinc finger protein GLIS2 homolog isoform X2 [Dr
>gi|1036997565|ref|XP_017149302.1| PREDICTED: zinc finger protein GLIS2 homolog isoform X1 [Dr
>gi|1036997549|ref|XP_017149301.1| PREDICTED: uncharacterized protein LOC108160064 [Drosophila
>gi|1037009825|ref|XP_017150040.1| PREDICTED: DCN1-like protein 3 [Drosophila miranda]
>gi|1037009845|ref|XP_017150042.1| PREDICTED: DCN1-like protein 3 [Drosophila miranda]
>gi|1037009863|ref|XP_017150043.1| PREDICTED: DCN1-like protein 3 [Drosophila miranda]
>gi|1036970233|ref|XP_017147622.1| PREDICTED: Usher syndrome type-1G protein homolog [Drosophil
>gi|1036970323|ref|XP_017147627.1| PREDICTED: S-phase kinase-associated protein 1 [Drosophila
>gi|1036970215|ref|XP_017147620.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosop
>gi|1036970287|ref|XP_017147625.1| PREDICTED: protein OSCP1 isoform X1 [Drosophila miranda]
>gi|1036970304|ref|XP_017147626.1| PREDICTED: protein OSCP1 isoform X2 [Drosophila miranda]
>gi|1036970269|ref|XP_017147624.1| PREDICTED: probable trans-2-enoyl-CoA reductase, mitochondri
>gi|1036970253|ref|XP_017147623.1| PREDICTED: nedd8-activating enzyme E1 catalytic subunit [Dr
>gi|1036970195|ref|XP_017147619.1| PREDICTED: LOW QUALITY PROTEIN: rac GTPase-activating prote
>gi|1036941717|ref|XP_017145920.1| PREDICTED: sortilin-related receptor [Drosophila miranda]
>gi|1037012883|ref|XP_017150224.1| PREDICTED: probable enoyl-CoA hydratase, mitochondrial [Dros
>gi|1037012869|ref|XP_017150223.1| PREDICTED: uncharacterized protein LOC108160618 [Drosophila
>gi|1037016850|ref|XP_017150460.1| PREDICTED: synaptogyrin-1 [Drosophila miranda]
>gi|1037016864|ref|XP_017150461.1| PREDICTED: uncharacterized protein LOC108160772 isoform X1
>gi|1037016882|ref|XP_017150462.1| PREDICTED: uncharacterized protein LOC108160772 isoform X2
>gi|1036944058|ref|XP_017146067.1| PREDICTED: cathepsin L1 [Drosophila miranda]
>gi|1037020237|ref|XP_017150680.1| PREDICTED: zingipain-1 [Drosophila miranda]
>gi|1037011251|ref|XP_017150123.1| PREDICTED: uncharacterized protein LOC108160560 [Drosophila
>gi|1037011269|ref|XP_017150124.1| PREDICTED: uncharacterized protein LOC108160560 [Drosophila
>gi|1037011287|ref|XP_017150125.1| PREDICTED: uncharacterized protein LOC108160561 [Drosophila
>gi|1037002328|ref|XP_017149592.1| PREDICTED: pre-mRNA-splicing regulator female-lethal(2)D iso
>gi|1037002344|ref|XP_017149593.1| PREDICTED: pre-mRNA-splicing regulator female-lethal(2)D iso
>gi|1037014400|ref|XP_017150317.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
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>gi|1037014416|ref|XP_017150319.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1037014369|ref|XP_017150315.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1037014336|ref|XP_017150313.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1037014385|ref|XP_017150316.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1037014351|ref|XP_017150314.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1037014432|ref|XP_017150320.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1036938087|ref|XP_017145703.1| PREDICTED: uncharacterized protein LOC108158065 [Drosophila
>gi|1037019274|ref|XP_017150620.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1037021890|ref|XP_017150784.1| PREDICTED: actin cytoskeleton-regulatory complex protein PA
>gi|1037021905|ref|XP_017150785.1| PREDICTED: actin cytoskeleton-regulatory complex protein PA
>gi|1037005892|ref|XP_017149800.1| PREDICTED: uncharacterized protein LOC108160349 [Drosophila
>gi|1037005908|ref|XP_017149801.1| PREDICTED: axoneme-associated protein mst101(2)-like [Droso
>gi|1036942491|ref|XP_017145972.1| PREDICTED: parafibromin isoform X2 [Drosophila miranda]
>gi|1036942455|ref|XP_017145970.1| PREDICTED: parafibromin isoform X1 [Drosophila miranda]
>gi|1036942473|ref|XP_017145971.1| PREDICTED: parafibromin isoform X1 [Drosophila miranda]
>gi|1036942277|ref|XP_017145958.1| PREDICTED: uncharacterized protein LOC108158237 [Drosophila
>gi|1036940495|ref|XP_017145854.1| PREDICTED: sulfhydryl oxidase 1-like [Drosophila miranda]
>gi|1036940647|ref|XP_017145864.1| PREDICTED: probable palmitoyltransferase ZDHHC24 [Drosophila
>gi|1036940683|ref|XP_017145866.1| PREDICTED: vacuolar protein-sorting-associated protein 25 [
>gi|1036940355|ref|XP_017145845.1| PREDICTED: latrophilin Cirl isoform X1 [Drosophila miranda]
>gi|1036940373|ref|XP_017145846.1| PREDICTED: latrophilin Cirl isoform X2 [Drosophila miranda]
>gi|1036940392|ref|XP_017145847.1| PREDICTED: latrophilin Cirl isoform X2 [Drosophila miranda]
>gi|1036940409|ref|XP_017145848.1| PREDICTED: latrophilin Cirl isoform X3 [Drosophila miranda]
>gi|1036940528|ref|XP_017145856.1| PREDICTED: uncharacterized protein LOC108158147 [Drosophila
>gi|1036938103|ref|XP_017145704.1| PREDICTED: actin-binding protein anillin-like [Drosophila m
>gi|1036940544|ref|XP_017145857.1| PREDICTED: putative ATPase N2B [Drosophila miranda]
>gi|1036940593|ref|XP_017145860.1| PREDICTED: deoxyribose-phosphate aldolase [Drosophila miran
>gi|1036940665|ref|XP_017145865.1| PREDICTED: uncharacterized protein LOC108158154 [Drosophila
>gi|1036940513|ref|XP_017145855.1| PREDICTED: probable nucleoporin Nup54 [Drosophila miranda]
>gi|1036940772|ref|XP_017145872.1| PREDICTED: uncharacterized protein LOC108158160 [Drosophila
>gi|1036940754|ref|XP_017145871.1| PREDICTED: LOW QUALITY PROTEIN: bolA-like protein 2 [Drosop
>gi|1036940561|ref|XP_017145858.1| PREDICTED: LOW QUALITY PROTEIN: mevalonate kinase [Drosophil
>gi|1036940736|ref|XP_017145870.1| PREDICTED: transcription elongation factor SPT4 [Drosophila
>gi|1036940718|ref|XP_017145868.1| PREDICTED: transcription elongation factor SPT4 [Drosophila
>gi|1036940700|ref|XP_017145867.1| PREDICTED: uncharacterized protein LOC108158156 [Drosophila
>gi|1036940611|ref|XP_017145862.1| PREDICTED: transmembrane protein 183 isoform X1 [Drosophila
>gi|1036940627|ref|XP_017145863.1| PREDICTED: transmembrane protein 183 isoform X2 [Drosophila
>gi|1036940576|ref|XP_017145859.1| PREDICTED: LOW QUALITY PROTEIN: cytosol aminopeptidase [Dros
>gi|1036940427|ref|XP_017145849.1| PREDICTED: pre-mRNA-splicing factor SYF1 [Drosophila miranda
>gi|1036940443|ref|XP_017145850.1| PREDICTED: CDK5 and ABL1 enzyme substrate 1 isoform X1 [Dros
>gi|1036940461|ref|XP_017145851.1| PREDICTED: CDK5 and ABL1 enzyme substrate 1 isoform X2 [Dros
>gi|1036940479|ref|XP_017145852.1| PREDICTED: CDK5 and ABL1 enzyme substrate 1 isoform X3 [Dros
>gi|1037019795|ref|XP_017150652.1| PREDICTED: uncharacterized protein LOC108160894 [Drosophila
>gi|1037021039|ref|XP_017150733.1| PREDICTED: L-lactate dehydrogenase B chain [Drosophila miran
>gi|1036985190|ref|XP_017148547.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm6 [Drosoph
>gi|1036985175|ref|XP_017148546.1| PREDICTED: uncharacterized protein LOC108159603 [Drosophila
>gi|1036985159|ref|XP_017148545.1| PREDICTED: charged multivesicular body protein 4c isoform X2
>gi|1036985143|ref|XP_017148544.1| PREDICTED: charged multivesicular body protein 4c isoform X1
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>gi|1036985127|ref|XP_017148543.1| PREDICTED: pre-mRNA-splicing factor 38A [Drosophila miranda]
>gi|1036985092|ref|XP_017148541.1| PREDICTED: proton-coupled folate transporter [Drosophila mi
>gi|1036985110|ref|XP_017148542.1| PREDICTED: proton-coupled folate transporter-like [Drosophil
>gi|1036985074|ref|XP_017148540.1| PREDICTED: uncharacterized protein LOC108159597 [Drosophila
>gi|1037021336|ref|XP_017150750.1| PREDICTED: proton-coupled folate transporter isoform X1 [Dro
>gi|1037021354|ref|XP_017150752.1| PREDICTED: proton-coupled folate transporter isoform X2 [Dro
>gi|1036959807|ref|XP_017147010.1| PREDICTED: DNA repair protein RAD51 homolog 4 [Drosophila mi
>gi|1036959825|ref|XP_017147011.1| PREDICTED: uncharacterized protein LOC108158825 [Drosophila
>gi|1036959690|ref|XP_017147004.1| PREDICTED: dnaJ homolog subfamily C member 13 isoform X2 [Dr
>gi|1036959710|ref|XP_017147005.1| PREDICTED: dnaJ homolog subfamily C member 13 isoform X3 [Dr
>gi|1036959654|ref|XP_017147002.1| PREDICTED: dnaJ homolog subfamily C member 13 isoform X1 [Dr
>gi|1036959672|ref|XP_017147003.1| PREDICTED: dnaJ homolog subfamily C member 13 isoform X1 [Dr
>gi|1036959763|ref|XP_017147009.1| PREDICTED: ras-related protein Rab-32 isoform X3 [Drosophila
>gi|1036959741|ref|XP_017147008.1| PREDICTED: ras-related protein Rab-44 isoform X2 [Drosophila
>gi|1036959723|ref|XP_017147007.1| PREDICTED: ras and EF-hand domain-containing protein homolog
>gi|1036959843|ref|XP_017147012.1| PREDICTED: mitochondrial folate transporter/carrier [Drosoph
>gi|1037000882|ref|XP_017149502.1| PREDICTED: protein SMG8 isoform X1 [Drosophila miranda]
>gi|1037000896|ref|XP_017149504.1| PREDICTED: protein SMG8 isoform X2 [Drosophila miranda]
>gi|1037000979|ref|XP_017149505.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1037002165|ref|XP_017149582.1| PREDICTED: protein scabrous [Drosophila miranda]
>gi|1037020255|ref|XP_017150682.1| PREDICTED: probable cytochrome P450 9h1 [Drosophila miranda]
>gi|1037020277|ref|XP_017150683.1| PREDICTED: odorant receptor 49b [Drosophila miranda]
>gi|1037022252|ref|XP_017150806.1| PREDICTED: venom allergen 5.01 isoform X2 [Drosophila miran
>gi|1037022234|ref|XP_017150805.1| PREDICTED: venom allergen 3 isoform X1 [Drosophila miranda]
>gi|1037021614|ref|XP_017150766.1| PREDICTED: venom allergen 3 [Drosophila miranda]
>gi|1037021642|ref|XP_017150768.1| PREDICTED: uncharacterized protein LOC108160978 isoform X2
>gi|1037021626|ref|XP_017150767.1| PREDICTED: uncharacterized protein LOC108160978 isoform X1
>gi|1036968976|ref|XP_017147545.1| PREDICTED: uncharacterized protein LOC108159090 isoform X10
>gi|1036968850|ref|XP_017147537.1| PREDICTED: uncharacterized protein LOC108159090 isoform X3
>gi|1036968808|ref|XP_017147535.1| PREDICTED: uncharacterized protein LOC108159090 isoform X1
>gi|1036968939|ref|XP_017147542.1| PREDICTED: uncharacterized protein LOC108159090 isoform X8
>gi|1036968828|ref|XP_017147536.1| PREDICTED: protein shisa-5 isoform X2 [Drosophila miranda]
>gi|1036968921|ref|XP_017147541.1| PREDICTED: uncharacterized protein LOC108159090 isoform X7
>gi|1036968904|ref|XP_017147540.1| PREDICTED: uncharacterized protein LOC108159090 isoform X6
>gi|1036968956|ref|XP_017147544.1| PREDICTED: uncharacterized protein LOC108159090 isoform X9
>gi|1036968888|ref|XP_017147539.1| PREDICTED: uncharacterized protein LOC108159090 isoform X5
>gi|1036968870|ref|XP_017147538.1| PREDICTED: uncharacterized protein LOC108159090 isoform X4
>gi|1036968790|ref|XP_017147534.1| PREDICTED: transcription factor BTF3 homolog 4 [Drosophila
>gi|1036969016|ref|XP_017147547.1| PREDICTED: carcinine transporter isoform X2 [Drosophila mira
>gi|1036968996|ref|XP_017147546.1| PREDICTED: carcinine transporter isoform X1 [Drosophila mira
>gi|1036969157|ref|XP_017147556.1| PREDICTED: protein lifeguard 1 isoform X2 [Drosophila miran
>gi|1036969181|ref|XP_017147557.1| PREDICTED: protein lifeguard 1 isoform X3 [Drosophila miran
>gi|1036969139|ref|XP_017147555.1| PREDICTED: protein lifeguard 1 isoform X1 [Drosophila miran
>gi|1036969129|ref|XP_017147554.1| PREDICTED: protein lifeguard 1 isoform X2 [Drosophila miran
>gi|1036969094|ref|XP_017147551.1| PREDICTED: protein lifeguard 1 isoform X1 [Drosophila miran
>gi|1036969111|ref|XP_017147553.1| PREDICTED: protein lifeguard 1 isoform X1 [Drosophila miran
>gi|1036969034|ref|XP_017147548.1| PREDICTED: aspartate--tRNA ligase, cytoplasmic isoform X1 [
>gi|1036969054|ref|XP_017147549.1| PREDICTED: aspartate--tRNA ligase, cytoplasmic isoform X2 [
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>gi|1036969074|ref|XP_017147550.1| PREDICTED: protein vestigial [Drosophila miranda]
>gi|1036942694|ref|XP_017145985.1| PREDICTED: ERI1 exoribonuclease 2-like [Drosophila miranda]
>gi|1036938121|ref|XP_017145705.1| PREDICTED: proteasome subunit alpha type-4-like [Drosophila
>gi|1037017810|ref|XP_017150523.1| PREDICTED: uncharacterized protein LOC108160811 [Drosophila
>gi|1037014450|ref|XP_017150321.1| PREDICTED: 40S ribosomal protein S18 [Drosophila miranda]
>gi|1037012769|ref|XP_017150216.1| PREDICTED: DNA replication inhibitor plutonium [Drosophila m
>gi|1037012803|ref|XP_017150219.1| PREDICTED: uncharacterized protein LOC108160614 isoform X2
>gi|1037012785|ref|XP_017150218.1| PREDICTED: uncharacterized protein LOC108160614 isoform X1
>gi|1036996340|ref|XP_017149226.1| PREDICTED: proliferating cell nuclear antigen [Drosophila m
>gi|1036996290|ref|XP_017149223.1| PREDICTED: hormone-sensitive lipase isoform X1 [Drosophila m
>gi|1036996307|ref|XP_017149224.1| PREDICTED: hormone-sensitive lipase isoform X2 [Drosophila m
>gi|1036996325|ref|XP_017149225.1| PREDICTED: hormone-sensitive lipase isoform X2 [Drosophila m
>gi|1037005932|ref|XP_017149803.1| PREDICTED: arginyl-tRNA--protein transferase 1 isoform X2 [I
>gi|1037005920|ref|XP_017149802.1| PREDICTED: arginyl-tRNA--protein transferase 1 isoform X1 [I
>gi|1037005942|ref|XP_017149804.1| PREDICTED: arginyl-tRNA--protein transferase 1 isoform X3 [I
>gi|1037005956|ref|XP_017149806.1| PREDICTED: sialin [Drosophila miranda]
>gi|1037005974|ref|XP_017149807.1| PREDICTED: sialin [Drosophila miranda]
>gi|1036998787|ref|XP_017149380.1| PREDICTED: 2-hydroxyacyl-CoA lyase 1 [Drosophila miranda]
>gi|1036998751|ref|XP_017149378.1| PREDICTED: EF-hand domain-containing family member C2 isofor
>gi|1036998733|ref|XP_017149377.1| PREDICTED: EF-hand domain-containing family member C2 isofor
>gi|1036998769|ref|XP_017149379.1| PREDICTED: EF-hand domain-containing family member C2 isofor
>gi|1036998838|ref|XP_017149383.1| PREDICTED: uncharacterized protein LOC108160104 [Drosophila
>gi|1036998805|ref|XP_017149381.1| PREDICTED: leucine-rich repeat and death domain-containing p
>gi|1036998822|ref|XP_017149382.1| PREDICTED: plant intracellular Ras-group-related LRR protei
>gi|1036952333|ref|XP_017146572.1| PREDICTED: uncharacterized protein LOC108158609 [Drosophila
>gi|1036938139|ref|XP_017145707.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036952301|ref|XP_017146570.1| PREDICTED: uncharacterized protein LOC108158607 isoform X2
>gi|1036952317|ref|XP_017146571.1| PREDICTED: uncharacterized protein LOC108158607 isoform X2
>gi|1036952283|ref|XP_017146569.1| PREDICTED: uncharacterized protein LOC108158607 isoform X1
>gi|1036952351|ref|XP_017146573.1| PREDICTED: PRADC1-like protein [Drosophila miranda]
>gi|1036952389|ref|XP_017146577.1| PREDICTED: CCHC-type zinc finger protein CG3800 [Drosophila
>gi|1036952225|ref|XP_017146566.1| PREDICTED: receptor expression-enhancing protein 1 isoform X
>gi|1036952175|ref|XP_017146563.1| PREDICTED: uncharacterized protein LOC108158605 isoform X7
>gi|1036952141|ref|XP_017146561.1| PREDICTED: uncharacterized protein LOC108158605 isoform X5
>gi|1036952101|ref|XP_017146559.1| PREDICTED: uncharacterized protein LOC108158605 isoform X4
>gi|1036952130|ref|XP_017146560.1| PREDICTED: uncharacterized protein LOC108158605 isoform X4
>gi|1036952083|ref|XP_017146558.1| PREDICTED: uncharacterized protein LOC108158605 isoform X3
>gi|1036952067|ref|XP_017146557.1| PREDICTED: uncharacterized protein LOC108158605 isoform X2
>gi|1036952049|ref|XP_017146556.1| PREDICTED: uncharacterized protein LOC108158605 isoform X1
>gi|1036952159|ref|XP_017146562.1| PREDICTED: uncharacterized protein LOC108158605 isoform X6
>gi|1036952209|ref|XP_017146565.1| PREDICTED: probable serine/threonine-protein kinase kinX isc
>gi|1036952269|ref|XP_017146568.1| PREDICTED: receptor expression-enhancing protein 1 isoform X
>gi|1036952193|ref|XP_017146564.1| PREDICTED: receptor expression-enhancing protein 2 isoform X
>gi|1036952251|ref|XP_017146567.1| PREDICTED: uncharacterized protein LOC108158605 isoform X11
>gi|1036952031|ref|XP_017146555.1| PREDICTED: nuclear pore complex protein Nup214 [Drosophila m
>gi|1036952359|ref|XP_017146574.1| PREDICTED: uncharacterized protein LOC108158611 isoform X1
>gi|1036952373|ref|XP_017146575.1| PREDICTED: uncharacterized protein LOC108158611 isoform X2
>gi|1036977664|ref|XP_017148079.1| PREDICTED: LOW QUALITY PROTEIN: protein slit [Drosophila mir
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>gi|1036996725|ref|XP_017149250.1| PREDICTED: uncharacterized protein LOC108160024 isoform X2
>gi|1036996709|ref|XP_017149249.1| PREDICTED: uncharacterized protein LOC108160024 isoform X1
>gi|1036986691|ref|XP_017148642.1| PREDICTED: death-associated inhibitor of apoptosis 2 [Drosop
>gi|1036986823|ref|XP_017148649.1| PREDICTED: uncharacterized protein LOC108159673 [Drosophila
>gi|1036986787|ref|XP_017148646.1| PREDICTED: myeloid leukemia factor isoform X4 [Drosophila m
>gi|1036986719|ref|XP_017148644.1| PREDICTED: myeloid leukemia factor isoform X2 [Drosophila m
>gi|1036986770|ref|XP_017148645.1| PREDICTED: myeloid leukemia factor isoform X3 [Drosophila m
>gi|1036986705|ref|XP_017148643.1| PREDICTED: myeloid leukemia factor isoform X1 [Drosophila m
>gi|1036986805|ref|XP_017148647.1| PREDICTED: myeloid leukemia factor isoform X5 [Drosophila m
>gi|1036977016|ref|XP_017148040.1| PREDICTED: ecdysone receptor isoform X2 [Drosophila miranda]
>gi|1036977034|ref|XP_017148041.1| PREDICTED: ecdysone receptor isoform X3 [Drosophila miranda]
>gi|1036976998|ref|XP_017148039.1| PREDICTED: ecdysone receptor isoform X1 [Drosophila miranda]
>gi|1036977066|ref|XP_017148043.1| PREDICTED: uncharacterized protein LOC108159347 [Drosophila
>gi|1036977052|ref|XP_017148042.1| PREDICTED: probable cytochrome P450 6w1 [Drosophila miranda]
>gi|1036943671|ref|XP_017146046.1| PREDICTED: macrophage mannose receptor 1-like [Drosophila m
>gi|1036938157|ref|XP_017145708.1| PREDICTED: macrophage mannose receptor 1 [Drosophila miranda
>gi|1036943530|ref|XP_017146036.1| PREDICTED: regenerating islet-derived protein 4-like [Drosop
>gi|1036995098|ref|XP_017149151.1| PREDICTED: patched domain-containing protein 3 [Drosophila r
>gi|1037018428|ref|XP_017150563.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037016975|ref|XP_017150468.1| PREDICTED: zinc finger protein 665 [Drosophila miranda]
>gi|1036975004|ref|XP_017147916.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036974986|ref|XP_017147915.1| PREDICTED: tetratricopeptide repeat protein 7B isoform X2 [D
>gi|1036974968|ref|XP_017147914.1| PREDICTED: tetratricopeptide repeat protein 7B isoform X1 [D
>gi|1036974950|ref|XP_017147913.1| PREDICTED: probable RNA methyltransferase bin3 [Drosophila r
>gi|1037010849|ref|XP_017150098.1| PREDICTED: phospholipase D2 isoform X1 [Drosophila miranda]
>gi|1037010869|ref|XP_017150099.1| PREDICTED: phospholipase D2 isoform X1 [Drosophila miranda]
>gi|1037010886|ref|XP_017150100.1| PREDICTED: phospholipase D2 isoform X2 [Drosophila miranda]
>gi|1037010901|ref|XP_017150101.1| PREDICTED: phospholipase D2 isoform X3 [Drosophila miranda]
>gi|1036972578|ref|XP_017147762.1| PREDICTED: zinc finger protein jing isoform X1 [Drosophila r
>gi|1036972629|ref|XP_017147766.1| PREDICTED: uncharacterized protein LOC108159205 isoform X2
>gi|1036972608|ref|XP_017147765.1| PREDICTED: uncharacterized protein LOC108159205 isoform X1
>gi|1036972594|ref|XP_017147763.1| PREDICTED: zinc finger protein jing isoform X2 [Drosophila r
>gi|1036972639|ref|XP_017147767.1| PREDICTED: uncharacterized protein LOC108159206 [Drosophila
>gi|1036986133|ref|XP_017148610.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit
>gi|1036986147|ref|XP_017148611.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit
>gi|1036986117|ref|XP_017148609.1| PREDICTED: heat shock protein 75 kDa, mitochondrial [Drosopl
>gi|1036993330|ref|XP_017149040.1| PREDICTED: AT-rich interactive domain-containing protein 2 :
>gi|1036993352|ref|XP_017149041.1| PREDICTED: AT-rich interactive domain-containing protein 2 :
>gi|1036993312|ref|XP_017149039.1| PREDICTED: AT-rich interactive domain-containing protein 2 :
>gi|1037010154|ref|XP_017150059.1| PREDICTED: senecionine N-oxygenase isoform X1 [Drosophila m
>gi|1037010174|ref|XP_017150060.1| PREDICTED: senecionine N-oxygenase isoform X2 [Drosophila m
>gi|1037010190|ref|XP_017150061.1| PREDICTED: senecionine N-oxygenase isoform X3 [Drosophila m
>gi|1037010206|ref|XP_017150063.1| PREDICTED: bcl-2-related ovarian killer protein homolog B [I
>gi|1037003556|ref|XP_017149663.1| PREDICTED: zinc finger CCHC domain-containing protein 8 hom
>gi|1037003574|ref|XP_017149665.1| PREDICTED: small integral membrane protein 14 [Drosophila m
>gi|1037003590|ref|XP_017149666.1| PREDICTED: small integral membrane protein 14 [Drosophila m
>gi|1037003606|ref|XP_017149667.1| PREDICTED: small integral membrane protein 14 [Drosophila m
>gi|1036938175|ref|XP_017145709.1| PREDICTED: uncharacterized protein LOC108158070 [Drosophila
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>gi|1037004422|ref|XP_017149711.1| PREDICTED: breast cancer type 2 susceptibility protein homol
>gi|1037004404|ref|XP_017149710.1| PREDICTED: breast cancer type 2 susceptibility protein homol
>gi|1037004386|ref|XP_017149709.1| PREDICTED: breast cancer type 2 susceptibility protein homol
>gi|1037004118|ref|XP_017149696.1| PREDICTED: polyadenylate-binding protein [Drosophila miranda
>gi|1037004136|ref|XP_017149697.1| PREDICTED: polyadenylate-binding protein [Drosophila miranda
>gi|1036938193|ref|XP_017145710.1| PREDICTED: uncharacterized protein LOC108158071 [Drosophila
>gi|1036938210|ref|XP_017145711.1| PREDICTED: uncharacterized protein LOC108158072 [Drosophila
>gi|1036938227|ref|XP_017145712.1| PREDICTED: uncharacterized protein LOC108158073 [Drosophila
>gi|1036995706|ref|XP_017149187.1| PREDICTED: LETM1 and EF-hand domain-containing protein anon-
>gi|1036995726|ref|XP_017149188.1| PREDICTED: LETM1 and EF-hand domain-containing protein anon-
>gi|1036995744|ref|XP_017149189.1| PREDICTED: LETM1 and EF-hand domain-containing protein anon-
>gi|1036995760|ref|XP_017149190.1| PREDICTED: LETM1 and EF-hand domain-containing protein anon-
>gi|1036995777|ref|XP_017149191.1| PREDICTED: LETM1 and EF-hand domain-containing protein anon-
>gi|1036943639|ref|XP_017146044.1| PREDICTED: uncharacterized protein LOC108158313 isoform X2
>gi|1036943622|ref|XP_017146042.1| PREDICTED: uncharacterized protein LOC108158313 isoform X1
>gi|1036942241|ref|XP_017145956.1| PREDICTED: proteasome subunit alpha type-7-1B [Drosophila m
>gi|1036941870|ref|XP_017145931.1| PREDICTED: suppressor-of-stellate-like protein isoform X1 [
>gi|1036941885|ref|XP_017145932.1| PREDICTED: suppressor-of-stellate-like protein isoform X2 [
>gi|1036941897|ref|XP_017145933.1| PREDICTED: suppressor-of-stellate-like protein isoform X3 [
>gi|1037020795|ref|XP_017150717.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036946462|ref|XP_017146210.1| PREDICTED: uncharacterized protein LOC108158438 [Drosophila
>gi|1036938245|ref|XP_017145713.1| PREDICTED: uncharacterized protein LOC108158074 [Drosophila
>gi|1037017794|ref|XP_017150522.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037017776|ref|XP_017150521.1| PREDICTED: dynein light chain roadblock-type 1 [Drosophila
>gi|1037017763|ref|XP_017150520.1| PREDICTED: leucine-rich repeat-containing protein 40 [Drosop
>gi|1037017749|ref|XP_017150519.1| PREDICTED: uncharacterized protein LOC108160806 [Drosophila
>gi|1036941681|ref|XP_017145918.1| PREDICTED: LOW QUALITY PROTEIN: luciferin 4-monooxygenase [
>gi|1036943877|ref|XP_017146058.1| PREDICTED: uncharacterized protein LOC108158328 [Drosophila
>gi|1037022338|ref|XP_017150811.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit alpha, mi
>gi|1036943719|ref|XP_017146048.1| PREDICTED: serine/threonine-protein phosphatase Pgam5, mito
>gi|1036938261|ref|XP_017145714.1| PREDICTED: granzyme C [Drosophila miranda]
>gi|1037002548|ref|XP_017149604.1| PREDICTED: uncharacterized protein LOC108160251 [Drosophila
>gi|1037002634|ref|XP_017149609.1| PREDICTED: LOW QUALITY PROTEIN: NSFL1 cofactor p47 [Drosoph
>gi|1037002616|ref|XP_017149607.1| PREDICTED: GPI mannosyltransferase 4-like isoform X2 [Droso
>gi|1037002590|ref|XP_017149605.1| PREDICTED: pupal cuticle protein G1A-like isoform X1 [Droso
>gi|1037002600|ref|XP_017149606.1| PREDICTED: pupal cuticle protein G1A-like isoform X1 [Droso
>gi|1036968522|ref|XP_017147518.1| PREDICTED: sodium-dependent serotonin transporter isoform X
>gi|1036968540|ref|XP_017147519.1| PREDICTED: sodium-dependent serotonin transporter isoform X
>gi|1036968558|ref|XP_017147520.1| PREDICTED: sodium-dependent serotonin transporter isoform X
>gi|1036968576|ref|XP_017147521.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 7 [
>gi|1036968487|ref|XP_017147515.1| PREDICTED: serine/threonine-protein kinase 10 isoform X3 [D
>gi|1036968504|ref|XP_017147516.1| PREDICTED: serine/threonine-protein kinase 10 isoform X4 [D
>gi|1036968453|ref|XP_017147513.1| PREDICTED: serine/threonine-protein kinase 10 isoform X1 [D
>gi|1036968469|ref|XP_017147514.1| PREDICTED: serine/threonine-protein kinase 10 isoform X2 [D
>gi|1037020039|ref|XP_017150669.1| PREDICTED: muscarinic acetylcholine receptor DM1 [Drosophila
>gi|1036942367|ref|XP_017145964.1| PREDICTED: uncharacterized protein LOC108158242 [Drosophila
>gi|1037007057|ref|XP_017149878.1| PREDICTED: serum response factor homolog [Drosophila miranda
>gi|1036938279|ref|XP_017145715.1| PREDICTED: uncharacterized protein LOC108158078, partial [D
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>gi|1037020186|ref|XP_017150677.1| PREDICTED: CCAAT/enhancer-binding protein [Drosophila mirand
>gi|1037008132|ref|XP_017149943.1| PREDICTED: tubulin beta-3 chain [Drosophila miranda]
>gi|1036998680|ref|XP_017149374.1| PREDICTED: long-chain fatty acid transport protein 4 isofor
>gi|1036998698|ref|XP_017149375.1| PREDICTED: long-chain fatty acid transport protein 4 isofor
>gi|1036998716|ref|XP_017149376.1| PREDICTED: long-chain fatty acid transport protein 4 isofor
>gi|1036998642|ref|XP_017149371.1| PREDICTED: protein CBFA2T3 isoform X1 [Drosophila miranda]
>gi|1036998664|ref|XP_017149372.1| PREDICTED: protein CBFA2T3 isoform X2 [Drosophila miranda]
>gi|1037008274|ref|XP_017149952.1| PREDICTED: sphingomyelin phosphodiesterase isoform X2 [Droso
>gi|1037008240|ref|XP_017149950.1| PREDICTED: sphingomyelin phosphodiesterase isoform X1 [Droso
>gi|1037008260|ref|XP_017149951.1| PREDICTED: sphingomyelin phosphodiesterase isoform X1 [Droso
>gi|1036945874|ref|XP_017146172.1| PREDICTED: gastrin/cholecystokinin type B receptor [Drosophi
>gi|1037019991|ref|XP_017150666.1| PREDICTED: uncharacterized protein LOC108160904 [Drosophila
>gi|1036936091|ref|XP_017145571.1| PREDICTED: uncharacterized protein LOC108157938 [Drosophila
>gi|1037022095|ref|XP_017150797.1| PREDICTED: uncharacterized protein LOC108161000 [Drosophila
>gi|1036963545|ref|XP_017147236.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1036963515|ref|XP_017147233.1| PREDICTED: ubiquitin-protein ligase E3C [Drosophila miranda]
>gi|1036963537|ref|XP_017147234.1| PREDICTED: 7-methylguanosine phosphate-specific 5'-nucleotic
>gi|1036963424|ref|XP_017147228.1| PREDICTED: uncharacterized protein LOC108158933 isoform X1
>gi|1036963442|ref|XP_017147229.1| PREDICTED: uncharacterized protein LOC108158933 isoform X2
>gi|1036963458|ref|XP_017147230.1| PREDICTED: uncharacterized protein LOC108158933 isoform X3
>gi|1036963479|ref|XP_017147231.1| PREDICTED: uncharacterized protein LOC108158933 isoform X4
>gi|1036963496|ref|XP_017147232.1| PREDICTED: uncharacterized protein LOC108158933 isoform X4
>gi|1037005095|ref|XP_017149748.1| PREDICTED: H/ACA ribonucleoprotein complex subunit 4 [Droso
>gi|1037006305|ref|XP_017149829.1| PREDICTED: ras-like GTP-binding protein Rho1 [Drosophila mi
>gi|1037011721|ref|XP_017150149.1| PREDICTED: GTP-binding protein Rit2 [Drosophila miranda]
>gi|1037003713|ref|XP_017149673.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X1
>gi|1037003853|ref|XP_017149682.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X9
>gi|1037003817|ref|XP_017149680.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X7
>gi|1037003871|ref|XP_017149683.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X10
>gi|1037003799|ref|XP_017149679.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X6
>gi|1037003749|ref|XP_017149675.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X3
>gi|1037003767|ref|XP_017149676.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X4
>gi|1037003731|ref|XP_017149674.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X2
>gi|1037003783|ref|XP_017149677.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X5
>gi|1037003835|ref|XP_017149681.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X8
>gi|1037003999|ref|XP_017149689.1| PREDICTED: aspartyl/asparaginyl beta-hydroxylase isoform X15
>gi|1037003887|ref|XP_017149684.1| PREDICTED: aspartic and glutamic acid-rich protein isoform
>gi|1037003905|ref|XP_017149685.1| PREDICTED: aspartic and glutamic acid-rich protein isoform
>gi|1037003921|ref|XP_017149686.1| PREDICTED: aspartic and glutamic acid-rich protein isoform
>gi|1037003972|ref|XP_017149688.1| PREDICTED: FK506-binding protein 5 isoform X14 [Drosophila
>gi|1037003954|ref|XP_017149687.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1036938296|ref|XP_017145717.1| PREDICTED: esterase 6-like [Drosophila miranda]
>gi|1037004084|ref|XP_017149694.1| PREDICTED: esterase-6-like [Drosophila miranda]
>gi|1037004038|ref|XP_017149690.1| PREDICTED: esterase E4 isoform X1 [Drosophila miranda]
>gi|1037004066|ref|XP_017149693.1| PREDICTED: esterase E4 isoform X1 [Drosophila miranda]
>gi|1037004054|ref|XP_017149691.1| PREDICTED: esterase E4 isoform X2 [Drosophila miranda]
>gi|1036938312|ref|XP_017145718.1| PREDICTED: uncharacterized protein LOC108158081 [Drosophila
>gi|1036995148|ref|XP_017149152.1| PREDICTED: protein spinster [Drosophila miranda]
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>gi|1037016578|ref|XP_017150443.1| PREDICTED: uncharacterized zinc finger protein CG12744 [Dros
>gi|1037016596|ref|XP_017150444.1| PREDICTED: uncharacterized zinc finger protein CG12744 [Dros
>gi|1036986841|ref|XP_017148650.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X1 [Drosoph
>gi|1036986881|ref|XP_017148652.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X3 [Drosoph
>gi|1036986860|ref|XP_017148651.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X2 [Drosoph
>gi|1036986902|ref|XP_017148653.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X4 [Drosoph
>gi|1036986920|ref|XP_017148654.1| PREDICTED: spermine oxidase [Drosophila miranda]
>gi|1036986942|ref|XP_017148655.1| PREDICTED: syntaxin-10 [Drosophila miranda]
>gi|1036941010|ref|XP_017145880.1| PREDICTED: phospholipid scramblase 3 isoform X1 [Drosophila
>gi|1036941028|ref|XP_017145881.1| PREDICTED: phospholipid scramblase 3 isoform X2 [Drosophila
>gi|1036962688|ref|XP_017147182.1| PREDICTED: uncharacterized protein LOC108158904 [Drosophila
>gi|1036962674|ref|XP_017147181.1| PREDICTED: uncharacterized protein LOC108158903 [Drosophila
>gi|1036962562|ref|XP_017147175.1| PREDICTED: uncharacterized zinc finger protein CG2678 [Dros
>gi|1036962642|ref|XP_017147179.1| PREDICTED: uncharacterized protein LOC108158902 [Drosophila
>gi|1036962618|ref|XP_017147178.1| PREDICTED: sex peptide receptor [Drosophila miranda]
>gi|1036962582|ref|XP_017147176.1| PREDICTED: Bardet-Biedl syndrome 4 protein homolog [Drosoph
>gi|1036962600|ref|XP_017147177.1| PREDICTED: Bardet-Biedl syndrome 4 protein homolog [Drosoph
>gi|1037020422|ref|XP_017150692.1| PREDICTED: probable cytochrome P450 12d1 proximal, mitochond
>gi|1037020438|ref|XP_017150693.1| PREDICTED: probable cytochrome P450 12d1 proximal, mitochond
>gi|1036996567|ref|XP_017149241.1| PREDICTED: Krueppel-like factor 7 [Drosophila miranda]
>gi|1036996585|ref|XP_017149242.1| PREDICTED: uncharacterized protein LOC108160018 [Drosophila
>gi|1037019650|ref|XP_017150642.1| PREDICTED: leukocyte elastase inhibitor [Drosophila miranda]
>gi|1036941183|ref|XP_017145891.1| PREDICTED: general odorant-binding protein 56a [Drosophila r
>gi|1036943216|ref|XP_017146017.1| PREDICTED: proton-coupled amino acid transporter 4 isoform X
>gi|1036943234|ref|XP_017146019.1| PREDICTED: proton-coupled amino acid transporter 4 isoform X
>gi|1036943250|ref|XP_017146020.1| PREDICTED: uncharacterized protein LOC108158292 isoform X3
>gi|1036995176|ref|XP_017149153.1| PREDICTED: ATP-citrate synthase isoform X1 [Drosophila mirar
>gi|1036995192|ref|XP_017149155.1| PREDICTED: ATP-citrate synthase isoform X1 [Drosophila mirar
>gi|1036995209|ref|XP_017149156.1| PREDICTED: ATP-citrate synthase isoform X1 [Drosophila mirar
>gi|1036995224|ref|XP_017149157.1| PREDICTED: ATP-citrate synthase isoform X2 [Drosophila mirar
>gi|1036995241|ref|XP_017149158.1| PREDICTED: ATP-citrate synthase isoform X3 [Drosophila mirar
>gi|1036995275|ref|XP_017149160.1| PREDICTED: transmembrane protein 208 isoform X2 [Drosophila
>gi|1036995258|ref|XP_017149159.1| PREDICTED: transmembrane protein 208 isoform X1 [Drosophila
>gi|1036962878|ref|XP_017147193.1| PREDICTED: peroxisomal membrane protein 11B [Drosophila mira
>gi|1036962836|ref|XP_017147191.1| PREDICTED: palmitoyltransferase ZDHHC3 [Drosophila miranda]
>gi|1036962706|ref|XP_017147183.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X1
>gi|1036962726|ref|XP_017147184.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X2
>gi|1036962743|ref|XP_017147185.1| PREDICTED: muscle M-line assembly protein unc-89 isoform X3
>gi|1036962818|ref|XP_017147190.1| PREDICTED: uncharacterized protein LOC108158907 [Drosophila
>gi|1036962762|ref|XP_017147186.1| PREDICTED: myosin light chain kinase, smooth muscle isoform
>gi|1036962780|ref|XP_017147187.1| PREDICTED: myosin light chain kinase, smooth muscle isoform
>gi|1036962800|ref|XP_017147188.1| PREDICTED: probable cytochrome P450 4aa1 [Drosophila miranda
>gi|1036962860|ref|XP_017147192.1| PREDICTED: trypsin alpha-3 [Drosophila miranda]
>gi|1036996022|ref|XP_017149206.1| PREDICTED: serrate RNA effector molecule homolog isoform X2
>gi|1036996004|ref|XP_017149205.1| PREDICTED: serrate RNA effector molecule homolog isoform X1
>gi|1036996040|ref|XP_017149207.1| PREDICTED: serrate RNA effector molecule homolog isoform X3
>gi|1037009014|ref|XP_017149995.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1037009026|ref|XP_017149996.1| PREDICTED: probable tRNA pseudouridine synthase 2 [Drosophil
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>gi|1037007571|ref|XP_017149908.1| PREDICTED: uncharacterized protein LOC108160420 isoform X1
>gi|1037007589|ref|XP_017149909.1| PREDICTED: uncharacterized protein LOC108160420 isoform X2
>gi|1037007603|ref|XP_017149910.1| PREDICTED: integumentary mucin C.1-like [Drosophila miranda]
>gi|1037007625|ref|XP_017149911.1| PREDICTED: integumentary mucin C.1-like [Drosophila miranda]
>gi|1037007640|ref|XP_017149912.1| PREDICTED: integumentary mucin C.1-like [Drosophila miranda]
>gi|1037007553|ref|XP_017149907.1| PREDICTED: WD repeat-containing protein 74 [Drosophila mira
>gi|1036938330|ref|XP_017145719.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036938348|ref|XP_017145720.1| PREDICTED: neuropeptide CCHamide-2 receptor [Drosophila mira
>gi|1037022411|ref|XP_017150816.1| PREDICTED: putative uncharacterized protein DDB_G0290521 [Dr
>gi|1036943773|ref|XP_017146051.1| PREDICTED: uncharacterized protein LOC108158322 [Drosophila
>gi|1037021990|ref|XP_017150791.1| PREDICTED: pro-resilin isoform X2 [Drosophila miranda]
>gi|1037021977|ref|XP_017150789.1| PREDICTED: pro-resilin isoform X1 [Drosophila miranda]
>gi|1036944852|ref|XP_017146112.1| PREDICTED: vitamin K epoxide reductase complex subunit 1 [Dr
>gi|1036983866|ref|XP_017148467.1| PREDICTED: LOW QUALITY PROTEIN: actin [Drosophila miranda]
>gi|1036983901|ref|XP_017148469.1| PREDICTED: superoxide dismutase [Mn], mitochondrial isoform
>gi|1036983884|ref|XP_017148468.1| PREDICTED: superoxide dismutase [Mn], mitochondrial isoform
>gi|1036984013|ref|XP_017148476.1| PREDICTED: uncharacterized protein LOC108159559 [Drosophila
>gi|1036983597|ref|XP_017148449.1| PREDICTED: kinesin-like protein unc-104 isoform X1 [Drosophi
>gi|1036983615|ref|XP_017148450.1| PREDICTED: kinesin-like protein unc-104 isoform X1 [Drosophi
>gi|1036983638|ref|XP_017148451.1| PREDICTED: kinesin-like protein unc-104 isoform X2 [Drosophi
>gi|1036983724|ref|XP_017148456.1| PREDICTED: kinesin-like protein unc-104 isoform X7 [Drosophi
>gi|1036983652|ref|XP_017148452.1| PREDICTED: kinesin-like protein unc-104 isoform X3 [Drosophi
>gi|1036983706|ref|XP_017148455.1| PREDICTED: kinesin-like protein unc-104 isoform X6 [Drosophi
>gi|1036983666|ref|XP_017148453.1| PREDICTED: kinesin-like protein unc-104 isoform X4 [Drosophi
>gi|1036983766|ref|XP_017148460.1| PREDICTED: kinesin-like protein unc-104 isoform X10 [Drosoph
>gi|1036983740|ref|XP_017148457.1| PREDICTED: kinesin-like protein unc-104 isoform X8 [Drosophi
>gi|1036983822|ref|XP_017148464.1| PREDICTED: kinesin-like protein unc-104 isoform X14 [Drosoph
>gi|1036983690|ref|XP_017148454.1| PREDICTED: kinesin-like protein unc-104 isoform X5 [Drosophi
>gi|1036983778|ref|XP_017148461.1| PREDICTED: kinesin-like protein unc-104 isoform X11 [Drosoph
>gi|1036983806|ref|XP_017148463.1| PREDICTED: kinesin-like protein unc-104 isoform X13 [Drosoph
>gi|1036983792|ref|XP_017148462.1| PREDICTED: kinesin-like protein unc-104 isoform X12 [Drosoph
>gi|1036983752|ref|XP_017148458.1| PREDICTED: kinesin-like protein unc-104 isoform X9 [Drosophi
>gi|1036983838|ref|XP_017148465.1| PREDICTED: kinesin-like protein unc-104 isoform X15 [Drosoph
>gi|1036983852|ref|XP_017148466.1| PREDICTED: kinesin-like protein unc-104 isoform X16 [Drosoph
>gi|1036983995|ref|XP_017148475.1| PREDICTED: uncharacterized protein LOC108159558 [Drosophila
>gi|1036983919|ref|XP_017148470.1| PREDICTED: uncharacterized protein LOC108159555 [Drosophila
>gi|1036983937|ref|XP_017148471.1| PREDICTED: uncharacterized protein LOC108159556 isoform X1
>gi|1036983955|ref|XP_017148472.1| PREDICTED: uncharacterized protein LOC108159556 isoform X2
>gi|1036983975|ref|XP_017148474.1| PREDICTED: accessory gland protein Acp53Ea-like [Drosophila
>gi|1036985371|ref|XP_017148560.1| PREDICTED: uncharacterized protein LOC108159616 [Drosophila
>gi|1036985391|ref|XP_017148561.1| PREDICTED: accessory gland protein Acp53Ea [Drosophila mira
>gi|1036985341|ref|XP_017148557.1| PREDICTED: uncharacterized protein LOC108159614 [Drosophila
>gi|1036985359|ref|XP_017148559.1| PREDICTED: uncharacterized protein LOC108159615 [Drosophila
>gi|1036985300|ref|XP_017148554.1| PREDICTED: putative polypeptide N-acetylgalactosaminyltrans
>gi|1036985316|ref|XP_017148555.1| PREDICTED: putative polypeptide N-acetylgalactosaminyltrans
>gi|1036985330|ref|XP_017148556.1| PREDICTED: chitinase-like protein CG5210 [Drosophila miranda
>gi|1036938366|ref|XP_017145721.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1037010506|ref|XP_017150082.1| PREDICTED: cilia- and flagella-associated protein 161 [Droso
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>gi|1037010534|ref|XP_017150084.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1037010520|ref|XP_017150083.1| PREDICTED: putative inorganic phosphate cotransporter isofo
>gi|1036944204|ref|XP_017146077.1| PREDICTED: LOW QUALITY PROTEIN: sodium channel protein Nach
>gi|1037013907|ref|XP_017150285.1| PREDICTED: alpha-amylase-related protein [Drosophila miranda
>gi|1037013957|ref|XP_017150289.1| PREDICTED: serine-enriched protein isoform X3 [Drosophila mi
>gi|1037013921|ref|XP_017150286.1| PREDICTED: serine-enriched protein isoform X1 [Drosophila mi
>gi|1037013941|ref|XP_017150287.1| PREDICTED: serine-enriched protein isoform X2 [Drosophila mi
>gi|1037010452|ref|XP_017150079.1| PREDICTED: ALK tyrosine kinase receptor [Drosophila miranda]
>gi|1037010470|ref|XP_017150080.1| PREDICTED: ALK tyrosine kinase receptor [Drosophila miranda]
>gi|1037010488|ref|XP_017150081.1| PREDICTED: ALK tyrosine kinase receptor [Drosophila miranda]
>gi|1036986161|ref|XP_017148612.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosoph
>gi|1036986177|ref|XP_017148613.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosoph
>gi|1036986193|ref|XP_017148614.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosoph
>gi|1036986274|ref|XP_017148620.1| PREDICTED: uncharacterized protein LOC108159657 [Drosophila
>gi|1036986209|ref|XP_017148615.1| PREDICTED: putative fatty acyl-CoA reductase CG8303 isoform
>gi|1036986239|ref|XP_017148618.1| PREDICTED: putative fatty acyl-CoA reductase CG8303 isoform
>gi|1036986221|ref|XP_017148617.1| PREDICTED: putative fatty acyl-CoA reductase CG8303 isoform
>gi|1036986257|ref|XP_017148619.1| PREDICTED: putative fatty acyl-CoA reductase CG8306 [Drosoph
>gi|1037003111|ref|XP_017149633.1| PREDICTED: LOW QUALITY PROTEIN: putative uncharacterized pro
>gi|1036998371|ref|XP_017149354.1| PREDICTED: uncharacterized protein LOC108160088 [Drosophila
>gi|1036998387|ref|XP_017149355.1| PREDICTED: uncharacterized protein LOC108160088 [Drosophila
>gi|1036998317|ref|XP_017149351.1| PREDICTED: endophilin-B1 isoform X2 [Drosophila miranda]
>gi|1036998353|ref|XP_017149353.1| PREDICTED: endophilin-B1 isoform X4 [Drosophila miranda]
>gi|1036998297|ref|XP_017149350.1| PREDICTED: endophilin-B1 isoform X1 [Drosophila miranda]
>gi|1036998335|ref|XP_017149352.1| PREDICTED: endophilin-B1 isoform X3 [Drosophila miranda]
>gi|1036984315|ref|XP_017148493.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036984391|ref|XP_017148498.1| PREDICTED: uncharacterized protein LOC108159569 [Drosophila
>gi|1036984409|ref|XP_017148499.1| PREDICTED: uncharacterized protein LOC108159569 [Drosophila
>gi|1036984425|ref|XP_017148500.1| PREDICTED: uncharacterized protein LOC108159570 isoform X1
>gi|1036984445|ref|XP_017148501.1| PREDICTED: uncharacterized protein LOC108159570 isoform X2
>gi|1036984333|ref|XP_017148495.1| PREDICTED: F-box only protein 42 isoform X1 [Drosophila mira
>gi|1036984351|ref|XP_017148496.1| PREDICTED: F-box only protein 42 isoform X2 [Drosophila mira
>gi|1036984371|ref|XP_017148497.1| PREDICTED: F-box only protein 42 isoform X2 [Drosophila mira
>gi|1036997427|ref|XP_017149293.1| PREDICTED: tRNA pseudouridine(38/39) synthase [Drosophila mi
>gi|1036997410|ref|XP_017149292.1| PREDICTED: uncharacterized protein LOC108160056 isoform X2
>gi|1036997372|ref|XP_017149290.1| PREDICTED: uncharacterized protein LOC108160056 isoform X1
>gi|1036997394|ref|XP_017149291.1| PREDICTED: uncharacterized protein LOC108160056 isoform X1
>gi|1036997463|ref|XP_017149295.1| PREDICTED: uncharacterized protein LOC108160060 [Drosophila
>gi|1036997445|ref|XP_017149294.1| PREDICTED: stress response protein NST1 [Drosophila miranda]
>gi|1036997483|ref|XP_017149296.1| PREDICTED: ribonuclease kappa [Drosophila miranda]
>gi|1036938382|ref|XP_017145722.1| PREDICTED: LOW QUALITY PROTEIN: cuticle protein 16.5 [Drosop
>gi|1036997176|ref|XP_017149279.1| PREDICTED: neuronal acetylcholine receptor subunit beta-3 [
>gi|1036997158|ref|XP_017149278.1| PREDICTED: golgin subfamily A member 2 [Drosophila miranda]
>gi|1036997194|ref|XP_017149280.1| PREDICTED: shugoshin [Drosophila miranda]
>gi|1036982354|ref|XP_017148375.1| PREDICTED: microfibril-associated glycoprotein 4 [Drosophila
>gi|1036982372|ref|XP_017148377.1| PREDICTED: ficolin-1 [Drosophila miranda]
>gi|1036982304|ref|XP_017148372.1| PREDICTED: WAS/WASL-interacting protein family member 1 isof
>gi|1036982251|ref|XP_017148368.1| PREDICTED: verprolin isoform X1 [Drosophila miranda]
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>gi|1036982266|ref|XP_017148369.1| PREDICTED: verprolin isoform X1 [Drosophila miranda]
>gi|1036982286|ref|XP_017148370.1| PREDICTED: verprolin isoform X2 [Drosophila miranda]
>gi|1036982334|ref|XP_017148374.1| PREDICTED: pleckstrin homology domain-containing family M me
>gi|1036982320|ref|XP_017148373.1| PREDICTED: uncharacterized protein KIAA0226-like isoform X1
>gi|1036995455|ref|XP_017149171.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1036995489|ref|XP_017149174.1| PREDICTED: cell surface glycoprotein 1 isoform X3 [Drosophil
>gi|1036995473|ref|XP_017149172.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1036995503|ref|XP_017149175.1| PREDICTED: probable peroxisomal acyl-coenzyme A oxidase 1 [D
>gi|1037002127|ref|XP_017149580.1| PREDICTED: uncharacterized protein LOC108160234 [Drosophila
>gi|1037002075|ref|XP_017149575.1| PREDICTED: probable peroxisomal acyl-coenzyme A oxidase 1 [D
>gi|1037002093|ref|XP_017149576.1| PREDICTED: selenocysteine-specific elongation factor isofor
>gi|1037002109|ref|XP_017149579.1| PREDICTED: selenocysteine-specific elongation factor isofor
>gi|1037002145|ref|XP_017149581.1| PREDICTED: TM2 domain-containing protein CG10795 [Drosophila
>gi|1036999918|ref|XP_017149443.1| PREDICTED: BMP-binding endothelial regulator protein [Drosop
>gi|1036999936|ref|XP_017149444.1| PREDICTED: BMP-binding endothelial regulator protein [Drosop
>gi|1036999952|ref|XP_017149445.1| PREDICTED: BMP-binding endothelial regulator protein [Drosop
>gi|1036999970|ref|XP_017149446.1| PREDICTED: BMP-binding endothelial regulator protein [Drosop
>gi|1036941942|ref|XP_017145937.1| PREDICTED: venom allergen 3 isoform X1 [Drosophila miranda]
>gi|1036941957|ref|XP_017145938.1| PREDICTED: venom allergen 3 isoform X2 [Drosophila miranda]
>gi|1036941987|ref|XP_017145940.1| PREDICTED: venom allergen 5 isoform X4 [Drosophila miranda]
>gi|1036941974|ref|XP_017145939.1| PREDICTED: venom allergen 5 isoform X3 [Drosophila miranda]
>gi|1036943386|ref|XP_017146028.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036962062|ref|XP_017147144.1| PREDICTED: protein PTHB1 [Drosophila miranda]
>gi|1036962078|ref|XP_017147145.1| PREDICTED: protein PTHB1 [Drosophila miranda]
>gi|1036962096|ref|XP_017147146.1| PREDICTED: glycerol-3-phosphate dehydrogenase, mitochondrial
>gi|1036962116|ref|XP_017147147.1| PREDICTED: glycerol-3-phosphate dehydrogenase, mitochondrial
>gi|1036962134|ref|XP_017147148.1| PREDICTED: uncharacterized protein LOC108158884 [Drosophila
>gi|1036962171|ref|XP_017147150.1| PREDICTED: protein N-terminal glutamine amidohydrolase [Dros
>gi|1036962153|ref|XP_017147149.1| PREDICTED: facilitated trehalose transporter Tret1 [Drosophi
>gi|1036962187|ref|XP_017147151.1| PREDICTED: paired box pox-neuro protein [Drosophila miranda]
>gi|1036962044|ref|XP_017147143.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X16 [Dros
>gi|1036961812|ref|XP_017147128.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X2 [Droso
>gi|1036961998|ref|XP_017147139.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X13 [Dros
>gi|1036961883|ref|XP_017147132.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X6 [Droso
>gi|1036962014|ref|XP_017147140.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X14 [Dros
>gi|1036961940|ref|XP_017147136.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X10 [Dros
>gi|1036961761|ref|XP_017147124.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X1 [Droso
>gi|1036961779|ref|XP_017147126.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X1 [Droso
>gi|1036961797|ref|XP_017147127.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X1 [Droso
>gi|1036961896|ref|XP_017147133.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X7 [Droso
>gi|1036961958|ref|XP_017147137.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X11 [Dros
>gi|1036961923|ref|XP_017147135.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X9 [Droso
>gi|1036961851|ref|XP_017147130.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X4 [Droso
>gi|1036961976|ref|XP_017147138.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X12 [Dros
>gi|1036961866|ref|XP_017147131.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X5 [Droso
>gi|1036962028|ref|XP_017147142.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X15 [Dros
>gi|1036961827|ref|XP_017147129.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X3 [Droso
>gi|1036961912|ref|XP_017147134.1| PREDICTED: PDZ and LIM domain protein Zasp isoform X8 [Droso
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>gi|1037001980|ref|XP_017149571.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1037001957|ref|XP_017149570.1| PREDICTED: ataxin-2 homolog isoform X1 [Drosophila miranda]
>gi|1037008792|ref|XP_017149980.1| PREDICTED: venom protease-like [Drosophila miranda]
>gi|1037008846|ref|XP_017149984.1| PREDICTED: uncharacterized protein LOC108160475 [Drosophila
>gi|1037008750|ref|XP_017149978.1| PREDICTED: venom protease-like isoform X2 [Drosophila miran
>gi|1037008732|ref|XP_017149977.1| PREDICTED: serine protease easter-like isoform X1 [Drosophil
>gi|1037008772|ref|XP_017149979.1| PREDICTED: serine protease easter-like isoform X3 [Drosophil
>gi|1037008864|ref|XP_017149985.1| PREDICTED: uncharacterized protein LOC108160476 [Drosophila
>gi|1037008826|ref|XP_017149983.1| PREDICTED: serine protease 44-like [Drosophila miranda]
>gi|1037008810|ref|XP_017149982.1| PREDICTED: chymotrypsin-like protease CTRL-1 [Drosophila mi
>gi|1036963745|ref|XP_017147247.1| PREDICTED: protein Cep89 homolog isoform X1 [Drosophila mira
>gi|1036963763|ref|XP_017147248.1| PREDICTED: protein Cep89 homolog isoform X2 [Drosophila mira
>gi|1036938399|ref|XP_017145723.1| PREDICTED: importin-11 [Drosophila miranda]
>gi|1036963817|ref|XP_017147251.1| PREDICTED: uncharacterized protein LOC108158942 [Drosophila
>gi|1036963801|ref|XP_017147250.1| PREDICTED: V-type proton ATPase subunit F 1 [Drosophila mira
>gi|1036963783|ref|XP_017147249.1| PREDICTED: mannose-1-phosphate guanyltransferase alpha [Dros
>gi|1036963610|ref|XP_017147240.1| PREDICTED: RNA-binding protein fusilli isoform X3 [Drosophil
>gi|1036963651|ref|XP_017147241.1| PREDICTED: RNA-binding protein fusilli isoform X4 [Drosophil
>gi|1036963557|ref|XP_017147237.1| PREDICTED: RNA-binding protein fusilli isoform X1 [Drosophil
>gi|1036963575|ref|XP_017147238.1| PREDICTED: RNA-binding protein fusilli isoform X1 [Drosophil
>gi|1036963588|ref|XP_017147239.1| PREDICTED: RNA-binding protein fusilli isoform X2 [Drosophil
>gi|1036963673|ref|XP_017147242.1| PREDICTED: RNA-binding protein fusilli isoform X5 [Drosophil
>gi|1036963689|ref|XP_017147243.1| PREDICTED: RNA-binding protein fusilli isoform X6 [Drosophil
>gi|1036963707|ref|XP_017147244.1| PREDICTED: RNA-binding protein fusilli isoform X6 [Drosophil
>gi|1036963727|ref|XP_017147246.1| PREDICTED: RNA-binding protein fusilli isoform X6 [Drosophil
>gi|1037016446|ref|XP_017150435.1| PREDICTED: uncharacterized protein LOC108160751 isoform X3
>gi|1037016383|ref|XP_017150433.1| PREDICTED: uncharacterized protein LOC108160751 isoform X1
>gi|1037016479|ref|XP_017150437.1| PREDICTED: uncharacterized protein LOC108160751 isoform X5
>gi|1037016428|ref|XP_017150434.1| PREDICTED: uncharacterized protein LOC108160751 isoform X2
>gi|1037016463|ref|XP_017150436.1| PREDICTED: uncharacterized protein LOC108160751 isoform X4
>gi|1036997058|ref|XP_017149271.1| PREDICTED: zinc finger HIT domain-containing protein 3 [Dros
>gi|1036996961|ref|XP_017149265.1| PREDICTED: beta-glucuronidase [Drosophila miranda]
>gi|1036996978|ref|XP_017149266.1| PREDICTED: beta-glucuronidase isoform X1 [Drosophila miranda
>gi|1036996996|ref|XP_017149267.1| PREDICTED: beta-glucuronidase isoform X2 [Drosophila miranda
>gi|1036997012|ref|XP_017149268.1| PREDICTED: beta-glucuronidase isoform X3 [Drosophila miranda
>gi|1036997030|ref|XP_017149269.1| PREDICTED: beta-glucuronidase isoform X4 [Drosophila miranda
>gi|1036997046|ref|XP_017149270.1| PREDICTED: beta-glucuronidase isoform X4 [Drosophila miranda
>gi|1036999425|ref|XP_017149416.1| PREDICTED: LOW QUALITY PROTEIN: exostosin-3 [Drosophila mira
>gi|1036999441|ref|XP_017149417.1| PREDICTED: ankyrin repeat domain-containing protein 13D [Dro
>gi|1037011703|ref|XP_017150148.1| PREDICTED: monoacylglycerol lipase ABHD12 isoform X2 [Drosop
>gi|1037011669|ref|XP_017150146.1| PREDICTED: monoacylglycerol lipase ABHD12 isoform X1 [Drosop
>gi|1037011685|ref|XP_017150147.1| PREDICTED: monoacylglycerol lipase ABHD12 isoform X1 [Drosop
>gi|1037005285|ref|XP_017149760.1| PREDICTED: LOW QUALITY PROTEIN: kinesin-like protein Klp61F
>gi|1037005213|ref|XP_017149755.1| PREDICTED: uncharacterized protein LOC108160335 isoform X2
>gi|1037005195|ref|XP_017149754.1| PREDICTED: uncharacterized protein LOC108160335 isoform X1
>gi|1037005231|ref|XP_017149756.1| PREDICTED: uncharacterized protein LOC108160335 isoform X3
>gi|1037005249|ref|XP_017149757.1| PREDICTED: uncharacterized protein LOC108160335 isoform X4
>gi|1037005267|ref|XP_017149759.1| PREDICTED: uncharacterized protein LOC108160335 isoform X4
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>gi|1036903214|ref|XP_017153651.1| PREDICTED: monocarboxylate transporter 13 [Drosophila miran
>gi|1036903232|ref|XP_017153652.1| PREDICTED: uncharacterized protein LOC108163084 isoform X1
>gi|1036903248|ref|XP_017153653.1| PREDICTED: uncharacterized protein LOC108163084 isoform X2
>gi|1036919864|ref|XP_017154580.1| PREDICTED: guanine nucleotide-binding protein subunit alpha
>gi|1036885326|ref|XP_017152653.1| PREDICTED: congested-like trachea protein isoform X2 [Droso
>gi|1036885273|ref|XP_017152650.1| PREDICTED: congested-like trachea protein isoform X1 [Droso
>gi|1036885292|ref|XP_017152651.1| PREDICTED: congested-like trachea protein isoform X1 [Droso
>gi|1036885309|ref|XP_017152652.1| PREDICTED: congested-like trachea protein isoform X1 [Droso
>gi|1036885254|ref|XP_017152649.1| PREDICTED: platelet glycoprotein V-like [Drosophila miranda]
>gi|1036885119|ref|XP_017152641.1| PREDICTED: meiosis arrest female protein 1 [Drosophila mira
>gi|1036885195|ref|XP_017152645.1| PREDICTED: scavenger receptor class B member 1 isoform X4 [
>gi|1036885176|ref|XP_017152644.1| PREDICTED: scavenger receptor class B member 1 isoform X3 [
>gi|1036885138|ref|XP_017152642.1| PREDICTED: scavenger receptor class B member 1 isoform X1 [
>gi|1036885157|ref|XP_017152643.1| PREDICTED: scavenger receptor class B member 1 isoform X2 [
>gi|1036885214|ref|XP_017152646.1| PREDICTED: scavenger receptor class B member 1 isoform X5 [
>gi|1036885233|ref|XP_017152648.1| PREDICTED: scavenger receptor class B member 1 isoform X5 [
>gi|1036869955|ref|XP_017151802.1| PREDICTED: carnitine O-acetyltransferase-like isoform X1 [Dr
>gi|1036869974|ref|XP_017151803.1| PREDICTED: carnitine O-acetyltransferase-like isoform X2 [Dr
>gi|1036869993|ref|XP_017151804.1| PREDICTED: carnitine O-acetyltransferase-like isoform X3 [Dr
>gi|1036870011|ref|XP_017151805.1| PREDICTED: carnitine O-acetyltransferase-like isoform X3 [Dr
>gi|1036870030|ref|XP_017151806.1| PREDICTED: carnitine O-acetyltransferase-like isoform X3 [Dr
>gi|1036870379|ref|XP_017151827.1| PREDICTED: uncharacterized protein LOC108161922 [Drosophila
>gi|1036870068|ref|XP_017151809.1| PREDICTED: kelch-like protein 10 isoform X2 [Drosophila mira
>gi|1036870049|ref|XP_017151807.1| PREDICTED: kelch-like protein 10 isoform X1 [Drosophila mira
>gi|1036870342|ref|XP_017151825.1| PREDICTED: glycoprotein hormone beta-5 [Drosophila miranda]
>gi|1036870305|ref|XP_017151823.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036870324|ref|XP_017151824.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036870215|ref|XP_017151817.1| PREDICTED: uncharacterized protein YJR142W isoform X1 [Droso
>gi|1036870234|ref|XP_017151818.1| PREDICTED: uncharacterized protein YJR142W isoform X2 [Droso
>gi|1036870253|ref|XP_017151819.1| PREDICTED: uncharacterized protein YJR142W isoform X3 [Droso
>gi|1036870360|ref|XP_017151826.1| PREDICTED: 60S ribosomal protein L21 [Drosophila miranda]
>gi|1036870179|ref|XP_017151815.1| PREDICTED: zinc finger protein 28-like isoform X1 [Drosophil
>gi|1036870197|ref|XP_017151816.1| PREDICTED: zinc finger and SCAN domain-containing protein 3
>gi|1036870160|ref|XP_017151814.1| PREDICTED: RNA polymerase I-specific transcription initiatio
>gi|1036870288|ref|XP_017151822.1| PREDICTED: deoxynucleotidyltransferase terminal-interacting
>gi|1036870270|ref|XP_017151820.1| PREDICTED: iron-sulfur protein NUBPL [Drosophila miranda]
>gi|1036870087|ref|XP_017151810.1| PREDICTED: uncharacterized protein LOC108161912 isoform X1
>gi|1036870123|ref|XP_017151812.1| PREDICTED: uncharacterized protein LOC108161912 isoform X3
>gi|1036870104|ref|XP_017151811.1| PREDICTED: uncharacterized protein LOC108161912 isoform X2
>gi|1036870142|ref|XP_017151813.1| PREDICTED: uncharacterized protein LOC108161912 isoform X4
>gi|1036884301|ref|XP_017152596.1| PREDICTED: 60S ribosomal protein L5 [Drosophila miranda]
>gi|1036884215|ref|XP_017152590.1| PREDICTED: vacuolar protein sorting-associated protein 41 ho
>gi|1036884284|ref|XP_017152595.1| PREDICTED: beta-1,3-galactosyltransferase 6 [Drosophila mira
>gi|1036884332|ref|XP_017152597.1| PREDICTED: uncharacterized protein LOC108162391 [Drosophila
>gi|1036884232|ref|XP_017152592.1| PREDICTED: serine/threonine-protein kinase GA29083 isoform
>gi|1036884249|ref|XP_017152593.1| PREDICTED: serine/threonine-protein kinase GA29083 isoform
>gi|1036884267|ref|XP_017152594.1| PREDICTED: serine/threonine-protein kinase GA29083 isoform
>gi|1036902006|ref|XP_017153588.1| PREDICTED: LOW QUALITY PROTEIN: sarcolemmal membrane-associa
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>gi|1036901950|ref|XP_017153583.1| PREDICTED: splicing factor 45 [Drosophila miranda]
>gi|1036901968|ref|XP_017153586.1| PREDICTED: splicing factor 45 [Drosophila miranda]
>gi|1036901988|ref|XP_017153587.1| PREDICTED: splicing factor 45 [Drosophila miranda]
>gi|1036886106|ref|XP_017152692.1| PREDICTED: protein-associating with the carboxyl-terminal do
>gi|1036886087|ref|XP_017152691.1| PREDICTED: CCR4-NOT transcription complex subunit 3 [Drosop
>gi|1036886159|ref|XP_017152696.1| PREDICTED: transmembrane protein 53 [Drosophila miranda]
>gi|1036886142|ref|XP_017152694.1| PREDICTED: probable cytosolic Fe-S cluster assembly factor C
>gi|1036886125|ref|XP_017152693.1| PREDICTED: probable cytosolic Fe-S cluster assembly factor C
>gi|1036907210|ref|XP_017153880.1| PREDICTED: uncharacterized protein LOC108163233 [Drosophila
>gi|1036907188|ref|XP_017153879.1| PREDICTED: insulin receptor [Drosophila miranda]
>gi|1036915995|ref|XP_017154363.1| PREDICTED: LOW QUALITY PROTEIN: chymotrypsin-like protease C
>gi|1036915977|ref|XP_017154362.1| PREDICTED: LOW QUALITY PROTEIN: LIM/homeobox protein Lhx3 [D
>gi|1036916031|ref|XP_017154366.1| PREDICTED: uncharacterized protein LOC108163528 [Drosophila
>gi|1036916013|ref|XP_017154365.1| PREDICTED: alpha-methyldopa hypersensitive protein-like [Dro
>gi|1036852664|ref|XP_017150835.1| PREDICTED: LOW QUALITY PROTEIN: alpha-methyldopa hypersensit
>gi|1036857818|ref|XP_017151145.1| PREDICTED: uncharacterized protein LOC108161405 [Drosophila
>gi|1036909147|ref|XP_017153979.1| PREDICTED: protein anon-37Cs [Drosophila miranda]
>gi|1036909165|ref|XP_017153980.1| PREDICTED: aromatic-L-amino-acid decarboxylase [Drosophila m
>gi|1036909183|ref|XP_017153981.1| PREDICTED: aromatic-L-amino-acid decarboxylase [Drosophila m
>gi|1036926394|ref|XP_017154951.1| PREDICTED: protein l(2)37Cc [Drosophila miranda]
>gi|1036870554|ref|XP_017151837.1| PREDICTED: putative pre-mRNA-splicing factor ATP-dependent R
>gi|1036870589|ref|XP_017151839.1| PREDICTED: general transcription factor 3C polypeptide 5 [Dr
>gi|1036870571|ref|XP_017151838.1| PREDICTED: asparagine--tRNA ligase, cytoplasmic [Drosophila
>gi|1036870607|ref|XP_017151840.1| PREDICTED: dynactin subunit 6 [Drosophila miranda]
>gi|1036870626|ref|XP_017151841.1| PREDICTED: probable DNA-directed RNA polymerases I and III s
>gi|1036870399|ref|XP_017151828.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870417|ref|XP_017151830.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870441|ref|XP_017151831.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870460|ref|XP_017151832.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870480|ref|XP_017151833.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870501|ref|XP_017151834.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870517|ref|XP_017151835.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036870535|ref|XP_017151836.1| PREDICTED: brain tumor protein [Drosophila miranda]
>gi|1036928539|ref|XP_017155068.1| PREDICTED: keratin-associated protein 5-4-like [Drosophila m
>gi|1036928519|ref|XP_017155067.1| PREDICTED: uncharacterized protein LOC108164004 [Drosophila
>gi|1036921072|ref|XP_017154654.1| PREDICTED: separin-like [Drosophila miranda]
>gi|1036904091|ref|XP_017153703.1| PREDICTED: keratin-associated protein 5-4 [Drosophila mirand
>gi|1036904109|ref|XP_017153704.1| PREDICTED: LOW QUALITY PROTEIN: tubulin gamma-2 chain [Droso
>gi|1036904056|ref|XP_017153700.1| PREDICTED: protein pigeon isoform X1 [Drosophila miranda]
>gi|1036904074|ref|XP_017153702.1| PREDICTED: protein pigeon isoform X2 [Drosophila miranda]
>gi|1036912406|ref|XP_017154150.1| PREDICTED: LOW QUALITY PROTEIN: tyrosine-protein kinase Drl
>gi|1036935463|ref|XP_017155444.1| PREDICTED: uncharacterized protein LOC108164306 [Drosophila
>gi|1036859258|ref|XP_017151212.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036913346|ref|XP_017154207.1| PREDICTED: tyrosine-protein kinase Dnt isoform X1 [Drosophil
>gi|1036913364|ref|XP_017154208.1| PREDICTED: tyrosine-protein kinase Dnt isoform X2 [Drosophil
>gi|1036934289|ref|XP_017155375.1| PREDICTED: uncharacterized protein LOC108164252 [Drosophila
>gi|1036934271|ref|XP_017155374.1| PREDICTED: uncharacterized protein LOC108164251 [Drosophila
>gi|1036859016|ref|XP_017151197.1| PREDICTED: parkin coregulated gene protein homolog [Drosophi
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>gi|1036860691|ref|XP_017151289.1| PREDICTED: C-type lectin 1 [Drosophila miranda]
>gi|1036929828|ref|XP_017155140.1| PREDICTED: LOW QUALITY PROTEIN: hyphally regulated cell wal
>gi|1036929848|ref|XP_017155142.1| PREDICTED: LOW QUALITY PROTEIN: fibroin heavy chain [Drosop
>gi|1036913328|ref|XP_017154206.1| PREDICTED: putative glycine-rich cell wall structural prote
>gi|1036913310|ref|XP_017154205.1| PREDICTED: keratin, type I cytoskeletal 9 isoform X1 [Droso
>gi|1036913275|ref|XP_017154202.1| PREDICTED: LOW QUALITY PROTEIN: peroxisomal acyl-coenzyme A
>gi|1036913293|ref|XP_017154203.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036900429|ref|XP_017153500.1| PREDICTED: LOW QUALITY PROTEIN: leupaxin [Drosophila mirand
>gi|1036900525|ref|XP_017153505.1| PREDICTED: perlucin-like [Drosophila miranda]
>gi|1036900465|ref|XP_017153502.1| PREDICTED: uncharacterized protein LOC108162997 isoform X2
>gi|1036900483|ref|XP_017153503.1| PREDICTED: uncharacterized protein LOC108162997 isoform X3
>gi|1036900446|ref|XP_017153501.1| PREDICTED: uncharacterized protein LOC108162997 isoform X1
>gi|1036900506|ref|XP_017153504.1| PREDICTED: uncharacterized protein LOC108162997 isoform X4
>gi|1036922581|ref|XP_017154746.1| PREDICTED: LOW QUALITY PROTEIN: tRNA-splicing ligase RtcB h
>gi|1036907009|ref|XP_017153868.1| PREDICTED: ras-related protein Rab-9B isoform X1 [Drosophila
>gi|1036907027|ref|XP_017153869.1| PREDICTED: ras-related protein Rab-9B isoform X2 [Drosophila
>gi|1036906962|ref|XP_017153865.1| PREDICTED: alpha-tocopherol transfer protein-like isoform X
>gi|1036906978|ref|XP_017153866.1| PREDICTED: alpha-tocopherol transfer protein-like isoform X
>gi|1036906992|ref|XP_017153867.1| PREDICTED: alpha-tocopherol transfer protein-like isoform X
>gi|1036906946|ref|XP_017153864.1| PREDICTED: heparin sulfate O-sulfotransferase [Drosophila m
>gi|1036852683|ref|XP_017150836.1| PREDICTED: ran GTPase-activating protein-like [Drosophila m
>gi|1036890874|ref|XP_017152957.1| PREDICTED: ran GTPase-activating protein-like [Drosophila m
>gi|1036890855|ref|XP_017152956.1| PREDICTED: LOW QUALITY PROTEIN: DNA topoisomerase 2 [Drosop
>gi|1036926648|ref|XP_017154966.1| PREDICTED: alpha-tocopherol transfer protein isoform X1 [Dr
>gi|1036926666|ref|XP_017154967.1| PREDICTED: alpha-tocopherol transfer protein isoform X2 [Dr
>gi|1036926684|ref|XP_017154968.1| PREDICTED: alpha-tocopherol transfer protein isoform X2 [Dr
>gi|1036926706|ref|XP_017154970.1| PREDICTED: LOW QUALITY PROTEIN: transcription factor MafB [
>gi|1036926048|ref|XP_017154933.1| PREDICTED: LOW QUALITY PROTEIN: nucleoside diphosphate-link
>gi|1036933915|ref|XP_017155358.1| PREDICTED: serine/arginine repetitive matrix protein 1 [Dros
>gi|1036902605|ref|XP_017153620.1| PREDICTED: uncharacterized protein LOC108163059 [Drosophila
>gi|1036902588|ref|XP_017153619.1| PREDICTED: nucleoside diphosphate-linked moiety X motif 19
>gi|1036902568|ref|XP_017153618.1| PREDICTED: nucleoside diphosphate-linked moiety X motif 19
>gi|1036902551|ref|XP_017153617.1| PREDICTED: cytoplasmic tRNA 2-thiolation protein 2 [Drosoph
>gi|1036902533|ref|XP_017153616.1| PREDICTED: RNA-binding protein 26 [Drosophila miranda]
>gi|1036878132|ref|XP_017152258.1| PREDICTED: LOW QUALITY PROTEIN: DNA topoisomerase 3-alpha [
>gi|1036878037|ref|XP_017152253.1| PREDICTED: LOW QUALITY PROTEIN: rho guanine nucleotide excha
>gi|1036878055|ref|XP_017152254.1| PREDICTED: transmembrane 9 superfamily member 4 [Drosophila
>gi|1036878094|ref|XP_017152256.1| PREDICTED: LOW QUALITY PROTEIN: glucose-6-phosphate 1-epime
>gi|1036878076|ref|XP_017152255.1| PREDICTED: mitogen-activated protein kinase 14B [Drosophila
>gi|1036878114|ref|XP_017152257.1| PREDICTED: protein FRA10AC1 homolog [Drosophila miranda]
>gi|1036893221|ref|XP_017153093.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofo
>gi|1036893239|ref|XP_017153094.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofo
>gi|1036921506|ref|XP_017154679.1| PREDICTED: zinc finger BED domain-containing protein 1 [Dros
>gi|1036904702|ref|XP_017153737.1| PREDICTED: brefeldin A-inhibited guanine nucleotide-exchange
>gi|1036922521|ref|XP_017154741.1| PREDICTED: carbonic anhydrase [Drosophila miranda]
>gi|1036922493|ref|XP_017154739.1| PREDICTED: tRNA-specific adenosine deaminase 1 [Drosophila
>gi|1036922508|ref|XP_017154740.1| PREDICTED: tRNA-specific adenosine deaminase 1 [Drosophila
>gi|1036852702|ref|XP_017150838.1| PREDICTED: LOW QUALITY PROTEIN: aldehyde dehydrogenase, mit
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>gi|1036923022|ref|XP_017154774.1| PREDICTED: UPF0428 protein CG16865 [Drosophila miranda]
>gi|1036922970|ref|XP_017154771.1| PREDICTED: trypsin-7-like isoform X1 [Drosophila miranda]
>gi|1036922988|ref|XP_017154772.1| PREDICTED: trypsin-7-like isoform X1 [Drosophila miranda]
>gi|1036923006|ref|XP_017154773.1| PREDICTED: kallikrein-15-like isoform X2 [Drosophila miranda
>gi|1036904684|ref|XP_017153736.1| PREDICTED: LOW QUALITY PROTEIN: protein son of sevenless [Di
>gi|1036929241|ref|XP_017155107.1| PREDICTED: cysteine sulfinic acid decarboxylase [Drosophila
>gi|1036920496|ref|XP_017154618.1| PREDICTED: DNA polymerase subunit gamma-1, mitochondrial [Di
>gi|1036905243|ref|XP_017153766.1| PREDICTED: actin-related protein 2/3 complex subunit 1A [Dro
>gi|1036905261|ref|XP_017153767.1| PREDICTED: actin-related protein 2/3 complex subunit 1A [Dro
>gi|1036905225|ref|XP_017153764.1| PREDICTED: origin recognition complex subunit 5 isoform X2
>gi|1036905207|ref|XP_017153763.1| PREDICTED: origin recognition complex subunit 5 isoform X1
>gi|1036905281|ref|XP_017153768.1| PREDICTED: LOW QUALITY PROTEIN: DNA polymerase subunit gamma
>gi|1036905301|ref|XP_017153769.1| PREDICTED: glutamyl-tRNA(Gln) amidotransferase subunit C, m:
>gi|1036890593|ref|XP_017152941.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB3 [Dro:
>gi|1036890630|ref|XP_017152943.1| PREDICTED: probable 28S ribosomal protein S23, mitochondrial
>gi|1036890536|ref|XP_017152938.1| PREDICTED: LOW QUALITY PROTEIN: centaurin-gamma-1A [Drosophi
>gi|1036890649|ref|XP_017152945.1| PREDICTED: uncharacterized protein LOC108162632 [Drosophila
>gi|1036852594|ref|XP_017150831.1| PREDICTED: uncharacterized protein LOC108161119 [Drosophila
>gi|1036890574|ref|XP_017152940.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036890555|ref|XP_017152939.1| PREDICTED: uncharacterized protein LOC108162627 [Drosophila
>gi|1036890611|ref|XP_017152942.1| PREDICTED: uncharacterized protein LOC108162630 [Drosophila
>gi|1036903542|ref|XP_017153670.1| PREDICTED: protein SMG5 isoform X2 [Drosophila miranda]
>gi|1036903524|ref|XP_017153669.1| PREDICTED: protein SMG5 isoform X1 [Drosophila miranda]
>gi|1036903562|ref|XP_017153671.1| PREDICTED: protein SMG5 isoform X3 [Drosophila miranda]
>gi|1036903580|ref|XP_017153673.1| PREDICTED: protein SMG5 isoform X3 [Drosophila miranda]
>gi|1036921470|ref|XP_017154677.1| PREDICTED: angiotensin-converting enzyme [Drosophila miranda
>gi|1036921489|ref|XP_017154678.1| PREDICTED: angiotensin-converting enzyme [Drosophila miranda
>gi|1036860394|ref|XP_017151272.1| PREDICTED: angiotensin-converting enzyme isoform X1 [Drosoph
>gi|1036860412|ref|XP_017151273.1| PREDICTED: angiotensin-converting enzyme isoform X2 [Drosoph
>gi|1036917819|ref|XP_017154470.1| PREDICTED: angiotensin-converting enzyme-like isoform X2 [Di
>gi|1036917801|ref|XP_017154469.1| PREDICTED: angiotensin-converting enzyme-like isoform X1 [Di
>gi|1036917839|ref|XP_017154471.1| PREDICTED: angiotensin-converting enzyme-like isoform X3 [Di
>gi|1036852720|ref|XP_017150839.1| PREDICTED: angiotensin-converting enzyme-like [Drosophila mi
>gi|1036852737|ref|XP_017150840.1| PREDICTED: angiotensin-converting enzyme-like [Drosophila mi
>gi|1036861260|ref|XP_017151322.1| PREDICTED: acylphosphatase-1 [Drosophila miranda]
>gi|1036921829|ref|XP_017154699.1| PREDICTED: angiotensin-converting enzyme [Drosophila miranda
>gi|1036921867|ref|XP_017154701.1| PREDICTED: MAGE-like protein 2 isoform X2 [Drosophila mirand
>gi|1036921847|ref|XP_017154700.1| PREDICTED: MAGE-like protein 2 isoform X1 [Drosophila mirand
>gi|1036921921|ref|XP_017154704.1| PREDICTED: uncharacterized protein LOC108163748 [Drosophila
>gi|1036921887|ref|XP_017154702.1| PREDICTED: chorion protein S38 [Drosophila miranda]
>gi|1036921905|ref|XP_017154703.1| PREDICTED: zinc finger protein 512B [Drosophila miranda]
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>gi|1036932102|ref|XP_017155256.1| PREDICTED: von Willebrand factor D and EGF domain-containing
>gi|1036923258|ref|XP_017154788.1| PREDICTED: LOW QUALITY PROTEIN: tenascin [Drosophila miranda
>gi|1036923310|ref|XP_017154791.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036923240|ref|XP_017154787.1| PREDICTED: von Willebrand factor D and EGF domain-containing
>gi|1036923292|ref|XP_017154790.1| PREDICTED: cell death abnormality protein 1 isoform X2 [Dro:
>gi|1036923274|ref|XP_017154789.1| PREDICTED: cell death abnormality protein 1 isoform X1 [Dro:
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>gi|1036923222|ref|XP_017154786.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036879314|ref|XP_017152320.1| PREDICTED: LOW QUALITY PROTEIN: multiple epidermal growth fa
>gi|1036879483|ref|XP_017152330.1| PREDICTED: leucine-rich repeat-containing G-protein coupled
>gi|1036879350|ref|XP_017152322.1| PREDICTED: LOW QUALITY PROTEIN: very long-chain-fatty-acid--
>gi|1036879464|ref|XP_017152329.1| PREDICTED: uncharacterized protein LOC108162220 [Drosophila
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>gi|1036879369|ref|XP_017152323.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036879421|ref|XP_017152327.1| PREDICTED: uncharacterized protein LOC108162219 isoform X2
>gi|1036879403|ref|XP_017152325.1| PREDICTED: uncharacterized protein LOC108162219 isoform X1
>gi|1036879440|ref|XP_017152328.1| PREDICTED: uncharacterized protein LOC108162219 isoform X3
>gi|1036879388|ref|XP_017152324.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036879295|ref|XP_017152319.1| PREDICTED: neuronal acetylcholine receptor subunit alpha-7 i
>gi|1036879238|ref|XP_017152316.1| PREDICTED: neuronal acetylcholine receptor subunit beta-2 is
>gi|1036879257|ref|XP_017152317.1| PREDICTED: neuronal acetylcholine receptor subunit beta-2 is
>gi|1036879276|ref|XP_017152318.1| PREDICTED: neuronal acetylcholine receptor subunit beta-2 is
>gi|1036852753|ref|XP_017150841.1| PREDICTED: BCL-6 corepressor-like protein 1 [Drosophila mira
>gi|1036928497|ref|XP_017155066.1| PREDICTED: leucine-rich repeat extensin-like protein 5 [Dros
>gi|1036928479|ref|XP_017155064.1| PREDICTED: LOW QUALITY PROTEIN: homeotic protein female ste
>gi|1036857374|ref|XP_017151122.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036932918|ref|XP_017155303.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036895497|ref|XP_017153221.1| PREDICTED: LOW QUALITY PROTEIN: dual specificity tyrosine-ph
>gi|1036893941|ref|XP_017153131.1| PREDICTED: laminin subunit alpha-1 isoform X1 [Drosophila mi
>gi|1036893956|ref|XP_017153132.1| PREDICTED: laminin subunit alpha-1 isoform X2 [Drosophila mi
>gi|1036893995|ref|XP_017153135.1| PREDICTED: uncharacterized protein LOC108162761 [Drosophila
>gi|1036893976|ref|XP_017153133.1| PREDICTED: laminin subunit alpha-1 isoform X3 [Drosophila mi
>gi|1036852772|ref|XP_017150842.1| PREDICTED: uncharacterized protein LOC108161131 [Drosophila
>gi|1036852790|ref|XP_017150843.1| PREDICTED: nardilysin-like [Drosophila miranda]
>gi|1036864385|ref|XP_017151489.1| PREDICTED: insulin-degrading enzyme-like [Drosophila miranda
>gi|1036864025|ref|XP_017151468.1| PREDICTED: non-lysosomal glucosylceramidase isoform X1 [Dros
>gi|1036864043|ref|XP_017151469.1| PREDICTED: non-lysosomal glucosylceramidase isoform X2 [Dros
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>gi|1036864169|ref|XP_017151476.1| PREDICTED: uncharacterized protein LOC108161680 [Drosophila
>gi|1036864315|ref|XP_017151485.1| PREDICTED: defense protein l(2)34Fc [Drosophila miranda]
>gi|1036864350|ref|XP_017151486.1| PREDICTED: uncharacterized protein LOC108161685 [Drosophila
>gi|1036864081|ref|XP_017151471.1| PREDICTED: nucleolar protein 10 [Drosophila miranda]
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>gi|1036864239|ref|XP_017151481.1| PREDICTED: ras-related protein Rab-14 isoform X2 [Drosophila
>gi|1036864188|ref|XP_017151477.1| PREDICTED: odorant receptor 13a isoform X1 [Drosophila miran
>gi|1036864203|ref|XP_017151478.1| PREDICTED: odorant receptor 13a isoform X2 [Drosophila miran
>gi|1036864135|ref|XP_017151474.1| PREDICTED: uncharacterized protein LOC108161677 isoform X2
>gi|1036864117|ref|XP_017151473.1| PREDICTED: uncharacterized protein LOC108161677 isoform X1
>gi|1036864099|ref|XP_017151472.1| PREDICTED: polypeptide N-acetylgalactosaminyltransferase 35A
>gi|1036864061|ref|XP_017151470.1| PREDICTED: DNA mismatch repair protein spellchecker 1 [Dros
>gi|1036864006|ref|XP_017151466.1| PREDICTED: voltage-dependent calcium channel subunit alpha-2
>gi|1036863988|ref|XP_017151465.1| PREDICTED: voltage-dependent calcium channel subunit alpha-2
>gi|1036864260|ref|XP_017151482.1| PREDICTED: uncharacterized protein LOC108161683 [Drosophila
>gi|1036864277|ref|XP_017151483.1| PREDICTED: uncharacterized protein LOC108161683 [Drosophila
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>gi|1036864296|ref|XP_017151484.1| PREDICTED: uncharacterized protein LOC108161683 [Drosophila
>gi|1036864423|ref|XP_017151491.1| PREDICTED: accessory gland-specific peptide 70A [Drosophila
>gi|1036852806|ref|XP_017150844.1| PREDICTED: uncharacterized protein LOC108161133 [Drosophila
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>gi|1036863970|ref|XP_017151464.1| PREDICTED: voltage-dependent calcium channel type D subunit
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>gi|1036925555|ref|XP_017154902.1| PREDICTED: uncharacterized protein LOC108163893 isoform X2
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>gi|1036925516|ref|XP_017154900.1| PREDICTED: EMI domain-containing protein 1-like [Drosophila
>gi|1036925574|ref|XP_017154903.1| PREDICTED: uncharacterized protein LOC108163894 [Drosophila
>gi|1036920335|ref|XP_017154608.1| PREDICTED: protein bicaudal C isoform X1 [Drosophila miranda
>gi|1036920353|ref|XP_017154609.1| PREDICTED: protein bicaudal C isoform X2 [Drosophila miranda
>gi|1036920371|ref|XP_017154610.1| PREDICTED: uncharacterized protein LOC108163690 [Drosophila
>gi|1036925930|ref|XP_017154927.1| PREDICTED: uncharacterized protein LOC108163910 [Drosophila
>gi|1036925981|ref|XP_017154931.1| PREDICTED: male-specific sperm protein Mst84Dc [Drosophila
>gi|1036925963|ref|XP_017154929.1| PREDICTED: uncharacterized protein LOC108163912 [Drosophila
>gi|1036925997|ref|XP_017154932.1| PREDICTED: keratin-associated protein 17-1 [Drosophila mira
>gi|1036925947|ref|XP_017154928.1| PREDICTED: keratin-associated protein 5-3 [Drosophila miran
>gi|1036860657|ref|XP_017151286.1| PREDICTED: uncharacterized protein LOC108161520 [Drosophila
>gi|1036930026|ref|XP_017155148.1| PREDICTED: borealin-like [Drosophila miranda]
>gi|1036930008|ref|XP_017155147.1| PREDICTED: protein CDV3 homolog [Drosophila miranda]
>gi|1036852843|ref|XP_017150846.1| PREDICTED: uncharacterized protein LOC108161136 [Drosophila
>gi|1036861373|ref|XP_017151329.1| PREDICTED: dynein light chain 1, axonemal [Drosophila mirand
>gi|1036852862|ref|XP_017150847.1| PREDICTED: uncharacterized protein LOC108161137 [Drosophila
>gi|1036852880|ref|XP_017150849.1| PREDICTED: retinoid-inducible serine carboxypeptidase-like
>gi|1036860432|ref|XP_017151274.1| PREDICTED: serine/threonine-protein phosphatase 2A activator
>gi|1036928192|ref|XP_017155049.1| PREDICTED: uncharacterized protein LOC108163991 [Drosophila
>gi|1036928210|ref|XP_017155050.1| PREDICTED: uncharacterized protein LOC108163992 [Drosophila
>gi|1036928267|ref|XP_017155053.1| PREDICTED: uncharacterized protein LOC108163994 [Drosophila
>gi|1036928287|ref|XP_017155054.1| PREDICTED: uncharacterized protein LOC108163994 [Drosophila
>gi|1036928305|ref|XP_017155055.1| PREDICTED: uncharacterized protein LOC108163994 [Drosophila
>gi|1036928228|ref|XP_017155051.1| PREDICTED: serine protease 33-like isoform X1 [Drosophila m
>gi|1036928250|ref|XP_017155052.1| PREDICTED: serine protease 33-like isoform X2 [Drosophila m
>gi|1036858868|ref|XP_017151190.1| PREDICTED: retinoid-inducible serine carboxypeptidase [Drosc
>gi|1036860283|ref|XP_017151269.1| PREDICTED: retinoid-inducible serine carboxypeptidase [Drosc
>gi|1036930076|ref|XP_017155150.1| PREDICTED: uncharacterized protein LOC108164077 isoform X1
>gi|1036930096|ref|XP_017155151.1| PREDICTED: uncharacterized protein LOC108164077 isoform X2
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>gi|1036930114|ref|XP_017155152.1| PREDICTED: inactive serine protease 54-like [Drosophila mira
>gi|1036930132|ref|XP_017155153.1| PREDICTED: inactive serine protease 54-like [Drosophila mira
>gi|1036927310|ref|XP_017155003.1| PREDICTED: thyrotropin-releasing hormone-degrading ectoenzy
>gi|1036934340|ref|XP_017155379.1| PREDICTED: tektin-4 [Drosophila miranda]
>gi|1036935512|ref|XP_017155447.1| PREDICTED: uncharacterized protein LOC108164310 [Drosophila
>gi|1036857689|ref|XP_017151138.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036861317|ref|XP_017151326.1| PREDICTED: uncharacterized protein LOC108161554 [Drosophila
>gi|1036852898|ref|XP_017150850.1| PREDICTED: uncharacterized protein LOC108161139 [Drosophila
>gi|1036934912|ref|XP_017155413.1| PREDICTED: uncharacterized protein LOC108164278 [Drosophila
>gi|1036852915|ref|XP_017150851.1| PREDICTED: serine/arginine repetitive matrix protein 2-like
>gi|1036912759|ref|XP_017154171.1| PREDICTED: uncharacterized protein LOC108163415 [Drosophila
>gi|1036912795|ref|XP_017154173.1| PREDICTED: leucine carboxyl methyltransferase 1 isoform X2
>gi|1036912777|ref|XP_017154172.1| PREDICTED: leucine carboxyl methyltransferase 1 isoform X1
>gi|1036889132|ref|XP_017152859.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1036889148|ref|XP_017152860.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1036889167|ref|XP_017152861.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1036889186|ref|XP_017152862.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1036889205|ref|XP_017152863.1| PREDICTED: acetylcholinesterase [Drosophila miranda]
>gi|1036889222|ref|XP_017152864.1| PREDICTED: P protein [Drosophila miranda]
>gi|1036889113|ref|XP_017152858.1| PREDICTED: superkiller viralicidic activity 2-like 2 [Drosp
>gi|1036929207|ref|XP_017155104.1| PREDICTED: uncharacterized protein LOC108164038 [Drosophila
>gi|1036893497|ref|XP_017153111.1| PREDICTED: G1/S-specific cyclin-E [Drosophila miranda]
>gi|1036931076|ref|XP_017155206.1| PREDICTED: uncharacterized protein LOC108164119 [Drosophila
>gi|1036924754|ref|XP_017154862.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1036924772|ref|XP_017154863.1| PREDICTED: voltage-dependent calcium channel subunit alpha-
>gi|1036924841|ref|XP_017154868.1| PREDICTED: uncharacterized protein LOC108163862 [Drosophila
>gi|1036924804|ref|XP_017154865.1| PREDICTED: LOW QUALITY PROTEIN: chymotrypsin-like protease
>gi|1036924788|ref|XP_017154864.1| PREDICTED: tryptase beta-2-like [Drosophila miranda]
>gi|1036924824|ref|XP_017154867.1| PREDICTED: uncharacterized protein LOC108163861 [Drosophila
>gi|1036916454|ref|XP_017154392.1| PREDICTED: inhibin beta A chain [Drosophila miranda]
>gi|1036916472|ref|XP_017154393.1| PREDICTED: inhibin beta A chain [Drosophila miranda]
>gi|1036916490|ref|XP_017154394.1| PREDICTED: pyruvate kinase [Drosophila miranda]
>gi|1036916508|ref|XP_017154395.1| PREDICTED: glucose-6-phosphatase 2 isoform X1 [Drosophila m
>gi|1036916539|ref|XP_017154397.1| PREDICTED: glucose-6-phosphatase 2 isoform X2 [Drosophila m
>gi|1036901703|ref|XP_017153571.1| PREDICTED: dual oxidase [Drosophila miranda]
>gi|1036901742|ref|XP_017153573.1| PREDICTED: dual oxidase [Drosophila miranda]
>gi|1036901760|ref|XP_017153574.1| PREDICTED: dual oxidase [Drosophila miranda]
>gi|1036901778|ref|XP_017153575.1| PREDICTED: dual oxidase [Drosophila miranda]
>gi|1036901798|ref|XP_017153576.1| PREDICTED: dual oxidase [Drosophila miranda]
>gi|1036932883|ref|XP_017155301.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galact
>gi|1036932900|ref|XP_017155302.1| PREDICTED: cuticle protein 18.6 [Drosophila miranda]
>gi|1036931513|ref|XP_017155227.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galact
>gi|1036931528|ref|XP_017155228.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galact
>gi|1036931495|ref|XP_017155226.1| PREDICTED: uncharacterized protein LOC108164136 [Drosophila
>gi|1036852932|ref|XP_017150852.1| PREDICTED: serine protease 42-like, partial [Drosophila mir
>gi|1036852948|ref|XP_017150853.1| PREDICTED: serine protease 42-like [Drosophila miranda]
>gi|1036935373|ref|XP_017155439.1| PREDICTED: protein phosphatase 1B [Drosophila miranda]
>gi|1036852966|ref|XP_017150854.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galact
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>gi|1036933620|ref|XP_017155343.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036909694|ref|XP_017154012.1| PREDICTED: ankyrin repeat domain-containing protein 29 [Dros
>gi|1036909676|ref|XP_017154011.1| PREDICTED: uncharacterized protein LOC108163306 isoform X3
>gi|1036909638|ref|XP_017154009.1| PREDICTED: uncharacterized protein LOC108163306 isoform X1
>gi|1036909656|ref|XP_017154010.1| PREDICTED: uncharacterized protein LOC108163306 isoform X2
>gi|1036909712|ref|XP_017154013.1| PREDICTED: 40S ribosomal protein S21 [Drosophila miranda]
>gi|1036890184|ref|XP_017152920.1| PREDICTED: uncharacterized protein LOC108162613 [Drosophila
>gi|1036890203|ref|XP_017152921.1| PREDICTED: uncharacterized protein LOC108162613 [Drosophila
>gi|1036890220|ref|XP_017152922.1| PREDICTED: interferon-related developmental regulator 2 [Dro
>gi|1036890238|ref|XP_017152923.1| PREDICTED: uncharacterized protein LOC108162616 [Drosophila
>gi|1036896925|ref|XP_017153299.1| PREDICTED: acyl-CoA-binding domain-containing protein 5 [Dro
>gi|1036896942|ref|XP_017153300.1| PREDICTED: peroxiredoxin-6 [Drosophila miranda]
>gi|1036896959|ref|XP_017153301.1| PREDICTED: N-alpha-acetyltransferase 38, NatC auxiliary subu
>gi|1036896978|ref|XP_017153302.1| PREDICTED: uncharacterized protein LOC108162865 [Drosophila
>gi|1036896906|ref|XP_017153298.1| PREDICTED: geranylgeranyl transferase type-2 subunit beta [D
>gi|1036917563|ref|XP_017154454.1| PREDICTED: ectonucleoside triphosphate diphosphohydrolase 5
>gi|1036917581|ref|XP_017154455.1| PREDICTED: ectonucleoside triphosphate diphosphohydrolase 5
>gi|1036917599|ref|XP_017154456.1| PREDICTED: ectonucleoside triphosphate diphosphohydrolase 5
>gi|1036935842|ref|XP_017155468.1| PREDICTED: uncharacterized protein LOC108164326 [Drosophila
>gi|1036935860|ref|XP_017155470.1| PREDICTED: uncharacterized protein LOC108164326 [Drosophila
>gi|1036867808|ref|XP_017151685.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila n
>gi|1036867827|ref|XP_017151686.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila n
>gi|1036867842|ref|XP_017151687.1| PREDICTED: AF4/FMR2 family member 4 isoform X3 [Drosophila n
>gi|1036867857|ref|XP_017151688.1| PREDICTED: AF4/FMR2 family member 4 isoform X3 [Drosophila n
>gi|1036852984|ref|XP_017150855.1| PREDICTED: uncharacterized protein LOC108161145 [Drosophila
>gi|1036867873|ref|XP_017151689.1| PREDICTED: LOW QUALITY PROTEIN: ELAV-like protein 2 [Drosoph
>gi|1036867900|ref|XP_017151690.1| PREDICTED: thymidylate synthase [Drosophila miranda]
>gi|1036905803|ref|XP_017153799.1| PREDICTED: recombination repair protein 1 isoform X2 [Droso
>gi|1036905789|ref|XP_017153798.1| PREDICTED: recombination repair protein 1 isoform X1 [Droso
>gi|1036905847|ref|XP_017153803.1| PREDICTED: tubulin gamma-1 chain [Drosophila miranda]
>gi|1036905816|ref|XP_017153801.1| PREDICTED: protein Hook homolog 1-like isoform X1 [Drosophil
>gi|1036905830|ref|XP_017153802.1| PREDICTED: protein Hook homolog 1-like isoform X2 [Drosophil
>gi|1036919203|ref|XP_017154543.1| PREDICTED: three-prime repair exonuclease 1 [Drosophila mira
>gi|1036919220|ref|XP_017154544.1| PREDICTED: protein-lysine N-methyltransferase Mettl10 [Dros
>gi|1036895909|ref|XP_017153245.1| PREDICTED: chromodomain-helicase-DNA-binding protein 1 isofo
>gi|1036895890|ref|XP_017153244.1| PREDICTED: chromodomain-helicase-DNA-binding protein 1 isofo
>gi|1036891079|ref|XP_017152969.1| PREDICTED: protein ABHD13 [Drosophila miranda]
>gi|1036891096|ref|XP_017152970.1| PREDICTED: protein ABHD13 [Drosophila miranda]
>gi|1036891039|ref|XP_017152967.1| PREDICTED: DNA repair and recombination protein RAD54-like
>gi|1036891060|ref|XP_017152968.1| PREDICTED: DNA repair and recombination protein RAD54-like
>gi|1036891022|ref|XP_017152966.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036908756|ref|XP_017153960.1| PREDICTED: uncharacterized protein LOC108163270 [Drosophila
>gi|1036915032|ref|XP_017154301.1| PREDICTED: ethanolamine-phosphate cytidylyltransferase isofo
>gi|1036915050|ref|XP_017154302.1| PREDICTED: ethanolamine-phosphate cytidylyltransferase isofo
>gi|1036915066|ref|XP_017154305.1| PREDICTED: uncharacterized protein LOC108163494 [Drosophila
>gi|1036888364|ref|XP_017152813.1| PREDICTED: actin-binding protein IPP [Drosophila miranda]
>gi|1036888417|ref|XP_017152817.1| PREDICTED: peptide chain release factor 1-like, mitochondria
>gi|1036888453|ref|XP_017152819.1| PREDICTED: exosome complex component RRP41 isoform X2 [Droso
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>gi|1036888435|ref|XP_017152818.1| PREDICTED: exosome complex component RRP41 isoform X1 [Dros
>gi|1036888399|ref|XP_017152815.1| PREDICTED: uncharacterized protein C19orf47 homolog isoform
>gi|1036888383|ref|XP_017152814.1| PREDICTED: uncharacterized protein C19orf47 homolog isoform
>gi|1036888471|ref|XP_017152820.1| PREDICTED: forkhead box protein B2 [Drosophila miranda]
>gi|1036857452|ref|XP_017151125.1| PREDICTED: uncharacterized protein LOC108161387 [Drosophila
>gi|1036857291|ref|XP_017151116.1| PREDICTED: uncharacterized protein LOC108161380 [Drosophila
>gi|1036853003|ref|XP_017150857.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop
>gi|1036924646|ref|XP_017154855.1| PREDICTED: proteasome subunit alpha type-1 [Drosophila mira
>gi|1036924682|ref|XP_017154857.1| PREDICTED: uncharacterized protein LOC108163854 [Drosophila
>gi|1036924664|ref|XP_017154856.1| PREDICTED: uncharacterized protein LOC108163853 [Drosophila
>gi|1036933466|ref|XP_017155336.1| PREDICTED: uncharacterized protein LOC108164214 [Drosophila
>gi|1036931479|ref|XP_017155225.1| PREDICTED: uncharacterized protein LOC108164135 [Drosophila
>gi|1036935391|ref|XP_017155440.1| PREDICTED: fibrinogen-like protein 1 [Drosophila miranda]
>gi|1036931285|ref|XP_017155214.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1036853022|ref|XP_017150858.1| PREDICTED: N-alpha-acetyltransferase 20 [Drosophila miranda]
>gi|1036860619|ref|XP_017151284.1| PREDICTED: N-alpha-acetyltransferase 20 [Drosophila miranda]
>gi|1036905603|ref|XP_017153786.1| PREDICTED: transcription factor SPT20 homolog isoform X6 [Di
>gi|1036905693|ref|XP_017153792.1| PREDICTED: uncharacterized protein LOC108163171 [Drosophila
>gi|1036905512|ref|XP_017153781.1| PREDICTED: transcription factor SPT20 homolog isoform X1 [Di
>gi|1036905530|ref|XP_017153782.1| PREDICTED: transcription factor SPT20 homolog isoform X2 [Di
>gi|1036905568|ref|XP_017153784.1| PREDICTED: transcription factor SPT20 homolog isoform X4 [Di
>gi|1036905586|ref|XP_017153785.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X5
>gi|1036905550|ref|XP_017153783.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036905620|ref|XP_017153787.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036905654|ref|XP_017153789.1| PREDICTED: trithorax group protein osa isoform X9 [Drosophi
>gi|1036905637|ref|XP_017153788.1| PREDICTED: uncharacterized protein LOC108163167 isoform X8
>gi|1036905675|ref|XP_017153790.1| PREDICTED: DNA fragmentation factor subunit beta [Drosophil
>gi|1036882641|ref|XP_017152499.1| PREDICTED: protein sickie isoform X3 [Drosophila miranda]
>gi|1036882821|ref|XP_017152509.1| PREDICTED: collectin-10-like [Drosophila miranda]
>gi|1036882802|ref|XP_017152508.1| PREDICTED: tetranectin-like protein [Drosophila miranda]
>gi|1036882587|ref|XP_017152497.1| PREDICTED: protein sickie isoform X1 [Drosophila miranda]
>gi|1036882840|ref|XP_017152511.1| PREDICTED: male-specific sperm protein Mst84Da-like [Drosop
>gi|1036882606|ref|XP_017152498.1| PREDICTED: protein sickie isoform X2 [Drosophila miranda]
>gi|1036882659|ref|XP_017152500.1| PREDICTED: protein sickie isoform X4 [Drosophila miranda]
>gi|1036882702|ref|XP_017152502.1| PREDICTED: protein sickie isoform X6 [Drosophila miranda]
>gi|1036882786|ref|XP_017152507.1| PREDICTED: actin-5C-like [Drosophila miranda]
>gi|1036882769|ref|XP_017152506.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036882684|ref|XP_017152501.1| PREDICTED: protein sickie isoform X5 [Drosophila miranda]
>gi|1036882731|ref|XP_017152504.1| PREDICTED: protein sickie isoform X7 [Drosophila miranda]
>gi|1036882750|ref|XP_017152505.1| PREDICTED: protein sickie isoform X7 [Drosophila miranda]
>gi|1036860320|ref|XP_017151270.1| PREDICTED: uncharacterized protein LOC108161505 [Drosophila
>gi|1036912709|ref|XP_017154168.1| PREDICTED: cytochrome c oxidase subunit 4 isoform 1, mitoch
>gi|1036912655|ref|XP_017154165.1| PREDICTED: uncharacterized protein LOC108163410 [Drosophila
>gi|1036912579|ref|XP_017154160.1| PREDICTED: protein PF14_0175-like [Drosophila miranda]
>gi|1036912597|ref|XP_017154161.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036912617|ref|XP_017154163.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X1
>gi|1036912637|ref|XP_017154164.1| PREDICTED: uncharacterized protein LOC108163409 isoform X2
>gi|1036912691|ref|XP_017154167.1| PREDICTED: uncharacterized protein LOC108163412 [Drosophila
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>gi|1036912739|ref|XP_017154169.1| PREDICTED: uncharacterized protein LOC108163414 [Drosophila
>gi|1036912673|ref|XP_017154166.1| PREDICTED: uncharacterized protein LOC108163411 [Drosophila
>gi|1036903655|ref|XP_017153676.1| PREDICTED: uncharacterized protein LOC108163099 [Drosophila
>gi|1036903637|ref|XP_017153675.1| PREDICTED: LOW QUALITY PROTEIN: condensin complex subunit 2-
>gi|1036903673|ref|XP_017153677.1| PREDICTED: uncharacterized protein LOC108163100 [Drosophila
>gi|1036903818|ref|XP_017153686.1| PREDICTED: uncharacterized protein LOC108163103 [Drosophila
>gi|1036903711|ref|XP_017153679.1| PREDICTED: condensin complex subunit 2-like isoform X2 [Dros
>gi|1036903782|ref|XP_017153683.1| PREDICTED: condensin complex subunit 2-like isoform X2 [Dros
>gi|1036903798|ref|XP_017153685.1| PREDICTED: condensin complex subunit 2-like isoform X2 [Dros
>gi|1036903691|ref|XP_017153678.1| PREDICTED: condensin complex subunit 2-like isoform X1 [Dros
>gi|1036903729|ref|XP_017153680.1| PREDICTED: condensin complex subunit 2-like isoform X1 [Dros
>gi|1036903747|ref|XP_017153681.1| PREDICTED: condensin complex subunit 2-like isoform X1 [Dros
>gi|1036903765|ref|XP_017153682.1| PREDICTED: condensin complex subunit 2-like isoform X1 [Dros
>gi|1036853041|ref|XP_017150859.1| PREDICTED: condensin complex subunit 2-like, partial [Drosop
>gi|1036857653|ref|XP_017151136.1| PREDICTED: short neuropeptide F [Drosophila miranda]
>gi|1036893620|ref|XP_017153112.1| PREDICTED: condensin complex subunit 2-like [Drosophila mira
>gi|1036893636|ref|XP_017153114.1| PREDICTED: ovarian-specific serine/threonine-protein kinase
>gi|1036893655|ref|XP_017153115.1| PREDICTED: ovarian-specific serine/threonine-protein kinase
>gi|1036893672|ref|XP_017153116.1| PREDICTED: protein valois [Drosophila miranda]
>gi|1036893687|ref|XP_017153117.1| PREDICTED: uncharacterized protein LOC108162749 [Drosophila
>gi|1036906273|ref|XP_017153830.1| PREDICTED: alkaline ceramidase [Drosophila miranda]
>gi|1036906259|ref|XP_017153829.1| PREDICTED: uncharacterized protein LOC108163195 [Drosophila
>gi|1036906289|ref|XP_017153831.1| PREDICTED: 6-pyruvoyl tetrahydrobiopterin synthase [Drosoph
>gi|1036906307|ref|XP_017153832.1| PREDICTED: 6-pyruvoyl tetrahydrobiopterin synthase [Drosoph
>gi|1036872798|ref|XP_017151968.1| PREDICTED: kinesin-like protein KIF14 isoform X1 [Drosophila
>gi|1036873047|ref|XP_017151983.1| PREDICTED: uncharacterized protein LOC108162007 isoform X3
>gi|1036873028|ref|XP_017151982.1| PREDICTED: uncharacterized protein LOC108162007 isoform X2
>gi|1036873003|ref|XP_017151981.1| PREDICTED: uncharacterized protein LOC108162007 isoform X1
>gi|1036872815|ref|XP_017151969.1| PREDICTED: kinesin-like protein KIF14 isoform X2 [Drosophila
>gi|1036872870|ref|XP_017151973.1| PREDICTED: PI-PLC X domain-containing protein 2 isoform X1
>gi|1036872888|ref|XP_017151974.1| PREDICTED: PI-PLC X domain-containing protein 2 isoform X2
>gi|1036872834|ref|XP_017151970.1| PREDICTED: SHC SH2 domain-binding protein 1 [Drosophila mira
>gi|1036872982|ref|XP_017151979.1| PREDICTED: transcription initiation factor TFIID subunit 13
>gi|1036872852|ref|XP_017151972.1| PREDICTED: pyridine nucleotide-disulfide oxidoreductase doma
>gi|1036872964|ref|XP_017151978.1| PREDICTED: 28S ribosomal protein S18b, mitochondrial [Drosop
>gi|1036872907|ref|XP_017151975.1| PREDICTED: transmembrane protein 189 [Drosophila miranda]
>gi|1036872927|ref|XP_017151976.1| PREDICTED: transmembrane protein 189 [Drosophila miranda]
>gi|1036872945|ref|XP_017151977.1| PREDICTED: transmembrane protein 189 [Drosophila miranda]
>gi|1036859369|ref|XP_017151218.1| PREDICTED: protein SPT2 homolog [Drosophila miranda]
>gi|1036859350|ref|XP_017151217.1| PREDICTED: uncharacterized protein LOC108161464 [Drosophila
>gi|1036904317|ref|XP_017153717.1| PREDICTED: intraflagellar transport protein 88 homolog [Dros
>gi|1036904357|ref|XP_017153720.1| PREDICTED: lysosome-associated membrane glycoprotein 1 [Dros
>gi|1036904412|ref|XP_017153723.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036904375|ref|XP_017153721.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036904394|ref|XP_017153722.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036904299|ref|XP_017153716.1| PREDICTED: cytosolic 10-formyltetrahydrofolate dehydrogenase
>gi|1036904338|ref|XP_017153718.1| PREDICTED: putative gustatory receptor 39b [Drosophila miran
>gi|1036902747|ref|XP_017153629.1| PREDICTED: dynein assembly factor 1, axonemal homolog [Droso
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>gi|1036902765|ref|XP_017153630.1| PREDICTED: CD63 antigen [Drosophila miranda]
>gi|1036902785|ref|XP_017153631.1| PREDICTED: CD63 antigen [Drosophila miranda]
>gi|1036859801|ref|XP_017151241.1| PREDICTED: protein dimmed [Drosophila miranda]
>gi|1036905391|ref|XP_017153774.1| PREDICTED: uncharacterized protein LOC108163164 isoform X1
>gi|1036905408|ref|XP_017153775.1| PREDICTED: activating transcription factor of chaperone iso
>gi|1036899313|ref|XP_017153435.1| PREDICTED: carbohydrate-responsive element-binding protein
>gi|1036899332|ref|XP_017153436.1| PREDICTED: carbohydrate-responsive element-binding protein
>gi|1036899350|ref|XP_017153437.1| PREDICTED: carbohydrate-responsive element-binding protein
>gi|1036852612|ref|XP_017150832.1| PREDICTED: gustatory and pheromone receptor 39a-like, partia
>gi|1036899371|ref|XP_017153439.1| PREDICTED: gustatory and pheromone receptor 39a-like [Droso
>gi|1036899390|ref|XP_017153440.1| PREDICTED: gustatory and pheromone receptor 39a-like [Droso
>gi|1036883865|ref|XP_017152573.1| PREDICTED: probable cytochrome P450 6t1 [Drosophila miranda]
>gi|1036883828|ref|XP_017152571.1| PREDICTED: protein FAM102A isoform X4 [Drosophila miranda]
>gi|1036883810|ref|XP_017152570.1| PREDICTED: uncharacterized protein LOC108162369 isoform X3
>gi|1036883792|ref|XP_017152568.1| PREDICTED: uncharacterized protein LOC108162369 isoform X2
>gi|1036883757|ref|XP_017152566.1| PREDICTED: uncharacterized protein LOC108162369 isoform X1
>gi|1036883774|ref|XP_017152567.1| PREDICTED: uncharacterized protein LOC108162369 isoform X1
>gi|1036883846|ref|XP_017152572.1| PREDICTED: uncharacterized protein LOC108162369 isoform X5
>gi|1036883884|ref|XP_017152574.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila r
>gi|1036883922|ref|XP_017152576.1| PREDICTED: ATP synthase mitochondrial F1 complex assembly fa
>gi|1036883903|ref|XP_017152575.1| PREDICTED: uncharacterized protein LOC108162372 [Drosophila
>gi|1036915420|ref|XP_017154328.1| PREDICTED: nuclear hormone receptor FTZ-F1 beta [Drosophila
>gi|1036885996|ref|XP_017152685.1| PREDICTED: extensin isoform X1 [Drosophila miranda]
>gi|1036886015|ref|XP_017152686.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1036885977|ref|XP_017152684.1| PREDICTED: microtubule-associated protein futsch [Drosophila
>gi|1036903357|ref|XP_017153660.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036903375|ref|XP_017153661.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036903303|ref|XP_017153657.1| PREDICTED: uncharacterized protein LOC108163087 [Drosophila
>gi|1036903319|ref|XP_017153658.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036903337|ref|XP_017153659.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036903266|ref|XP_017153654.1| PREDICTED: glutamate receptor ionotropic, kainate 2 [Drosopl
>gi|1036903431|ref|XP_017153665.1| PREDICTED: uncharacterized protein C4orf22 homolog [Drosophi
>gi|1036903395|ref|XP_017153662.1| PREDICTED: uncharacterized protein C4orf22 homolog isoform }
>gi|1036903413|ref|XP_017153663.1| PREDICTED: uncharacterized protein C4orf22 homolog isoform }
>gi|1036903284|ref|XP_017153656.1| PREDICTED: glutamate receptor ionotropic, kainate 2 [Drosopl
>gi|1036934073|ref|XP_017155368.1| PREDICTED: glutamate receptor ionotropic, kainate 2 [Drosopl
>gi|1036857799|ref|XP_017151144.1| PREDICTED: uncharacterized protein LOC108161404 [Drosophila
>gi|1036933947|ref|XP_017155360.1| PREDICTED: uncharacterized serine-rich protein C215.13-like
>gi|1036933965|ref|XP_017155361.1| PREDICTED: uncharacterized serine-rich protein C215.13-like
>gi|1036934357|ref|XP_017155380.1| PREDICTED: stress response protein NST1 [Drosophila miranda]
>gi|1036934375|ref|XP_017155381.1| PREDICTED: stress response protein NST1 [Drosophila miranda]
>gi|1036934391|ref|XP_017155382.1| PREDICTED: stress response protein NST1 [Drosophila miranda]
>gi|1036932705|ref|XP_017155290.1| PREDICTED: limbic system-associated membrane protein [Drosop
>gi|1036918936|ref|XP_017154532.1| PREDICTED: cytochrome P450 6a22 [Drosophila miranda]
>gi|1036918897|ref|XP_017154529.1| PREDICTED: uncharacterized protein LOC108163632 isoform X2
>gi|1036918862|ref|XP_017154527.1| PREDICTED: cell adhesion molecule 2 isoform X1 [Drosophila r
>gi|1036918880|ref|XP_017154528.1| PREDICTED: cell adhesion molecule 2 isoform X1 [Drosophila r
>gi|1036918958|ref|XP_017154533.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
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>gi|1036918916|ref|XP_017154531.1| PREDICTED: 60 kDa heat shock protein homolog 2, mitochondria
>gi|1036860002|ref|XP_017151252.1| PREDICTED: uncharacterized protein LOC108161490 [Drosophila
>gi|1036862079|ref|XP_017151368.1| PREDICTED: uncharacterized protein LOC108161593 [Drosophila
>gi|1036860638|ref|XP_017151285.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036861900|ref|XP_017151358.1| PREDICTED: cyclin-dependent kinase-like 1 isoform X2 [Drosop
>gi|1036861919|ref|XP_017151359.1| PREDICTED: cyclin-dependent kinase-like 1 isoform X3 [Drosop
>gi|1036861881|ref|XP_017151357.1| PREDICTED: cyclin-dependent kinase-like 1 isoform X1 [Drosop
>gi|1036861937|ref|XP_017151360.1| PREDICTED: cyclin-dependent kinase-like 1 isoform X4 [Drosop
>gi|1036865855|ref|XP_017151571.1| PREDICTED: uncharacterized protein LOC108161749 [Drosophila
>gi|1036853059|ref|XP_017150860.1| PREDICTED: uncharacterized protein LOC108161149 [Drosophila
>gi|1036917948|ref|XP_017154477.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917891|ref|XP_017154473.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917913|ref|XP_017154474.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917930|ref|XP_017154475.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917873|ref|XP_017154472.1| PREDICTED: septin-interacting protein 1-like [Drosophila mi
>gi|1036917187|ref|XP_017154430.1| PREDICTED: voltage-dependent anion-selective channel [Drosop
>gi|1036917165|ref|XP_017154429.1| PREDICTED: voltage-dependent anion-selective channel [Drosop
>gi|1036917146|ref|XP_017154428.1| PREDICTED: voltage-dependent anion-selective channel-like [
>gi|1036909622|ref|XP_017154007.1| PREDICTED: LOW QUALITY PROTEIN: leukocyte receptor cluster
>gi|1036920570|ref|XP_017154623.1| PREDICTED: LOW QUALITY PROTEIN: nitric oxide synthase [Droso
>gi|1036920592|ref|XP_017154624.1| PREDICTED: lysosomal aspartic protease-like [Drosophila mira
>gi|1036928559|ref|XP_017155069.1| PREDICTED: outer dense fiber protein 3-like protein 2 [Droso
>gi|1036928581|ref|XP_017155070.1| PREDICTED: LOW QUALITY PROTEIN: outer dense fiber protein 3-
>gi|1036928599|ref|XP_017155071.1| PREDICTED: LOW QUALITY PROTEIN: mucin-5AC [Drosophila miran
>gi|1036863455|ref|XP_017151443.1| PREDICTED: uncharacterized protein LOC108161652 [Drosophila
>gi|1036923968|ref|XP_017154819.1| PREDICTED: ribosome biogenesis protein WDR12 homolog [Drosop
>gi|1036917205|ref|XP_017154431.1| PREDICTED: ubiquitin-conjugating enzyme E2-24 kDa [Drosophil
>gi|1036917223|ref|XP_017154432.1| PREDICTED: ubiquitin-conjugating enzyme E2-24 kDa [Drosophil
>gi|1036917241|ref|XP_017154433.1| PREDICTED: ubiquitin-conjugating enzyme E2-24 kDa [Drosophil
>gi|1036917259|ref|XP_017154435.1| PREDICTED: LOW QUALITY PROTEIN: low-density lipoprotein rece
>gi|1036909093|ref|XP_017153976.1| PREDICTED: protein phosphatase 1 regulatory subunit 14B [Dro
>gi|1036909111|ref|XP_017153977.1| PREDICTED: protein phosphatase 1 regulatory subunit 14B [Dro
>gi|1036909129|ref|XP_017153978.1| PREDICTED: zinc finger protein 34 [Drosophila miranda]
>gi|1036898587|ref|XP_017153396.1| PREDICTED: LOW QUALITY PROTEIN: nuclear pore complex protei
>gi|1036898652|ref|XP_017153401.1| PREDICTED: LOW QUALITY PROTEIN: protein dpy-30 homolog [Dros
>gi|1036898633|ref|XP_017153399.1| PREDICTED: ER membrane protein complex subunit 3 [Drosophila
>gi|1036898617|ref|XP_017153398.1| PREDICTED: cilia- and flagella-associated protein 20 [Drosop
>gi|1036898602|ref|XP_017153397.1| PREDICTED: protein RTF2 homolog [Drosophila miranda]
>gi|1036931149|ref|XP_017155207.1| PREDICTED: lipase member H-A [Drosophila miranda]
>gi|1036853078|ref|XP_017150861.1| PREDICTED: perlucin-like protein [Drosophila miranda]
>gi|1036927574|ref|XP_017155021.1| PREDICTED: lipase 1 [Drosophila miranda]
>gi|1036927556|ref|XP_017155020.1| PREDICTED: aminomethyltransferase, mitochondrial [Drosophila
>gi|1036927539|ref|XP_017155019.1| PREDICTED: lipase 1 [Drosophila miranda]
>gi|1036935599|ref|XP_017155455.1| PREDICTED: uncharacterized protein LOC108164315 [Drosophila
>gi|1036935582|ref|XP_017155454.1| PREDICTED: keratin, type I cytoskeletal 9-like [Drosophila
>gi|1036908912|ref|XP_017153966.1| PREDICTED: glycine-rich cell wall structural protein [Drosop
>gi|1036908967|ref|XP_017153969.1| PREDICTED: glycine-rich cell wall structural protein 2-like
>gi|1036908932|ref|XP_017153967.1| PREDICTED: acanthoscurrin-2-like [Drosophila miranda]
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>gi|1036909006|ref|XP_017153971.1| PREDICTED: glycine-rich cell wall structural protein 2-like
>gi|1036908984|ref|XP_017153970.1| PREDICTED: alanine and glycine-rich protein-like [Drosophila
>gi|1036853097|ref|XP_017150863.1| PREDICTED: LOW QUALITY PROTEIN: mucin-5AC, partial [Drosophi
>gi|1036909024|ref|XP_017153973.1| PREDICTED: LOW QUALITY PROTEIN: lipase 3 [Drosophila miranda
>gi|1036909042|ref|XP_017153974.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036908949|ref|XP_017153968.1| PREDICTED: uncharacterized protein LOC108163278 [Drosophila
>gi|1036853117|ref|XP_017150864.1| PREDICTED: LOW QUALITY PROTEIN: lipase 3-like [Drosophila mi
>gi|1036890688|ref|XP_017152947.1| PREDICTED: uncharacterized protein LOC108162634 [Drosophila
>gi|1036890707|ref|XP_017152948.1| PREDICTED: lipase 1 [Drosophila miranda]
>gi|1036890668|ref|XP_017152946.1| PREDICTED: LOW QUALITY PROTEIN: E3 ubiquitin-protein ligase
>gi|1036890742|ref|XP_017152950.1| PREDICTED: acylphosphatase-2 [Drosophila miranda]
>gi|1036890761|ref|XP_017152951.1| PREDICTED: uncharacterized protein LOC108162638 [Drosophila
>gi|1036890724|ref|XP_017152949.1| PREDICTED: uncharacterized protein LOC108162636 [Drosophila
>gi|1036935018|ref|XP_017155419.1| PREDICTED: lipase 1 [Drosophila miranda]
>gi|1036929295|ref|XP_017155110.1| PREDICTED: LOW QUALITY PROTEIN: lipase 3-like [Drosophila mi
>gi|1036908794|ref|XP_017153962.1| PREDICTED: leucine-rich repeat protein 1 [Drosophila miranda
>gi|1036908774|ref|XP_017153961.1| PREDICTED: LOW QUALITY PROTEIN: long-chain fatty acid transp
>gi|1036897341|ref|XP_017153323.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036897358|ref|XP_017153324.1| PREDICTED: peptidyl-tRNA hydrolase ICT1, mitochondrial [Dros
>gi|1036897324|ref|XP_017153322.1| PREDICTED: LOW QUALITY PROTEIN: ATP-binding cassette sub-fam
>gi|1036901142|ref|XP_017153541.1| PREDICTED: LOW QUALITY PROTEIN: ATP-binding cassette sub-fam
>gi|1036901122|ref|XP_017153540.1| PREDICTED: LOW QUALITY PROTEIN: cell division cycle protein
>gi|1036901160|ref|XP_017153542.1| PREDICTED: LOW QUALITY PROTEIN: group XV phospholipase A2-li
>gi|1036906327|ref|XP_017153833.1| PREDICTED: LOW QUALITY PROTEIN: ras-interacting protein RIP3
>gi|1036915456|ref|XP_017154331.1| PREDICTED: transmembrane inner ear expressed protein isofor
>gi|1036915473|ref|XP_017154332.1| PREDICTED: transmembrane inner ear expressed protein isofor
>gi|1036915438|ref|XP_017154330.1| PREDICTED: actin-related protein 2/3 complex subunit 2 [Dros
>gi|1036915128|ref|XP_017154309.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036915143|ref|XP_017154310.1| PREDICTED: calmodulin-lysine N-methyltransferase [Drosophila
>gi|1036915158|ref|XP_017154311.1| PREDICTED: lymphokine-activated killer T-cell-originated pro
>gi|1036853136|ref|XP_017150865.1| PREDICTED: uncharacterized protein LOC108161154 [Drosophila
>gi|1036885345|ref|XP_017152654.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036885392|ref|XP_017152656.1| PREDICTED: LOW QUALITY PROTEIN: la protein homolog [Drosophi
>gi|1036885375|ref|XP_017152655.1| PREDICTED: LOW QUALITY PROTEIN: protein spire [Drosophila mi
>gi|1036911992|ref|XP_017154124.1| PREDICTED: protein UXT [Drosophila miranda]
>gi|1036911924|ref|XP_017154121.1| PREDICTED: cingulin isoform X2 [Drosophila miranda]
>gi|1036911906|ref|XP_017154120.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036911941|ref|XP_017154123.1| PREDICTED: synaptonemal complex protein 1 isoform X3 [Drosop
>gi|1036909075|ref|XP_017153975.1| PREDICTED: LOW QUALITY PROTEIN: cullin-3 [Drosophila miranda
>gi|1036933222|ref|XP_017155322.1| PREDICTED: ribonuclease UK114 [Drosophila miranda]
>gi|1036853154|ref|XP_017150866.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036923363|ref|XP_017154795.1| PREDICTED: uncharacterized protein LOC108163815 [Drosophila
>gi|1036923346|ref|XP_017154794.1| PREDICTED: uncharacterized protein LOC108163814 [Drosophila
>gi|1036853171|ref|XP_017150867.1| PREDICTED: uncharacterized protein LOC108161156 [Drosophila
>gi|1036934593|ref|XP_017155396.1| PREDICTED: uncharacterized protein LOC108164265 [Drosophila
>gi|1036927455|ref|XP_017155013.1| PREDICTED: protein escargot [Drosophila miranda]
>gi|1036861743|ref|XP_017151352.1| PREDICTED: uncharacterized protein LOC108161575 [Drosophila
>gi|1036933397|ref|XP_017155332.1| PREDICTED: zinc finger and SCAN domain-containing protein 12
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>gi|1036934782|ref|XP_017155406.1| PREDICTED: LOW QUALITY PROTEIN: protein-glutamate O-methylt
>gi|1036928998|ref|XP_017155094.1| PREDICTED: LOW QUALITY PROTEIN: protein snail [Drosophila mi
>gi|1036859296|ref|XP_017151214.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036859315|ref|XP_017151215.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036859334|ref|XP_017151216.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036916049|ref|XP_017154367.1| PREDICTED: LOW QUALITY PROTEIN: serine palmitoyltransferase
>gi|1036922394|ref|XP_017154732.1| PREDICTED: uncharacterized protein LOC108163768 [Drosophila
>gi|1036922376|ref|XP_017154731.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036857634|ref|XP_017151135.1| PREDICTED: LOW QUALITY PROTEIN: zinc metalloproteinase nas-4
>gi|1036935087|ref|XP_017155422.1| PREDICTED: high choriolytic enzyme 1-like [Drosophila mirand
>gi|1036858775|ref|XP_017151184.1| PREDICTED: zinc metalloproteinase nas-4-like [Drosophila mi
>gi|1036934510|ref|XP_017155390.1| PREDICTED: LOW QUALITY PROTEIN: zinc metalloproteinase nas-
>gi|1036853190|ref|XP_017150868.1| PREDICTED: LOW QUALITY PROTEIN: astacin-like metalloprotease
>gi|1036934492|ref|XP_017155389.1| PREDICTED: odorant receptor 35a [Drosophila miranda]
>gi|1036860675|ref|XP_017151288.1| PREDICTED: zinc metalloproteinase nas-4-like [Drosophila mi
>gi|1036853209|ref|XP_017150869.1| PREDICTED: LOW QUALITY PROTEIN: zinc metalloproteinase nas-4
>gi|1036933760|ref|XP_017155294.1| PREDICTED: uncharacterized protein LOC108164187 [Drosophila
>gi|1036924109|ref|XP_017154827.1| PREDICTED: synaptotagmin 1 isoform X1 [Drosophila miranda]
>gi|1036924158|ref|XP_017154828.1| PREDICTED: synaptotagmin 1 isoform X1 [Drosophila miranda]
>gi|1036924174|ref|XP_017154829.1| PREDICTED: synaptotagmin 1 isoform X1 [Drosophila miranda]
>gi|1036924192|ref|XP_017154830.1| PREDICTED: synaptotagmin 1 isoform X1 [Drosophila miranda]
>gi|1036924210|ref|XP_017154831.1| PREDICTED: synaptotagmin 1 isoform X2 [Drosophila miranda]
>gi|1036924248|ref|XP_017154833.1| PREDICTED: synaptotagmin 1 isoform X4 [Drosophila miranda]
>gi|1036924228|ref|XP_017154832.1| PREDICTED: synaptotagmin 1 isoform X3 [Drosophila miranda]
>gi|1036924266|ref|XP_017154834.1| PREDICTED: synaptotagmin 1 isoform X5 [Drosophila miranda]
>gi|1036922713|ref|XP_017154755.1| PREDICTED: rab-like protein 3 [Drosophila miranda]
>gi|1036922733|ref|XP_017154756.1| PREDICTED: rab-like protein 3 [Drosophila miranda]
>gi|1036922789|ref|XP_017154759.1| PREDICTED: histidine triad nucleotide-binding protein 1 iso
>gi|1036922771|ref|XP_017154758.1| PREDICTED: histidine triad nucleotide-binding protein 1 iso
>gi|1036922751|ref|XP_017154757.1| PREDICTED: LOW QUALITY PROTEIN: transcription initiation fac
>gi|1036865116|ref|XP_017151529.1| PREDICTED: transcription initiation factor TFIID subunit 10
>gi|1036865078|ref|XP_017151526.1| PREDICTED: gamma-secretase subunit Aph-1 [Drosophila miranda
>gi|1036865057|ref|XP_017151525.1| PREDICTED: sorting nexin-21 [Drosophila miranda]
>gi|1036865039|ref|XP_017151524.1| PREDICTED: transportin-3 [Drosophila miranda]
>gi|1036865135|ref|XP_017151530.1| PREDICTED: SNAPIN protein homolog [Drosophila miranda]
>gi|1036865097|ref|XP_017151528.1| PREDICTED: hemK methyltransferase family member 2 [Drosophil
>gi|1036853228|ref|XP_017150870.1| PREDICTED: uncharacterized protein LOC108161160 [Drosophila
>gi|1036865269|ref|XP_017151538.1| PREDICTED: prothymosin alpha-B [Drosophila miranda]
>gi|1036865251|ref|XP_017151537.1| PREDICTED: phosphoglycerate kinase [Drosophila miranda]
>gi|1036865232|ref|XP_017151536.1| PREDICTED: pre-mRNA-splicing factor CWC25 homolog [Drosophil
>gi|1036853247|ref|XP_017150871.1| PREDICTED: LOW QUALITY PROTEIN: phosphoglycerate kinase [Dro
>gi|1036865154|ref|XP_017151531.1| PREDICTED: LOW QUALITY PROTEIN: hepatocyte growth factor-reg
>gi|1036865175|ref|XP_017151532.1| PREDICTED: ATP-binding cassette sub-family G member 1 [Droso
>gi|1036865194|ref|XP_017151533.1| PREDICTED: ATP-binding cassette sub-family G member 1 [Droso
>gi|1036865213|ref|XP_017151534.1| PREDICTED: ATP-binding cassette sub-family G member 1 [Droso
>gi|1036931697|ref|XP_017155239.1| PREDICTED: LOW QUALITY PROTEIN: transmembrane and TPR repeat
>gi|1036931673|ref|XP_017155238.1| PREDICTED: TBP-related factor [Drosophila miranda]
>gi|1036933204|ref|XP_017155321.1| PREDICTED: putative gustatory receptor 23a, isoforms A/C [Dr
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>gi|1036933043|ref|XP_017155311.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036933124|ref|XP_017155316.1| PREDICTED: uncharacterized protein LOC108164200 isoform X6
>gi|1036933110|ref|XP_017155315.1| PREDICTED: uncharacterized protein LOC108164200 isoform X5
>gi|1036933079|ref|XP_017155313.1| PREDICTED: uncharacterized protein LOC108164200 isoform X3
>gi|1036933061|ref|XP_017155312.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036933096|ref|XP_017155314.1| PREDICTED: uncharacterized protein LOC108164200 isoform X4
>gi|1036933171|ref|XP_017155319.1| PREDICTED: uncharacterized protein LOC108164200 isoform X9
>gi|1036933152|ref|XP_017155318.1| PREDICTED: uncharacterized protein LOC108164200 isoform X8
>gi|1036933138|ref|XP_017155317.1| PREDICTED: uncharacterized protein LOC108164200 isoform X7
>gi|1036860729|ref|XP_017151291.1| PREDICTED: probable proteasome subunit beta type-2 [Drosophi
>gi|1036853266|ref|XP_017150872.1| PREDICTED: LOW QUALITY PROTEIN: odorant receptor 23a [Drosop
>gi|1036932811|ref|XP_017155296.1| PREDICTED: uncharacterized protein LOC108164190 isoform X1
>gi|1036932829|ref|XP_017155297.1| PREDICTED: uncharacterized protein LOC108164190 isoform X2
>gi|1036853285|ref|XP_017150874.1| PREDICTED: uncharacterized protein LOC108161163 [Drosophila
>gi|1036853302|ref|XP_017150875.1| PREDICTED: uncharacterized protein LOC108161164 [Drosophila
>gi|1036860375|ref|XP_017151271.1| PREDICTED: uncharacterized protein LOC108161506 [Drosophila
>gi|1036863694|ref|XP_017151451.1| PREDICTED: uncharacterized protein LOC108161664 [Drosophila
>gi|1036881447|ref|XP_017152437.1| PREDICTED: uncharacterized protein LOC108162288 [Drosophila
>gi|1036881315|ref|XP_017152429.1| PREDICTED: dynamin-1-like protein [Drosophila miranda]
>gi|1036881409|ref|XP_017152435.1| PREDICTED: actin-related protein 2/3 complex subunit 5 [Dros
>gi|1036881428|ref|XP_017152436.1| PREDICTED: probable NADH dehydrogenase [ubiquinone] 1 alpha
>gi|1036881333|ref|XP_017152430.1| PREDICTED: LOW QUALITY PROTEIN: early boundary activity prot
>gi|1036881352|ref|XP_017152431.1| PREDICTED: LOW QUALITY PROTEIN: protein insensitive [Drosoph
>gi|1036881390|ref|XP_017152433.1| PREDICTED: uncharacterized protein LOC108162285 [Drosophila
>gi|1036881371|ref|XP_017152432.1| PREDICTED: cell cycle checkpoint protein RAD1 [Drosophila mi
>gi|1036881466|ref|XP_017152438.1| PREDICTED: probable cytochrome P450 309a2 [Drosophila mirand
>gi|1036881484|ref|XP_017152439.1| PREDICTED: LOW QUALITY PROTEIN: probable cytochrome P450 309
>gi|1036857963|ref|XP_017151153.1| PREDICTED: ficolin-1-like [Drosophila miranda]
>gi|1036853319|ref|XP_017150876.1| PREDICTED: uncharacterized protein LOC108161165 [Drosophila
>gi|1036861336|ref|XP_017151327.1| PREDICTED: uncharacterized protein C4orf22 homolog [Drosophi
>gi|1036861354|ref|XP_017151328.1| PREDICTED: uncharacterized protein C4orf22 homolog [Drosophi
>gi|1036922142|ref|XP_017154716.1| PREDICTED: LOW QUALITY PROTEIN: out at first protein [Drosop
>gi|1036920177|ref|XP_017154599.1| PREDICTED: LOW QUALITY PROTEIN: protein sly1 homolog [Drosop
>gi|1036914448|ref|XP_017154265.1| PREDICTED: protein decapentaplegic [Drosophila miranda]
>gi|1036914466|ref|XP_017154266.1| PREDICTED: protein decapentaplegic [Drosophila miranda]
>gi|1036914484|ref|XP_017154267.1| PREDICTED: protein decapentaplegic [Drosophila miranda]
>gi|1036925700|ref|XP_017154911.1| PREDICTED: LOW QUALITY PROTEIN: glycerate kinase [Drosophila
>gi|1036926817|ref|XP_017154976.1| PREDICTED: LOW QUALITY PROTEIN: pancreatic triacylglycerol l
>gi|1036853338|ref|XP_017150877.1| PREDICTED: pancreatic triacylglycerol lipase [Drosophila mir
>gi|1036933930|ref|XP_017155359.1| PREDICTED: LOW QUALITY PROTEIN: pancreatic triacylglycerol l
>gi|1036932138|ref|XP_017155259.1| PREDICTED: vesicular glutamate transporter 1 [Drosophila mir
>gi|1036932146|ref|XP_017155260.1| PREDICTED: vesicular glutamate transporter 1 [Drosophila mir
>gi|1036932160|ref|XP_017155261.1| PREDICTED: vesicular glutamate transporter 1 [Drosophila mir
>gi|1036932178|ref|XP_017155262.1| PREDICTED: vesicular glutamate transporter 1 [Drosophila mir
>gi|1036921160|ref|XP_017154660.1| PREDICTED: uncharacterized protein LOC108163721 [Drosophila
>gi|1036921142|ref|XP_017154659.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein At4g
>gi|1036853357|ref|XP_017150878.1| PREDICTED: uncharacterized protein LOC108161167 [Drosophila
>gi|1036920514|ref|XP_017154620.1| PREDICTED: LOW QUALITY PROTEIN: EGF domain-specific O-linked
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>gi|1036909220|ref|XP_017153984.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036909238|ref|XP_017153985.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036909260|ref|XP_017153986.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036909278|ref|XP_017153987.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036909294|ref|XP_017153988.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036909312|ref|XP_017153989.1| PREDICTED: uncharacterized protein LOC108163294 [Drosophila
>gi|1036886033|ref|XP_017152688.1| PREDICTED: LOW QUALITY PROTEIN: piezo-type mechanosensitive
>gi|1036886052|ref|XP_017152689.1| PREDICTED: uncharacterized protein LOC108162456 isoform X1
>gi|1036886071|ref|XP_017152690.1| PREDICTED: uncharacterized protein LOC108162456 isoform X2
>gi|1036883434|ref|XP_017152547.1| PREDICTED: uncharacterized protein LOC108162352 isoform X4
>gi|1036883418|ref|XP_017152546.1| PREDICTED: protein Hook homolog 1 isoform X3 [Drosophila mi
>gi|1036883384|ref|XP_017152543.1| PREDICTED: protein Hook homolog 1 isoform X1 [Drosophila mi
>gi|1036883400|ref|XP_017152544.1| PREDICTED: protein Hook homolog 1 isoform X2 [Drosophila mi
>gi|1036883490|ref|XP_017152550.1| PREDICTED: LOW QUALITY PROTEIN: 26S proteasome non-ATPase re
>gi|1036883526|ref|XP_017152552.1| PREDICTED: histone H3.3 [Drosophila miranda]
>gi|1036883580|ref|XP_017152555.1| PREDICTED: gametocyte-specific factor 1 [Drosophila miranda]
>gi|1036883453|ref|XP_017152548.1| PREDICTED: CMP-sialic acid transporter 1 [Drosophila miranda
>gi|1036883472|ref|XP_017152549.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036883346|ref|XP_017152541.1| PREDICTED: receptor-interacting serine/threonine-protein ki
>gi|1036883365|ref|XP_017152542.1| PREDICTED: ankyrin repeat, PH and SEC7 domain containing pr
>gi|1036883509|ref|XP_017152551.1| PREDICTED: ankyrin repeat domain-containing protein 39 [Dro
>gi|1036883544|ref|XP_017152553.1| PREDICTED: uncharacterized protein LOC108162359 [Drosophila
>gi|1036883562|ref|XP_017152554.1| PREDICTED: uncharacterized protein LOC108162360 [Drosophila
>gi|1036898431|ref|XP_017153386.1| PREDICTED: reticulocalbin-2 isoform X1 [Drosophila miranda]
>gi|1036898465|ref|XP_017153388.1| PREDICTED: LOW QUALITY PROTEIN: malate dehydrogenase, mitoch
>gi|1036898448|ref|XP_017153387.1| PREDICTED: reticulocalbin-2 isoform X2 [Drosophila miranda]
>gi|1036898482|ref|XP_017153390.1| PREDICTED: SPRY domain-containing SOCS box protein 3 [Drosop
>gi|1036898499|ref|XP_017153391.1| PREDICTED: uncharacterized protein LOC108162920 [Drosophila
>gi|1036898414|ref|XP_017153385.1| PREDICTED: LOW QUALITY PROTEIN: collagen alpha-1(IV) chain
>gi|1036869603|ref|XP_017151781.1| PREDICTED: collagen alpha-1(III) chain [Drosophila miranda]
>gi|1036869717|ref|XP_017151788.1| PREDICTED: selT-like protein [Drosophila miranda]
>gi|1036869774|ref|XP_017151792.1| PREDICTED: general transcription factor IIH subunit 5 isofo
>gi|1036869757|ref|XP_017151791.1| PREDICTED: general transcription factor IIH subunit 5 isofo
>gi|1036869738|ref|XP_017151789.1| PREDICTED: zinc finger HIT domain-containing protein 1 [Dro
>gi|1036869698|ref|XP_017151787.1| PREDICTED: LOW QUALITY PROTEIN: reticulon-1 [Drosophila mira
>gi|1036869679|ref|XP_017151786.1| PREDICTED: UHRF1-binding protein 1-like isoform X4 [Drosophi
>gi|1036869641|ref|XP_017151784.1| PREDICTED: UHRF1-binding protein 1 isoform X2 [Drosophila m
>gi|1036869660|ref|XP_017151785.1| PREDICTED: UHRF1-binding protein 1-like isoform X3 [Drosophi
>gi|1036869622|ref|XP_017151783.1| PREDICTED: UHRF1-binding protein 1 isoform X1 [Drosophila m
>gi|1036922611|ref|XP_017154748.1| PREDICTED: LOW QUALITY PROTEIN: UBX domain-containing prote
>gi|1036871883|ref|XP_017151915.1| PREDICTED: inosine triphosphate pyrophosphatase [Drosophila
>gi|1036871867|ref|XP_017151914.1| PREDICTED: mannose-P-dolichol utilization defect 1 protein
>gi|1036871770|ref|XP_017151907.1| PREDICTED: GDP-mannose 4,6 dehydratase [Drosophila miranda]
>gi|1036871820|ref|XP_017151911.1| PREDICTED: LOW QUALITY PROTEIN: 39S ribosomal protein L28, m
>gi|1036871899|ref|XP_017151916.1| PREDICTED: LOW QUALITY PROTEIN: DAZ-associated protein 2 [Dr
>gi|1036871801|ref|XP_017151909.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036871835|ref|XP_017151912.1| PREDICTED: protein SCO1 homolog, mitochondrial [Drosophila
>gi|1036871786|ref|XP_017151908.1| PREDICTED: DNA-directed RNA polymerases I and III subunit RP
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>gi|1036871852|ref|XP_017151913.1| PREDICTED: uncharacterized protein LOC108161965 [Drosophila
>gi|1036871751|ref|XP_017151906.1| PREDICTED: LOW QUALITY PROTEIN: protein KBP homolog [Drosoph
>gi|1036871712|ref|XP_017151904.1| PREDICTED: LOW QUALITY PROTEIN: cilia- and flagella-associat
>gi|1036871732|ref|XP_017151905.1| PREDICTED: LOW QUALITY PROTEIN: SWI/SNF-related matrix-assoc
>gi|1036934191|ref|XP_017155372.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036926833|ref|XP_017154978.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036926851|ref|XP_017154979.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 subunit C2 [Dro
>gi|1036906134|ref|XP_017153822.1| PREDICTED: LOW QUALITY PROTEIN: protein mothers against dpp
>gi|1036921937|ref|XP_017154705.1| PREDICTED: probable tyrosyl-DNA phosphodiesterase [Drosophi
>gi|1036920063|ref|XP_017154592.1| PREDICTED: abhydrolase domain-containing protein 2 [Drosophi
>gi|1036920081|ref|XP_017154593.1| PREDICTED: uncharacterized protein LOC108163678 [Drosophila
>gi|1036920101|ref|XP_017154594.1| PREDICTED: uncharacterized protein LOC108163679 [Drosophila
>gi|1036857909|ref|XP_017151150.1| PREDICTED: uncharacterized protein LOC108161409 isoform X2
>gi|1036857890|ref|XP_017151148.1| PREDICTED: uncharacterized protein LOC108161409 isoform X1
>gi|1036853376|ref|XP_017150879.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036933555|ref|XP_017155341.1| PREDICTED: uncharacterized protein LOC108164219 [Drosophila
>gi|1036933573|ref|XP_017155342.1| PREDICTED: uncharacterized protein LOC108164219 [Drosophila
>gi|1036924575|ref|XP_017154853.1| PREDICTED: uncharacterized protein LOC108163847 [Drosophila
>gi|1036924557|ref|XP_017154852.1| PREDICTED: LOW QUALITY PROTEIN: otopetrin-2 [Drosophila mira
>gi|1036935807|ref|XP_017155466.1| PREDICTED: uncharacterized protein LOC108164324 [Drosophila
>gi|1036911835|ref|XP_017154119.1| PREDICTED: uncharacterized protein LOC108163377 [Drosophila
>gi|1036911800|ref|XP_017154116.1| PREDICTED: eukaryotic translation initiation factor 4E-like
>gi|1036853394|ref|XP_017150880.1| PREDICTED: uncharacterized protein LOC108161171 [Drosophila
>gi|1036853413|ref|XP_017150881.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036853432|ref|XP_017150883.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036911818|ref|XP_017154118.1| PREDICTED: LOW QUALITY PROTEIN: pupal cuticle protein Edg-78
>gi|1036911783|ref|XP_017154115.1| PREDICTED: LOW QUALITY PROTEIN: DNA-directed RNA polymerase
>gi|1036890129|ref|XP_017152917.1| PREDICTED: LOW QUALITY PROTEIN: multiple epidermal growth fa
>gi|1036874961|ref|XP_017152088.1| PREDICTED: LOW QUALITY PROTEIN: protein purity of essence [
>gi|1036892670|ref|XP_017153063.1| PREDICTED: LOW QUALITY PROTEIN: TBP-related factor [Drosoph
>gi|1036892650|ref|XP_017153062.1| PREDICTED: LOW QUALITY PROTEIN: CD109 antigen [Drosophila m
>gi|1036883740|ref|XP_017152565.1| PREDICTED: LOW QUALITY PROTEIN: neuroplastin [Drosophila mi
>gi|1036883721|ref|XP_017152564.1| PREDICTED: trithorax group protein osa-like [Drosophila mira
>gi|1036906435|ref|XP_017153839.1| PREDICTED: LOW QUALITY PROTEIN: protein MON2 homolog [Droso
>gi|1036889367|ref|XP_017152873.1| PREDICTED: centrosome-associated zinc finger protein CP190-
>gi|1036889313|ref|XP_017152870.1| PREDICTED: beta-1,3-galactosyltransferase 1-like isoform X1
>gi|1036889330|ref|XP_017152871.1| PREDICTED: beta-1,3-galactosyltransferase 1-like isoform X2
>gi|1036889348|ref|XP_017152872.1| PREDICTED: beta-1,3-galactosyltransferase 1-like isoform X3
>gi|1036889258|ref|XP_017152867.1| PREDICTED: beta-1,3-galactosyltransferase 5-like isoform X1
>gi|1036889296|ref|XP_017152869.1| PREDICTED: uncharacterized protein LOC108162574 isoform X3
>gi|1036889277|ref|XP_017152868.1| PREDICTED: beta-1,3-galactosyltransferase 1-like isoform X2
>gi|1036889240|ref|XP_017152865.1| PREDICTED: LOW QUALITY PROTEIN: UDP-GlcNAc:betaGal beta-1,3-
>gi|1036889478|ref|XP_017152880.1| PREDICTED: beta-1,3-galactosyltransferase 5-like [Drosophila
>gi|1036889403|ref|XP_017152875.1| PREDICTED: beta-1,3-galactosyltransferase 5-like isoform X2
>gi|1036889442|ref|XP_017152878.1| PREDICTED: beta-1,3-galactosyltransferase 5-like isoform X4
>gi|1036889385|ref|XP_017152874.1| PREDICTED: beta-1,3-galactosyltransferase 5-like isoform X1
>gi|1036889423|ref|XP_017152877.1| PREDICTED: beta-1,3-galactosyltransferase 5-like isoform X3
>gi|1036889460|ref|XP_017152879.1| PREDICTED: LOW QUALITY PROTEIN: beta-1,3-galactosyltransfera
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>gi|1036871918|ref|XP_017151917.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036871988|ref|XP_017151922.1| PREDICTED: complement component 1 Q subcomponent-binding pro
>gi|1036872005|ref|XP_017151923.1| PREDICTED: radial spoke head 1 homolog [Drosophila miranda]
>gi|1036871971|ref|XP_017151920.1| PREDICTED: LOW QUALITY PROTEIN: UDP-N-acetylglucosamine--do:
>gi|1036871952|ref|XP_017151919.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036871935|ref|XP_017151918.1| PREDICTED: sorting nexin-29 [Drosophila miranda]
>gi|1036872023|ref|XP_017151924.1| PREDICTED: LOW QUALITY PROTEIN: polycomb protein Sfmbt [Dros
>gi|1036905926|ref|XP_017153809.1| PREDICTED: NAD-dependent histone deacetylase Sir2 [Drosophil
>gi|1036921560|ref|XP_017154683.1| PREDICTED: dnaJ homolog subfamily A member 2 [Drosophila mi:
>gi|1036921524|ref|XP_017154681.1| PREDICTED: protein hunchback [Drosophila miranda]
>gi|1036921542|ref|XP_017154682.1| PREDICTED: protein hunchback [Drosophila miranda]
>gi|1036853451|ref|XP_017150884.1| PREDICTED: glycogen synthase kinase-3 alpha [Drosophila mira
>gi|1036859818|ref|XP_017151242.1| PREDICTED: uncharacterized protein LOC108161483 isoform X1
>gi|1036859835|ref|XP_017151243.1| PREDICTED: uncharacterized protein LOC108161483 isoform X2
>gi|1036932687|ref|XP_017155289.1| PREDICTED: beta-1,3-galactosyltransferase 5-like [Drosophila
>gi|1036932649|ref|XP_017155287.1| PREDICTED: uncharacterized protein LOC108164180 [Drosophila
>gi|1036932667|ref|XP_017155288.1| PREDICTED: uncharacterized protein LOC108164182 [Drosophila
>gi|1036930190|ref|XP_017155155.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036934017|ref|XP_017155365.1| PREDICTED: uncharacterized protein LOC108164241 [Drosophila
>gi|1036933999|ref|XP_017155364.1| PREDICTED: tetratricopeptide repeat protein 30 homolog [Dros
>gi|1036853470|ref|XP_017150885.1| PREDICTED: uncharacterized protein LOC108161175 [Drosophila
>gi|1036891222|ref|XP_017152979.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036926066|ref|XP_017154935.1| PREDICTED: uncharacterized protein LOC108163918 [Drosophila
>gi|1036884971|ref|XP_017152633.1| PREDICTED: LOW QUALITY PROTEIN: ubiquitin conjugation facto:
>gi|1036884950|ref|XP_017152632.1| PREDICTED: LOW QUALITY PROTEIN: target of rapamycin [Drosoph
>gi|1036907989|ref|XP_017153918.1| PREDICTED: LOW QUALITY PROTEIN: V-type proton ATPase cataly1
>gi|1036908025|ref|XP_017153920.1| PREDICTED: uncharacterized protein LOC108163260 isoform X2
>gi|1036908007|ref|XP_017153919.1| PREDICTED: protein YIPF5 isoform X1 [Drosophila miranda]
>gi|1036931336|ref|XP_017155217.1| PREDICTED: LOW QUALITY PROTEIN: V-type proton ATPase cataly1
>gi|1036917331|ref|XP_017154439.1| PREDICTED: V-type proton ATPase catalytic subunit A isoform
>gi|1036917349|ref|XP_017154440.1| PREDICTED: V-type proton ATPase catalytic subunit A isoform
>gi|1036926795|ref|XP_017154975.1| PREDICTED: LOW QUALITY PROTEIN: adenylate cyclase type 7-li
>gi|1036864478|ref|XP_017151492.1| PREDICTED: cylicin-2 [Drosophila miranda]
>gi|1036863713|ref|XP_017151452.1| PREDICTED: uncharacterized protein LOC108161666 [Drosophila
>gi|1036926572|ref|XP_017154962.1| PREDICTED: adenylate cyclase type 2-like isoform X1 [Drosoph
>gi|1036926592|ref|XP_017154963.1| PREDICTED: adenylate cyclase type 2-like isoform X2 [Drosoph
>gi|1036853488|ref|XP_017150887.1| PREDICTED: adenylate cyclase type 5-like [Drosophila miranda
>gi|1036926610|ref|XP_017154964.1| PREDICTED: adenylate cyclase type 8-like isoform X1 [Drosoph
>gi|1036926630|ref|XP_017154965.1| PREDICTED: uncharacterized protein LOC108163939 isoform X2
>gi|1036859646|ref|XP_017151233.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036859665|ref|XP_017151234.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036927753|ref|XP_017155031.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036861447|ref|XP_017151333.1| PREDICTED: uncharacterized protein LOC108161561 isoform X1
>gi|1036861464|ref|XP_017151334.1| PREDICTED: uncharacterized protein LOC108161561 isoform X2
>gi|1036859201|ref|XP_017151209.1| PREDICTED: LOW QUALITY PROTEIN: beta-1,4-glucuronyltransfera
>gi|1036930218|ref|XP_017155157.1| PREDICTED: zinc finger protein 189-like isoform X2 [Drosophi
>gi|1036930204|ref|XP_017155156.1| PREDICTED: zinc finger protein 510-like isoform X1 [Drosophi
>gi|1036853505|ref|XP_017150888.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
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>gi|1036930232|ref|XP_017155158.1| PREDICTED: uncharacterized protein LOC108164084 [Drosophila
>gi|1036930248|ref|XP_017155159.1| PREDICTED: uncharacterized protein LOC108164084 [Drosophila
>gi|1036861661|ref|XP_017151347.1| PREDICTED: LOW QUALITY PROTEIN: multidrug resistance-associ
>gi|1036861521|ref|XP_017151338.1| PREDICTED: PIH1 domain-containing protein 1 isoform X2 [Dros
>gi|1036861540|ref|XP_017151339.1| PREDICTED: PIH1 domain-containing protein 1 isoform X3 [Dros
>gi|1036861502|ref|XP_017151337.1| PREDICTED: pre-mRNA 3' end processing protein WDR33 isoform
>gi|1036861559|ref|XP_017151340.1| PREDICTED: probable cyclin-dependent serine/threonine-prote
>gi|1036861578|ref|XP_017151341.1| PREDICTED: GATA zinc finger domain-containing protein 14 iso
>gi|1036861625|ref|XP_017151345.1| PREDICTED: phosphatidylethanolamine-binding protein 1 [Droso
>gi|1036861609|ref|XP_017151344.1| PREDICTED: LOW QUALITY PROTEIN: magnesium transporter NIPA2
>gi|1036861678|ref|XP_017151348.1| PREDICTED: LOW QUALITY PROTEIN: spermatogenesis-associated p
>gi|1036861593|ref|XP_017151342.1| PREDICTED: thiamine transporter 1 [Drosophila miranda]
>gi|1036861642|ref|XP_017151346.1| PREDICTED: proteasome subunit alpha type-2-like [Drosophila
>gi|1036925819|ref|XP_017154920.1| PREDICTED: PRKCA-binding protein isoform X2 [Drosophila mira
>gi|1036925801|ref|XP_017154919.1| PREDICTED: PRKCA-binding protein isoform X1 [Drosophila mira
>gi|1036925838|ref|XP_017154921.1| PREDICTED: uncharacterized protein LOC108163905 [Drosophila
>gi|1036911753|ref|XP_017154113.1| PREDICTED: intraflagellar transport protein 43 homolog [Dros
>gi|1036911765|ref|XP_017154114.1| PREDICTED: PITH domain-containing protein GA19395 [Drosophil
>gi|1036911739|ref|XP_017154112.1| PREDICTED: T-complex protein 1 subunit delta [Drosophila mi
>gi|1036924893|ref|XP_017154870.1| PREDICTED: uncharacterized protein LOC108163865 [Drosophila
>gi|1036924963|ref|XP_017154874.1| PREDICTED: barrier-to-autointegration factor-like [Drosophil
>gi|1036924983|ref|XP_017154875.1| PREDICTED: barrier-to-autointegration factor-like [Drosophil
>gi|1036925003|ref|XP_017154876.1| PREDICTED: barrier-to-autointegration factor-like [Drosophil
>gi|1036924945|ref|XP_017154873.1| PREDICTED: uncharacterized protein LOC108163868 [Drosophila
>gi|1036924911|ref|XP_017154871.1| PREDICTED: protein ariadne-1-like [Drosophila miranda]
>gi|1036924927|ref|XP_017154872.1| PREDICTED: protein ariadne-1-like [Drosophila miranda]
>gi|1036863532|ref|XP_017151446.1| PREDICTED: tripartite motif-containing protein 40-like [Dros
>gi|1036863476|ref|XP_017151444.1| PREDICTED: tripartite motif-containing protein 40-like [Dros
>gi|1036859459|ref|XP_017151224.1| PREDICTED: protein phosphatase inhibitor 2 [Drosophila mira
>gi|1036853522|ref|XP_017150889.1| PREDICTED: uncharacterized protein LOC108161179 [Drosophila
>gi|1036853539|ref|XP_017150890.1| PREDICTED: uncharacterized protein CG31750 [Drosophila mira
>gi|1036853555|ref|XP_017150891.1| PREDICTED: putative gustatory receptor 36a [Drosophila mira
>gi|1036862779|ref|XP_017151407.1| PREDICTED: uncharacterized protein LOC108161619 [Drosophila
>gi|1036935250|ref|XP_017155431.1| PREDICTED: uncharacterized protein LOC108164295 [Drosophila
>gi|1036934157|ref|XP_017155370.1| PREDICTED: uncharacterized protein LOC108164246 [Drosophila
>gi|1036853572|ref|XP_017150892.1| PREDICTED: transcription initiation factor TFIID subunit 5-1
>gi|1036924284|ref|XP_017154836.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1036853588|ref|XP_017150894.1| PREDICTED: uncharacterized protein LOC108161184 [Drosophila
>gi|1036862852|ref|XP_017151410.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosopl
>gi|1036888672|ref|XP_017152832.1| PREDICTED: uncharacterized protein LOC108162553 isoform X2
>gi|1036888655|ref|XP_017152831.1| PREDICTED: uncharacterized protein LOC108162553 isoform X1
>gi|1036888689|ref|XP_017152834.1| PREDICTED: uncharacterized protein LOC108162553 isoform X3
>gi|1036888707|ref|XP_017152835.1| PREDICTED: uncharacterized protein LOC108162553 isoform X3
>gi|1036888638|ref|XP_017152830.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein kin
>gi|1036853605|ref|XP_017150895.1| PREDICTED: integumentary mucin C.1-like, partial [Drosophila
>gi|1036888744|ref|XP_017152836.1| PREDICTED: uncharacterized protein LOC108162554 [Drosophila
>gi|1036888762|ref|XP_017152837.1| PREDICTED: LOW QUALITY PROTEIN: integrin beta-nu [Drosophila
>gi|1036922413|ref|XP_017154734.1| PREDICTED: estradiol 17-beta-dehydrogenase 11 [Drosophila m
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>gi|1036932776|ref|XP_017155295.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036857414|ref|XP_017151123.1| PREDICTED: uncharacterized protein LOC108161385 [Drosophila
>gi|1036932252|ref|XP_017155266.1| PREDICTED: uncharacterized protein LOC108164166 [Drosophila
>gi|1036932270|ref|XP_017155267.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036866993|ref|XP_017151637.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036867012|ref|XP_017151638.1| PREDICTED: uncharacterized protein LOC108161799 [Drosophila
>gi|1036867104|ref|XP_017151644.1| PREDICTED: uncharacterized protein LOC108161801 isoform X4
>gi|1036867048|ref|XP_017151640.1| PREDICTED: uncharacterized protein LOC108161801 isoform X1
>gi|1036867066|ref|XP_017151642.1| PREDICTED: uncharacterized protein LOC108161801 isoform X2
>gi|1036867086|ref|XP_017151643.1| PREDICTED: uncharacterized protein LOC108161801 isoform X3
>gi|1036867126|ref|XP_017151645.1| PREDICTED: uncharacterized protein LOC108161801 isoform X5
>gi|1036867145|ref|XP_017151646.1| PREDICTED: uncharacterized protein LOC108161801 isoform X5
>gi|1036866976|ref|XP_017151636.1| PREDICTED: WD repeat-containing protein 75 [Drosophila miran
>gi|1036867164|ref|XP_017151647.1| PREDICTED: uncharacterized protein LOC108161802 [Drosophila
>gi|1036867031|ref|XP_017151639.1| PREDICTED: malectin [Drosophila miranda]
>gi|1036866942|ref|XP_017151634.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036866960|ref|XP_017151635.1| PREDICTED: LOW QUALITY PROTEIN: intersectin-1 [Drosophila m
>gi|1036925126|ref|XP_017154877.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036910439|ref|XP_017154044.1| PREDICTED: probable ATP-dependent RNA helicase DDX47 [Droso
>gi|1036910401|ref|XP_017154042.1| PREDICTED: uncharacterized protein LOC108163323 isoform X3
>gi|1036910421|ref|XP_017154043.1| PREDICTED: uncharacterized protein LOC108163323 isoform X4
>gi|1036910383|ref|XP_017154041.1| PREDICTED: uncharacterized protein LOC108163323 isoform X2
>gi|1036910365|ref|XP_017154040.1| PREDICTED: uncharacterized protein LOC108163323 isoform X1
>gi|1036891563|ref|XP_017152999.1| PREDICTED: transcription factor E2F2 [Drosophila miranda]
>gi|1036891597|ref|XP_017153001.1| PREDICTED: uncharacterized protein LOC108162670 isoform X1
>gi|1036891616|ref|XP_017153002.1| PREDICTED: uncharacterized protein LOC108162670 isoform X2
>gi|1036891635|ref|XP_017153003.1| PREDICTED: M-phase phosphoprotein 6 [Drosophila miranda]
>gi|1036891580|ref|XP_017153000.1| PREDICTED: ubiquinone biosynthesis O-methyltransferase [Dros
>gi|1036891545|ref|XP_017152997.1| PREDICTED: phospholipase D3 [Drosophila miranda]
>gi|1036891527|ref|XP_017152996.1| PREDICTED: exonuclease mut-7 homolog [Drosophila miranda]
>gi|1036893713|ref|XP_017153119.1| PREDICTED: LOW QUALITY PROTEIN: nucleolar complex protein 2
>gi|1036893742|ref|XP_017153122.1| PREDICTED: mimitin, mitochondrial [Drosophila miranda]
>gi|1036893727|ref|XP_017153120.1| PREDICTED: LOW QUALITY PROTEIN: phospholipase D3 [Drosophila
>gi|1036893699|ref|XP_017153118.1| PREDICTED: probable aconitate hydratase, mitochondrial [Dros
>gi|1036933981|ref|XP_017155362.1| PREDICTED: uncharacterized protein LOC108164239 [Drosophila
>gi|1036869397|ref|XP_017151769.1| PREDICTED: GMP synthase [glutamine-hydrolyzing] [Drosophila
>gi|1036869251|ref|XP_017151760.1| PREDICTED: protein MCM10 homolog [Drosophila miranda]
>gi|1036869269|ref|XP_017151761.1| PREDICTED: protein MCM10 homolog [Drosophila miranda]
>gi|1036869194|ref|XP_017151756.1| PREDICTED: uncharacterized protein LOC108161877 [Drosophila
>gi|1036869213|ref|XP_017151757.1| PREDICTED: uncharacterized protein LOC108161877 [Drosophila
>gi|1036869232|ref|XP_017151758.1| PREDICTED: uncharacterized protein LOC108161877 [Drosophila
>gi|1036869380|ref|XP_017151768.1| PREDICTED: uncharacterized protein LOC108161885 [Drosophila
>gi|1036869362|ref|XP_017151766.1| PREDICTED: uncharacterized protein LOC108161882 [Drosophila
>gi|1036869303|ref|XP_017151763.1| PREDICTED: uncharacterized protein LOC108161880 [Drosophila
>gi|1036869345|ref|XP_017151765.1| PREDICTED: glucosamine-6-phosphate isomerase isoform X2 [Dro
>gi|1036869322|ref|XP_017151764.1| PREDICTED: glucosamine-6-phosphate isomerase isoform X1 [Dro
>gi|1036869287|ref|XP_017151762.1| PREDICTED: lamin Dm0 [Drosophila miranda]
>gi|1036867693|ref|XP_017151678.1| PREDICTED: ATP-dependent RNA helicase WM6 [Drosophila mirand
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>gi|1036867636|ref|XP_017151675.1| PREDICTED: glycoprotein endo-alpha-1,2-mannosidase [Drosophi
>gi|1036867733|ref|XP_017151680.1| PREDICTED: uncharacterized protein LOC108161833 [Drosophila
>gi|1036867579|ref|XP_017151671.1| PREDICTED: uncharacterized protein LOC108161824 [Drosophila
>gi|1036867655|ref|XP_017151676.1| PREDICTED: ubiquinone biosynthesis monooxygenase COQ6, mito
>gi|1036867617|ref|XP_017151674.1| PREDICTED: glycosyltransferase 25 family member [Drosophila
>gi|1036867751|ref|XP_017151681.1| PREDICTED: cytochrome c oxidase assembly protein COX11, mito
>gi|1036867674|ref|XP_017151677.1| PREDICTED: putative elongator complex protein 4 [Drosophila
>gi|1036867598|ref|XP_017151673.1| PREDICTED: LOW QUALITY PROTEIN: vacuolar protein sorting-ass
>gi|1036867789|ref|XP_017151684.1| PREDICTED: thioredoxin domain-containing protein 17 [Drosoph
>gi|1036867770|ref|XP_017151682.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036867714|ref|XP_017151679.1| PREDICTED: uncharacterized protein LOC108161831 [Drosophila
>gi|1036860451|ref|XP_017151276.1| PREDICTED: uncharacterized protein LOC108161510 [Drosophila
>gi|1036925285|ref|XP_017154887.1| PREDICTED: LOW QUALITY PROTEIN: T-box protein H15 [Drosophi
>gi|1036935479|ref|XP_017155445.1| PREDICTED: lysophospholipid acyltransferase 5-like [Drosoph
>gi|1036930643|ref|XP_017155181.1| PREDICTED: LOW QUALITY PROTEIN: T-box protein H15 [Drosophi
>gi|1036935876|ref|XP_017155471.1| PREDICTED: LOW QUALITY PROTEIN: cytosolic non-specific dipep
>gi|1036882112|ref|XP_017152475.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein pho
>gi|1036853622|ref|XP_017150896.1| PREDICTED: cytosolic non-specific dipeptidase-like [Drosophi
>gi|1036882131|ref|XP_017152476.1| PREDICTED: acyl-CoA synthetase family member 2, mitochondria
>gi|1036881984|ref|XP_017152467.1| PREDICTED: mucin-4 isoform X1 [Drosophila miranda]
>gi|1036882003|ref|XP_017152468.1| PREDICTED: uncharacterized protein LOC108162309 isoform X2
>gi|1036882021|ref|XP_017152469.1| PREDICTED: uncharacterized protein LOC108162309 isoform X3
>gi|1036882040|ref|XP_017152470.1| PREDICTED: filamin A-interacting protein 1-like isoform X4
>gi|1036882058|ref|XP_017152471.1| PREDICTED: filamin A-interacting protein 1-like isoform X5
>gi|1036882076|ref|XP_017152472.1| PREDICTED: filamin A-interacting protein 1-like isoform X6
>gi|1036882093|ref|XP_017152474.1| PREDICTED: filamin A-interacting protein 1-like isoform X7
>gi|1036902893|ref|XP_017153636.1| PREDICTED: LOW QUALITY PROTEIN: laminin subunit beta-1 [Dros
>gi|1036865876|ref|XP_017151572.1| PREDICTED: LOW QUALITY PROTEIN: protein split ends [Drosophi
>gi|1036926554|ref|XP_017154961.1| PREDICTED: E3 SUMO-protein ligase NSE2-like [Drosophila mira
>gi|1036927365|ref|XP_017155008.1| PREDICTED: 39S ribosomal protein L10, mitochondrial [Drosoph
>gi|1036927328|ref|XP_017155004.1| PREDICTED: homeobox protein Mohawk isoform X1 [Drosophila mi
>gi|1036927345|ref|XP_017155005.1| PREDICTED: homeobox protein Mohawk isoform X2 [Drosophila mi
>gi|1036864576|ref|XP_017151498.1| PREDICTED: TBC1 domain family member 17 [Drosophila miranda]
>gi|1036864616|ref|XP_017151501.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036864594|ref|XP_017151499.1| PREDICTED: uncharacterized protein LOC108161698 [Drosophila
>gi|1036926470|ref|XP_017154955.1| PREDICTED: ribonuclease P protein subunit p30 [Drosophila mi
>gi|1036881222|ref|XP_017152423.1| PREDICTED: nuclear RNA export factor 1 [Drosophila miranda]
>gi|1036881187|ref|XP_017152421.1| PREDICTED: testis-expressed sequence 2 protein-like [Drosoph
>gi|1036881279|ref|XP_017152427.1| PREDICTED: tRNA-dihydrouridine(16/17) synthase [NAD(P)(+)]-1
>gi|1036881168|ref|XP_017152420.1| PREDICTED: uncharacterized protein LOC108162273 [Drosophila
>gi|1036881204|ref|XP_017152422.1| PREDICTED: uncharacterized protein LOC108162275 [Drosophila
>gi|1036881260|ref|XP_017152425.1| PREDICTED: stress response protein NST1 [Drosophila miranda]
>gi|1036881298|ref|XP_017152428.1| PREDICTED: uncharacterized protein LOC108162280 [Drosophila
>gi|1036881241|ref|XP_017152424.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036925497|ref|XP_017154899.1| PREDICTED: protein smoothened-like [Drosophila miranda]
>gi|1036870799|ref|XP_017151851.1| PREDICTED: putative pre-mRNA-splicing factor ATP-dependent
>gi|1036870853|ref|XP_017151854.1| PREDICTED: general transcription factor 3C polypeptide 5-li
>gi|1036870817|ref|XP_017151852.1| PREDICTED: asparagine--tRNA ligase, cytoplasmic-like [Drosop
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>gi|1036870836|ref|XP_017151853.1| PREDICTED: asparagine--tRNA ligase, cytoplasmic-like [Drosop
>gi|1036870872|ref|XP_017151855.1| PREDICTED: dynactin subunit 6-like [Drosophila miranda]
>gi|1036870891|ref|XP_017151857.1| PREDICTED: probable DNA-directed RNA polymerases I and III s
>gi|1036870645|ref|XP_017151843.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870667|ref|XP_017151844.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870686|ref|XP_017151845.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870705|ref|XP_017151846.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870724|ref|XP_017151847.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870743|ref|XP_017151848.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870761|ref|XP_017151849.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036870780|ref|XP_017151850.1| PREDICTED: brain tumor protein-like [Drosophila miranda]
>gi|1036928617|ref|XP_017155072.1| PREDICTED: keratin-associated protein 5-4-like [Drosophila r
>gi|1036889691|ref|XP_017152893.1| PREDICTED: uncharacterized protein LOC108162588 [Drosophila
>gi|1036889582|ref|XP_017152886.1| PREDICTED: LOW QUALITY PROTEIN: protein smoothened-like [Dro
>gi|1036889673|ref|XP_017152892.1| PREDICTED: androgen-induced gene 1 protein [Drosophila miran
>gi|1036889655|ref|XP_017152891.1| PREDICTED: androgen-induced gene 1 protein isoform X3 [Dros
>gi|1036889619|ref|XP_017152889.1| PREDICTED: androgen-induced gene 1 protein isoform X1 [Dros
>gi|1036889637|ref|XP_017152890.1| PREDICTED: androgen-induced gene 1 protein isoform X2 [Dros
>gi|1036889601|ref|XP_017152887.1| PREDICTED: protein amnionless [Drosophila miranda]
>gi|1036858997|ref|XP_017151196.1| PREDICTED: uncharacterized protein CG13380-like isoform X2
>gi|1036858978|ref|XP_017151195.1| PREDICTED: uncharacterized protein CG13380-like isoform X1
>gi|1036923094|ref|XP_017154778.1| PREDICTED: protein PET117 homolog, mitochondrial [Drosophila
>gi|1036923076|ref|XP_017154777.1| PREDICTED: splicing factor U2af 38 kDa subunit [Drosophila r
>gi|1036920280|ref|XP_017154605.1| PREDICTED: stress-induced-phosphoprotein 1 isoform X1 [Dros
>gi|1036920298|ref|XP_017154606.1| PREDICTED: stress-induced-phosphoprotein 1 isoform X2 [Dros
>gi|1036922340|ref|XP_017154729.1| PREDICTED: phosphatidylinositol 3-kinase regulatory subunit
>gi|1036897377|ref|XP_017153325.1| PREDICTED: dedicator of cytokinesis protein 11 isoform X1 [
>gi|1036897396|ref|XP_017153327.1| PREDICTED: dedicator of cytokinesis protein 11 isoform X2 [
>gi|1036897415|ref|XP_017153328.1| PREDICTED: dedicator of cytokinesis protein 11 isoform X3 [
>gi|1036897472|ref|XP_017153330.1| PREDICTED: uncharacterized protein LOC108162885 isoform X2
>gi|1036897453|ref|XP_017153329.1| PREDICTED: uncharacterized protein LOC108162885 isoform X1
>gi|1036897491|ref|XP_017153331.1| PREDICTED: uncharacterized protein LOC108162885 isoform X3
>gi|1036863400|ref|XP_017151441.1| PREDICTED: uncharacterized protein LOC108161649 [Drosophila
>gi|1036935228|ref|XP_017155430.1| PREDICTED: uncharacterized protein LOC108164294 [Drosophila
>gi|1036859571|ref|XP_017151228.1| PREDICTED: putative gustatory receptor 28b isoform X1 [Dros
>gi|1036859590|ref|XP_017151229.1| PREDICTED: putative gustatory receptor 28b isoform X2 [Dros
>gi|1036859609|ref|XP_017151231.1| PREDICTED: putative gustatory receptor 28b isoform X3 [Dros
>gi|1036853637|ref|XP_017150897.1| PREDICTED: putative gustatory receptor 28a [Drosophila miran
>gi|1036930539|ref|XP_017155174.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036930556|ref|XP_017155175.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036905979|ref|XP_017153813.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036906009|ref|XP_017153814.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036906027|ref|XP_017153815.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036906044|ref|XP_017153816.1| PREDICTED: clavesin-1 [Drosophila miranda]
>gi|1036905959|ref|XP_017153812.1| PREDICTED: dihydrolipoyllysine-residue acetyltransferase con
>gi|1036905942|ref|XP_017153810.1| PREDICTED: dihydrolipoyllysine-residue acetyltransferase con
>gi|1036896759|ref|XP_017153289.1| PREDICTED: histone acetyltransferase KAT7 [Drosophila miranc
>gi|1036896832|ref|XP_017153293.1| PREDICTED: SOSS complex subunit B homolog [Drosophila mirand
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>gi|1036896851|ref|XP_017153294.1| PREDICTED: SOSS complex subunit B homolog [Drosophila miran
>gi|1036896869|ref|XP_017153296.1| PREDICTED: SOSS complex subunit B homolog [Drosophila miran
>gi|1036896888|ref|XP_017153297.1| PREDICTED: SOSS complex subunit B homolog [Drosophila miran
>gi|1036896816|ref|XP_017153292.1| PREDICTED: probable trehalose-phosphate phosphatase C [Dros
>gi|1036896778|ref|XP_017153290.1| PREDICTED: probable trehalose-phosphate phosphatase C [Dros
>gi|1036896797|ref|XP_017153291.1| PREDICTED: probable trehalose-phosphate phosphatase C [Dros
>gi|1036930436|ref|XP_017155169.1| PREDICTED: ras-related and estrogen-regulated growth inhibi
>gi|1036930418|ref|XP_017155168.1| PREDICTED: armadillo repeat-containing protein 4 [Drosophila
>gi|1036878778|ref|XP_017152290.1| PREDICTED: WD repeat-containing protein 82 [Drosophila miran
>gi|1036878752|ref|XP_017152289.1| PREDICTED: mitochondrial ribosome-associated GTPase 2 [Dros
>gi|1036878799|ref|XP_017152291.1| PREDICTED: haloacid dehalogenase-like hydrolase domain-conta
>gi|1036878643|ref|XP_017152282.1| PREDICTED: alanine--tRNA ligase, cytoplasmic [Drosophila mi
>gi|1036878661|ref|XP_017152283.1| PREDICTED: alanine--tRNA ligase, cytoplasmic [Drosophila mi
>gi|1036878817|ref|XP_017152292.1| PREDICTED: uncharacterized protein LOC108162197 [Drosophila
>gi|1036878833|ref|XP_017152294.1| PREDICTED: uncharacterized protein LOC108162198 [Drosophila
>gi|1036878852|ref|XP_017152295.1| PREDICTED: uncharacterized protein LOC108162198 [Drosophila
>gi|1036878871|ref|XP_017152296.1| PREDICTED: uncharacterized protein LOC108162198 [Drosophila
>gi|1036878890|ref|XP_017152297.1| PREDICTED: uncharacterized protein LOC108162198 [Drosophila
>gi|1036878909|ref|XP_017152299.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1036878679|ref|XP_017152285.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036878697|ref|XP_017152286.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036878714|ref|XP_017152287.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036878733|ref|XP_017152288.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036878928|ref|XP_017152300.1| PREDICTED: uncharacterized protein LOC108162200 [Drosophila
>gi|1036853652|ref|XP_017150898.1| PREDICTED: uncharacterized protein LOC108161188 [Drosophila
>gi|1036932440|ref|XP_017155277.1| PREDICTED: uncharacterized protein LOC108164172 [Drosophila
>gi|1036915210|ref|XP_017154315.1| PREDICTED: uncharacterized protein LOC108163501 [Drosophila
>gi|1036915192|ref|XP_017154314.1| PREDICTED: uncharacterized protein LOC108163500 [Drosophila
>gi|1036915174|ref|XP_017154313.1| PREDICTED: protein fuzzy homolog [Drosophila miranda]
>gi|1036890148|ref|XP_017152918.1| PREDICTED: exportin-1 [Drosophila miranda]
>gi|1036890167|ref|XP_017152919.1| PREDICTED: alpha-N-acetylglucosaminidase [Drosophila miranda
>gi|1036924628|ref|XP_017154854.1| PREDICTED: uncharacterized protein LOC108163850 [Drosophila
>gi|1036905426|ref|XP_017153776.1| PREDICTED: mucin-17 isoform X1 [Drosophila miranda]
>gi|1036905444|ref|XP_017153778.1| PREDICTED: mucin-17 isoform X1 [Drosophila miranda]
>gi|1036905462|ref|XP_017153779.1| PREDICTED: myristoylated alanine-rich C-kinase substrate iso
>gi|1036882935|ref|XP_017152516.1| PREDICTED: protein gurken isoform X1 [Drosophila miranda]
>gi|1036882954|ref|XP_017152517.1| PREDICTED: protein gurken isoform X2 [Drosophila miranda]
>gi|1036882898|ref|XP_017152514.1| PREDICTED: uncharacterized protein LOC108162333 [Drosophila
>gi|1036882973|ref|XP_017152518.1| PREDICTED: DNA polymerase epsilon subunit 3 [Drosophila mira
>gi|1036882859|ref|XP_017152512.1| PREDICTED: DNA repair protein complementing XP-G cells homol
>gi|1036882880|ref|XP_017152513.1| PREDICTED: DNA repair protein complementing XP-G cells homol
>gi|1036882917|ref|XP_017152515.1| PREDICTED: uncharacterized protein LOC108162334 [Drosophila
>gi|1036913800|ref|XP_017154234.1| PREDICTED: uncharacterized protein LOC108163456 [Drosophila
>gi|1036905355|ref|XP_017153772.1| PREDICTED: cullin-2 [Drosophila miranda]
>gi|1036905373|ref|XP_017153773.1| PREDICTED: cullin-2 [Drosophila miranda]
>gi|1036905319|ref|XP_017153770.1| PREDICTED: adenylate cyclase type 3 isoform X1 [Drosophila m
>gi|1036905337|ref|XP_017153771.1| PREDICTED: adenylate cyclase type 3 isoform X2 [Drosophila m
>gi|1036912207|ref|XP_017154138.1| PREDICTED: cytochrome c1 [Drosophila miranda]
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>gi|1036910218|ref|XP_017154031.1| PREDICTED: choline/ethanolamine kinase isoform X2 [Drosophil
>gi|1036910198|ref|XP_017154030.1| PREDICTED: choline/ethanolamine kinase isoform X1 [Drosophil
>gi|1036910233|ref|XP_017154032.1| PREDICTED: uncharacterized protein LOC108163317 isoform X3
>gi|1036909604|ref|XP_017154006.1| PREDICTED: male-specific sperm protein Mst84Da [Drosophila r
>gi|1036909555|ref|XP_017154004.1| PREDICTED: uncharacterized protein LOC108163303 isoform X1
>gi|1036909571|ref|XP_017154005.1| PREDICTED: uncharacterized protein LOC108163303 isoform X2
>gi|1036853669|ref|XP_017150899.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036918348|ref|XP_017154500.1| PREDICTED: uncharacterized protein LOC108163616 isoform X2
>gi|1036918366|ref|XP_017154501.1| PREDICTED: uncharacterized protein LOC108163616 isoform X3
>gi|1036918330|ref|XP_017154498.1| PREDICTED: uncharacterized protein LOC108163616 isoform X1
>gi|1036918240|ref|XP_017154493.1| PREDICTED: ADAMTS-like protein 1 isoform X1 [Drosophila mira
>gi|1036918276|ref|XP_017154495.1| PREDICTED: ADAMTS-like protein 1 isoform X3 [Drosophila mira
>gi|1036918294|ref|XP_017154496.1| PREDICTED: ADAMTS-like protein 1 isoform X4 [Drosophila mira
>gi|1036918312|ref|XP_017154497.1| PREDICTED: ADAMTS-like protein 1 isoform X5 [Drosophila mira
>gi|1036918258|ref|XP_017154494.1| PREDICTED: ADAMTS-like protein 1 isoform X2 [Drosophila mira
>gi|1036898706|ref|XP_017153403.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036898725|ref|XP_017153404.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036898744|ref|XP_017153405.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036898763|ref|XP_017153406.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036898781|ref|XP_017153407.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036898806|ref|XP_017153408.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036898670|ref|XP_017153402.1| PREDICTED: uncharacterized protein LOC108162931 [Drosophila
>gi|1036923988|ref|XP_017154820.1| PREDICTED: uncharacterized protein LOC108163831 isoform X1
>gi|1036924005|ref|XP_017154821.1| PREDICTED: uncharacterized protein LOC108163831 isoform X1
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>gi|1036924041|ref|XP_017154824.1| PREDICTED: uncharacterized protein LOC108163831 isoform X2
>gi|1036924059|ref|XP_017154825.1| PREDICTED: uncharacterized protein LOC108163831 isoform X3
>gi|1036924077|ref|XP_017154826.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X4
>gi|1036919542|ref|XP_017154560.1| PREDICTED: serine/threonine-protein kinase SBK1 [Drosophila
>gi|1036919558|ref|XP_017154561.1| PREDICTED: uncharacterized protein LOC108163653 [Drosophila
>gi|1036919490|ref|XP_017154557.1| PREDICTED: tubulin glycylase 3A [Drosophila miranda]
>gi|1036919508|ref|XP_017154558.1| PREDICTED: tubulin glycylase 3B [Drosophila miranda]
>gi|1036919526|ref|XP_017154559.1| PREDICTED: tubulin glycylase 3B [Drosophila miranda]
>gi|1036858511|ref|XP_017151171.1| PREDICTED: RNA-binding protein 39 isoform X2 [Drosophila mi
>gi|1036858452|ref|XP_017151168.1| PREDICTED: trafficking kinesin-binding protein milt isoform
>gi|1036858414|ref|XP_017151165.1| PREDICTED: trafficking kinesin-binding protein milt isoform
>gi|1036858433|ref|XP_017151167.1| PREDICTED: trafficking kinesin-binding protein milt isoform
>gi|1036858473|ref|XP_017151169.1| PREDICTED: trafficking kinesin-binding protein milt isoform
>gi|1036858530|ref|XP_017151172.1| PREDICTED: GATA zinc finger domain-containing protein 14-lik
>gi|1036858701|ref|XP_017151182.1| PREDICTED: protein tyrosine phosphatase type IVA 1-like [Dro
>gi|1036858720|ref|XP_017151183.1| PREDICTED: protein tyrosine phosphatase type IVA 1-like [Dro
>gi|1036922595|ref|XP_017154747.1| PREDICTED: uncharacterized protein LOC108163779 [Drosophila
>gi|1036929190|ref|XP_017155103.1| PREDICTED: cdc25-like protein phosphatase twine [Drosophila
>gi|1036935566|ref|XP_017155451.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1036885411|ref|XP_017152657.1| PREDICTED: zinc finger protein Elbow [Drosophila miranda]
>gi|1036885794|ref|XP_017152674.1| PREDICTED: partner of bursicon [Drosophila miranda]
>gi|1036853703|ref|XP_017150901.1| PREDICTED: protein toll-like [Drosophila miranda]
>gi|1036885601|ref|XP_017152669.1| PREDICTED: protein toll-like [Drosophila miranda]
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>gi|1036885619|ref|XP_017152670.1| PREDICTED: protein toll-like [Drosophila miranda]
>gi|1036885632|ref|XP_017152671.1| PREDICTED: protein toll-like [Drosophila miranda]
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>gi|1036885563|ref|XP_017152666.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036885582|ref|XP_017152668.1| PREDICTED: DNA ligase 1 [Drosophila miranda]
>gi|1036885811|ref|XP_017152675.1| PREDICTED: ficolin-1-like [Drosophila miranda]
>gi|1036853720|ref|XP_017150902.1| PREDICTED: uncharacterized protein LOC108161193 [Drosophila
>gi|1036853739|ref|XP_017150903.1| PREDICTED: putative leucine-rich repeat-containing protein
>gi|1036885432|ref|XP_017152658.1| PREDICTED: zinc finger protein Noc [Drosophila miranda]
>gi|1036885544|ref|XP_017152665.1| PREDICTED: uncharacterized protein LOC108162439 [Drosophila
>gi|1036853758|ref|XP_017150905.1| PREDICTED: protein toll-like [Drosophila miranda]
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>gi|1036885468|ref|XP_017152661.1| PREDICTED: uncharacterized protein LOC108162438 isoform X1
>gi|1036885487|ref|XP_017152662.1| PREDICTED: uncharacterized protein LOC108162438 isoform X2
>gi|1036885506|ref|XP_017152663.1| PREDICTED: uncharacterized protein LOC108162438 isoform X3
>gi|1036885525|ref|XP_017152664.1| PREDICTED: uncharacterized protein LOC108162438 isoform X4
>gi|1036893109|ref|XP_017153086.1| PREDICTED: protein outspread isoform X1 [Drosophila miranda]
>gi|1036893128|ref|XP_017153088.1| PREDICTED: protein outspread isoform X2 [Drosophila miranda]
>gi|1036893147|ref|XP_017153089.1| PREDICTED: protein outspread isoform X3 [Drosophila miranda]
>gi|1036893166|ref|XP_017153090.1| PREDICTED: protein outspread isoform X4 [Drosophila miranda]
>gi|1036893203|ref|XP_017153092.1| PREDICTED: alcohol dehydrogenase [Drosophila miranda]
>gi|1036893184|ref|XP_017153091.1| PREDICTED: alcohol dehydrogenase-related 31 kDa protein [Dro
>gi|1036859069|ref|XP_017151200.1| PREDICTED: heterogeneous nuclear ribonucleoprotein A1-like
>gi|1036859087|ref|XP_017151201.1| PREDICTED: homeobox protein 13 [Drosophila miranda]
>gi|1036853794|ref|XP_017150907.1| PREDICTED: uncharacterized protein LOC108161197 [Drosophila
>gi|1036924326|ref|XP_017154838.1| PREDICTED: uncharacterized protein LOC108163837 [Drosophila
>gi|1036924304|ref|XP_017154837.1| PREDICTED: uncharacterized protein LOC108163836 [Drosophila
>gi|1036934609|ref|XP_017155398.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1-like [Drosophil
>gi|1036860229|ref|XP_017151266.1| PREDICTED: uncharacterized protein LOC108161501 isoform X1
>gi|1036860248|ref|XP_017151267.1| PREDICTED: uncharacterized protein LOC108161501 isoform X2
>gi|1036919705|ref|XP_017154570.1| PREDICTED: protein Turandot X-like [Drosophila miranda]
>gi|1036919790|ref|XP_017154575.1| PREDICTED: holotricin-3 isoform X3 [Drosophila miranda]
>gi|1036919758|ref|XP_017154573.1| PREDICTED: glycine-rich cell wall structural protein isofor
>gi|1036919772|ref|XP_017154574.1| PREDICTED: holotricin-3 isoform X2 [Drosophila miranda]
>gi|1036919724|ref|XP_017154571.1| PREDICTED: uncharacterized protein LOC108163661 [Drosophila
>gi|1036919740|ref|XP_017154572.1| PREDICTED: uncharacterized protein LOC108163663 [Drosophila
>gi|1036919808|ref|XP_017154577.1| PREDICTED: glycine-rich cell wall structural protein-like [
>gi|1036919828|ref|XP_017154578.1| PREDICTED: uncharacterized protein LOC108163666 [Drosophila
>gi|1036931717|ref|XP_017155240.1| PREDICTED: uncharacterized protein LOC108164147 [Drosophila
>gi|1036864524|ref|XP_017151495.1| PREDICTED: keratin-associated protein 21-1-like [Drosophila
>gi|1036860174|ref|XP_017151262.1| PREDICTED: uncharacterized protein LOC108161499 isoform X1
>gi|1036860192|ref|XP_017151263.1| PREDICTED: uncharacterized protein LOC108161499 isoform X2
>gi|1036858034|ref|XP_017151157.1| PREDICTED: COP9 signalosome complex subunit 1 [Drosophila mi
>gi|1036934728|ref|XP_017155402.1| PREDICTED: ATPase family AAA domain-containing protein 1 [Dr
>gi|1036897800|ref|XP_017153351.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036897818|ref|XP_017153352.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
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>gi|1036897837|ref|XP_017153353.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036897856|ref|XP_017153354.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
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>gi|1036897893|ref|XP_017153356.1| PREDICTED: uncharacterized protein LOC108162897 isoform X1
>gi|1036897930|ref|XP_017153357.1| PREDICTED: uncharacterized protein LOC108162897 isoform X2
>gi|1036897734|ref|XP_017153346.1| PREDICTED: uncharacterized protein LOC108162892 [Drosophila
>gi|1036897966|ref|XP_017153359.1| PREDICTED: LOW QUALITY PROTEIN: E3 ubiquitin-protein ligase
>gi|1036853811|ref|XP_017150908.1| PREDICTED: RING finger protein 5-like [Drosophila miranda]
>gi|1036897874|ref|XP_017153355.1| PREDICTED: ER lumen protein-retaining receptor-like [Drosoph
>gi|1036897750|ref|XP_017153347.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
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>gi|1036910500|ref|XP_017154046.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036910518|ref|XP_017154047.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036910556|ref|XP_017154050.1| PREDICTED: protein rolling stone [Drosophila miranda]
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>gi|1036918038|ref|XP_017154482.1| PREDICTED: neural cell adhesion molecule 1 [Drosophila miran
>gi|1036918020|ref|XP_017154481.1| PREDICTED: protein strawberry notch [Drosophila miranda]
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>gi|1036886808|ref|XP_017152728.1| PREDICTED: dual oxidase maturation factor 1 [Drosophila mira
>gi|1036886899|ref|XP_017152735.1| PREDICTED: dynactin subunit 5 [Drosophila miranda]
>gi|1036886861|ref|XP_017152731.1| PREDICTED: ankyrin repeat domain-containing protein 49 [Dros
>gi|1036886915|ref|XP_017152736.1| PREDICTED: protein Dr1 [Drosophila miranda]
>gi|1036886844|ref|XP_017152730.1| PREDICTED: probable tRNA(His) guanylyltransferase [Drosophil
>gi|1036886827|ref|XP_017152729.1| PREDICTED: mRNA cap guanine-N7 methyltransferase [Drosophila
>gi|1036886880|ref|XP_017152734.1| PREDICTED: uncharacterized protein LOC108162492 [Drosophila
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>gi|1036916892|ref|XP_017154413.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1036916910|ref|XP_017154414.1| PREDICTED: anamorsin homolog [Drosophila miranda]
>gi|1036919670|ref|XP_017154568.1| PREDICTED: suppressor of hairless protein [Drosophila mirand
>gi|1036894832|ref|XP_017153182.1| PREDICTED: myosin-VIIa [Drosophila miranda]
>gi|1036894851|ref|XP_017153184.1| PREDICTED: myosin-VIIa [Drosophila miranda]
>gi|1036924539|ref|XP_017154851.1| PREDICTED: transcription elongation factor S-II [Drosophila
>gi|1036888295|ref|XP_017152810.1| PREDICTED: plasminogen activator inhibitor 1 RNA-binding pro
>gi|1036888259|ref|XP_017152807.1| PREDICTED: ATP-dependent RNA helicase vasa [Drosophila miran
>gi|1036888278|ref|XP_017152809.1| PREDICTED: ATP-dependent RNA helicase vasa [Drosophila miran
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>gi|1036888177|ref|XP_017152802.1| PREDICTED: anoctamin-8 isoform X2 [Drosophila miranda]
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>gi|1036888225|ref|XP_017152805.1| PREDICTED: anoctamin-8 isoform X5 [Drosophila miranda]
>gi|1036888162|ref|XP_017152801.1| PREDICTED: anoctamin-8 isoform X1 [Drosophila miranda]
>gi|1036888193|ref|XP_017152803.1| PREDICTED: anoctamin-8 isoform X3 [Drosophila miranda]
>gi|1036888242|ref|XP_017152806.1| PREDICTED: zinc finger protein 585A [Drosophila miranda]
>gi|1036888311|ref|XP_017152811.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-]
>gi|1036888329|ref|XP_017152812.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-]
>gi|1036903153|ref|XP_017153649.1| PREDICTED: protein shuttle craft isoform X1 [Drosophila mira
>gi|1036903175|ref|XP_017153650.1| PREDICTED: protein shuttle craft isoform X2 [Drosophila mira
>gi|1036925771|ref|XP_017154917.1| PREDICTED: chaoptin isoform X2 [Drosophila miranda]
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>gi|1036861956|ref|XP_017151362.1| PREDICTED: uncharacterized protein LOC108161584 [Drosophila
>gi|1036853829|ref|XP_017150909.1| PREDICTED: FK506-binding protein 3-like, partial [Drosophila
>gi|1036910923|ref|XP_017154069.1| PREDICTED: uncharacterized protein LOC108163341 [Drosophila
>gi|1036910941|ref|XP_017154070.1| PREDICTED: uncharacterized protein LOC108163341 [Drosophila
>gi|1036910961|ref|XP_017154071.1| PREDICTED: uncharacterized protein LOC108163341 [Drosophila
>gi|1036853848|ref|XP_017150910.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036910871|ref|XP_017154066.1| PREDICTED: zinc transporter 2 isoform X1 [Drosophila miranda
>gi|1036910888|ref|XP_017154067.1| PREDICTED: zinc transporter 2 isoform X2 [Drosophila miranda
>gi|1036910905|ref|XP_017154068.1| PREDICTED: zinc transporter 2 isoform X3 [Drosophila miranda
>gi|1036920695|ref|XP_017154631.1| PREDICTED: SET and MYND domain-containing protein 4 isoform
>gi|1036920662|ref|XP_017154629.1| PREDICTED: SET and MYND domain-containing protein 4 isoform
>gi|1036920678|ref|XP_017154630.1| PREDICTED: SET and MYND domain-containing protein 4 isoform
>gi|1036924485|ref|XP_017154847.1| PREDICTED: protein OS-9 [Drosophila miranda]
>gi|1036863641|ref|XP_017151449.1| PREDICTED: uncharacterized protein LOC108161662 isoform X1
>gi|1036863660|ref|XP_017151450.1| PREDICTED: uncharacterized protein LOC108161662 isoform X2
>gi|1036858052|ref|XP_017151159.1| PREDICTED: uncharacterized protein LOC108161415 [Drosophila
>gi|1036857342|ref|XP_017151119.1| PREDICTED: trypsin [Drosophila miranda]
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>gi|1036926520|ref|XP_017154959.1| PREDICTED: uncharacterized protein LOC108163935 isoform X3
>gi|1036926505|ref|XP_017154958.1| PREDICTED: uncharacterized protein LOC108163935 isoform X2
>gi|1036926536|ref|XP_017154960.1| PREDICTED: serine proteinase stubble [Drosophila miranda]
>gi|1036911686|ref|XP_017154108.1| PREDICTED: ribosomal RNA processing protein 1 homolog [Droso
>gi|1036911705|ref|XP_017154109.1| PREDICTED: glutaredoxin 3 [Drosophila miranda]
>gi|1036921018|ref|XP_017154651.1| PREDICTED: immunoglobulin-binding protein 1 [Drosophila mira
>gi|1036921036|ref|XP_017154652.1| PREDICTED: DET1- and DDB1-associated protein 1 [Drosophila m
>gi|1036921054|ref|XP_017154653.1| PREDICTED: DET1- and DDB1-associated protein 1 [Drosophila m
>gi|1036929954|ref|XP_017155144.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036921250|ref|XP_017154666.1| PREDICTED: transcription factor TFIIIB component B'' homolog
>gi|1036921268|ref|XP_017154667.1| PREDICTED: transcription factor TFIIIB component B'' homolog
>gi|1036889531|ref|XP_017152883.1| PREDICTED: uncharacterized protein LOC108162581 [Drosophila
>gi|1036889513|ref|XP_017152882.1| PREDICTED: protein disulfide-isomerase A5 [Drosophila mirand
>gi|1036889545|ref|XP_017152884.1| PREDICTED: dnaJ-like protein 60 isoform X1 [Drosophila miran
>gi|1036889564|ref|XP_017152885.1| PREDICTED: dnaJ-like protein 60 isoform X2 [Drosophila miran
>gi|1036889495|ref|XP_017152881.1| PREDICTED: coatomer subunit beta' [Drosophila miranda]
>gi|1036883616|ref|XP_017152558.1| PREDICTED: interferon-inducible double-stranded RNA-dependen
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>gi|1036883635|ref|XP_017152559.1| PREDICTED: interferon-inducible double-stranded RNA-depender
>gi|1036883654|ref|XP_017152560.1| PREDICTED: interferon-inducible double-stranded RNA-depender
>gi|1036883690|ref|XP_017152562.1| PREDICTED: inhibitor of growth protein 4 isoform X2 [Drosop]
>gi|1036883672|ref|XP_017152561.1| PREDICTED: inhibitor of growth protein 5 isoform X1 [Drosop]
>gi|1036883598|ref|XP_017152557.1| PREDICTED: uncharacterized protein LOC108162362 [Drosophila
>gi|1036883704|ref|XP_017152563.1| PREDICTED: 60S ribosomal protein L24 [Drosophila miranda]
>gi|1036931615|ref|XP_017155235.1| PREDICTED: uncharacterized protein LOC108164142 [Drosophila
>gi|1036934456|ref|XP_017155387.1| PREDICTED: dynein intermediate chain 3, ciliary [Drosophila
>gi|1036917454|ref|XP_017154447.1| PREDICTED: uncharacterized protein LOC108163585 [Drosophila
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>gi|1036859628|ref|XP_017151232.1| PREDICTED: vitelline membrane protein Vm34Ca [Drosophila mi
>gi|1036862239|ref|XP_017151378.1| PREDICTED: uncharacterized protein LOC108161601 [Drosophila
>gi|1036913482|ref|XP_017154216.1| PREDICTED: uncharacterized protein LOC108163442 [Drosophila
>gi|1036913415|ref|XP_017154211.1| PREDICTED: sialin-like [Drosophila miranda]
>gi|1036913449|ref|XP_017154213.1| PREDICTED: fibrinogen-like protein 1 [Drosophila miranda]
>gi|1036913399|ref|XP_017154210.1| PREDICTED: protein toll [Drosophila miranda]
>gi|1036913467|ref|XP_017154214.1| PREDICTED: ficolin-2-like [Drosophila miranda]
>gi|1036913431|ref|XP_017154212.1| PREDICTED: very-long-chain (3R)-3-hydroxyacyl-CoA dehydratas
>gi|1036934764|ref|XP_017155405.1| PREDICTED: 60 kDa heat shock protein homolog 2, mitochondria
>gi|1036922625|ref|XP_017154749.1| PREDICTED: uncharacterized protein LOC108163781 isoform X1
>gi|1036922639|ref|XP_017154750.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1036863569|ref|XP_017151448.1| PREDICTED: uncharacterized protein LOC108161659 [Drosophila
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>gi|1036897042|ref|XP_017153306.1| PREDICTED: angiopoietin-related protein 7-like isoform X1 []
>gi|1036897060|ref|XP_017153307.1| PREDICTED: fibrinogen-like protein 1 isoform X2 [Drosophila
>gi|1036897095|ref|XP_017153308.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop]
>gi|1036897114|ref|XP_017153309.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop]
>gi|1036897273|ref|XP_017153319.1| PREDICTED: uncharacterized protein LOC108162877 [Drosophila
>gi|1036897223|ref|XP_017153316.1| PREDICTED: uncharacterized protein LOC108162874 [Drosophila
>gi|1036897256|ref|XP_017153318.1| PREDICTED: uncharacterized protein LOC108162876 [Drosophila
>gi|1036897239|ref|XP_017153317.1| PREDICTED: uncharacterized protein LOC108162875 [Drosophila
>gi|1036897290|ref|XP_017153320.1| PREDICTED: uncharacterized protein LOC108162878 [Drosophila
>gi|1036897307|ref|XP_017153321.1| PREDICTED: uncharacterized protein LOC108162879 [Drosophila
>gi|1036897150|ref|XP_017153311.1| PREDICTED: uncharacterized protein LOC108162870 isoform X1
>gi|1036897171|ref|XP_017153312.1| PREDICTED: uncharacterized protein LOC108162870 isoform X2
>gi|1036897190|ref|XP_017153313.1| PREDICTED: uncharacterized protein LOC108162872 [Drosophila
>gi|1036897131|ref|XP_017153310.1| PREDICTED: uncharacterized protein LOC108162869 [Drosophila
>gi|1036853868|ref|XP_017150912.1| PREDICTED: 1-acylglycerol-3-phosphate O-acyltransferase ABHI
>gi|1036929016|ref|XP_017155096.1| PREDICTED: uncharacterized protein LOC108164031 [Drosophila
>gi|1036879117|ref|XP_017152309.1| PREDICTED: disintegrin and metalloproteinase domain-containi
>gi|1036879136|ref|XP_017152310.1| PREDICTED: disintegrin and metalloproteinase domain-containi
>gi|1036879077|ref|XP_017152307.1| PREDICTED: disintegrin and metalloproteinase domain-containi
>gi|1036879096|ref|XP_017152308.1| PREDICTED: disintegrin and metalloproteinase domain-containi
>gi|1036879174|ref|XP_017152312.1| PREDICTED: angiopoietin-related protein 7-like [Drosophila r
>gi|1036879200|ref|XP_017152313.1| PREDICTED: uncharacterized protein LOC108162210 [Drosophila
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>gi|1036879155|ref|XP_017152311.1| PREDICTED: ficolin-2-like [Drosophila miranda]
>gi|1036879219|ref|XP_017152315.1| PREDICTED: RING finger protein PFF0165c-like [Drosophila mi
>gi|1036926376|ref|XP_017154950.1| PREDICTED: ras suppressor protein 1 [Drosophila miranda]
>gi|1036860062|ref|XP_017151256.1| PREDICTED: ficolin-2-like isoform X1 [Drosophila miranda]
>gi|1036860081|ref|XP_017151257.1| PREDICTED: ficolin-1-like isoform X2 [Drosophila miranda]
>gi|1036927648|ref|XP_017155025.1| PREDICTED: uncharacterized protein LOC108163977 isoform X3
>gi|1036927700|ref|XP_017155027.1| PREDICTED: uncharacterized protein LOC108163977 isoform X5
>gi|1036927682|ref|XP_017155026.1| PREDICTED: uncharacterized protein LOC108163977 isoform X4
>gi|1036927630|ref|XP_017155024.1| PREDICTED: uncharacterized protein LOC108163977 isoform X2
>gi|1036927610|ref|XP_017155023.1| PREDICTED: uncharacterized protein LOC108163977 isoform X1
>gi|1036925874|ref|XP_017154923.1| PREDICTED: uncharacterized protein LOC108163907 [Drosophila
>gi|1036925856|ref|XP_017154922.1| PREDICTED: LOW QUALITY PROTEIN: glycerol-3-phosphate dehydro
>gi|1036925912|ref|XP_017154926.1| PREDICTED: ficolin-1-like [Drosophila miranda]
>gi|1036925892|ref|XP_017154925.1| PREDICTED: serine/arginine-rich splicing factor 4-like [Dros
>gi|1036857946|ref|XP_017151152.1| PREDICTED: E3 ubiquitin-protein ligase NRDP1 [Drosophila mir
>gi|1036935445|ref|XP_017155443.1| PREDICTED: cysteine and histidine-rich protein 1-like [Droso
>gi|1036863328|ref|XP_017151436.1| PREDICTED: H/ACA ribonucleoprotein complex subunit 1 [Drosop
>gi|1036857837|ref|XP_017151146.1| PREDICTED: uncharacterized protein LOC108161406 [Drosophila
>gi|1036925716|ref|XP_017154912.1| PREDICTED: uncharacterized protein LOC108163900 isoform X1
>gi|1036925730|ref|XP_017154913.1| PREDICTED: protein lev-9 isoform X2 [Drosophila miranda]
>gi|1036922535|ref|XP_017154743.1| PREDICTED: uncharacterized protein LOC108163776 [Drosophila
>gi|1036922551|ref|XP_017154744.1| PREDICTED: uncharacterized protein LOC108163776 [Drosophila
>gi|1036922565|ref|XP_017154745.1| PREDICTED: probable dolichol-phosphate mannosyltransferase
>gi|1036918384|ref|XP_017154502.1| PREDICTED: centrosomal protein of 104 kDa isoform X1 [Drosop
>gi|1036918402|ref|XP_017154503.1| PREDICTED: centrosomal protein of 104 kDa isoform X2 [Drosop
>gi|1036918438|ref|XP_017154505.1| PREDICTED: cytochrome b561 domain-containing protein 2 [Dros
>gi|1036918420|ref|XP_017154504.1| PREDICTED: centrosomal protein of 104 kDa isoform X3 [Drosop
>gi|1036913617|ref|XP_017154224.1| PREDICTED: ethanolaminephosphotransferase 1 isoform X1 [Dros
>gi|1036913635|ref|XP_017154225.1| PREDICTED: ethanolaminephosphotransferase 1 isoform X2 [Dros
>gi|1036913670|ref|XP_017154227.1| PREDICTED: uncharacterized protein LOC108163450 [Drosophila
>gi|1036913652|ref|XP_017154226.1| PREDICTED: uncharacterized protein LOC108163449 [Drosophila
>gi|1036861760|ref|XP_017151353.1| PREDICTED: uncharacterized protein LOC108161576 [Drosophila
>gi|1036889076|ref|XP_017152856.1| PREDICTED: protein ref(2)P [Drosophila miranda]
>gi|1036889094|ref|XP_017152857.1| PREDICTED: uncharacterized protein LOC108162568 [Drosophila
>gi|1036889058|ref|XP_017152855.1| PREDICTED: LOW QUALITY PROTEIN: pregnancy zone protein [Dros
>gi|1036891932|ref|XP_017153021.1| PREDICTED: equilibrative nucleoside transporter 1 [Drosophil
>gi|1036891951|ref|XP_017153022.1| PREDICTED: tRNA (adenine(58)-N(1))-methyltransferase non-cat
>gi|1036891970|ref|XP_017153023.1| PREDICTED: tRNA (adenine(58)-N(1))-methyltransferase non-cat
>gi|1036892008|ref|XP_017153025.1| PREDICTED: cytochrome c oxidase subunit 5B, mitochondrial [
>gi|1036892027|ref|XP_017153026.1| PREDICTED: cytochrome c oxidase subunit 5B, mitochondrial [
>gi|1036891989|ref|XP_017153024.1| PREDICTED: intraflagellar transport protein 52 homolog [Dros
>gi|1036891894|ref|XP_017153019.1| PREDICTED: developmental protein eyes absent isoform X1 [Dro
>gi|1036891913|ref|XP_017153020.1| PREDICTED: developmental protein eyes absent isoform X2 [Dro
>gi|1036906571|ref|XP_017153847.1| PREDICTED: male-specific sperm protein Mst84Da-like [Drosoph
>gi|1036906535|ref|XP_017153845.1| PREDICTED: carbohydrate sulfotransferase 4 isoform X1 [Droso
>gi|1036906553|ref|XP_017153846.1| PREDICTED: carbohydrate sulfotransferase 2 isoform X2 [Droso
>gi|1036896606|ref|XP_017153280.1| PREDICTED: uncharacterized protein LOC108162847 [Drosophila
>gi|1036896551|ref|XP_017153278.1| PREDICTED: coiled-coil domain-containing protein 102A isofor
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>gi|1036896532|ref|XP_017153277.1| PREDICTED: coiled-coil domain-containing protein 102A isofo
>gi|1036896588|ref|XP_017153279.1| PREDICTED: glutaryl-CoA dehydrogenase, mitochondrial [Drosop
>gi|1036896663|ref|XP_017153283.1| PREDICTED: uncharacterized protein CG9548 [Drosophila mirand
>gi|1036896625|ref|XP_017153281.1| PREDICTED: coatomer subunit epsilon [Drosophila miranda]
>gi|1036896644|ref|XP_017153282.1| PREDICTED: ATP-dependent (S)-NAD(P)H-hydrate dehydratase [Di
>gi|1036909387|ref|XP_017153994.1| PREDICTED: kynurenine formamidase [Drosophila miranda]
>gi|1036909405|ref|XP_017153995.1| PREDICTED: kynurenine formamidase [Drosophila miranda]
>gi|1036909351|ref|XP_017153991.1| PREDICTED: proteoglycan 4 isoform X2 [Drosophila miranda]
>gi|1036909330|ref|XP_017153990.1| PREDICTED: proteoglycan 4 isoform X1 [Drosophila miranda]
>gi|1036909369|ref|XP_017153993.1| PREDICTED: proteoglycan 4 isoform X3 [Drosophila miranda]
>gi|1036910251|ref|XP_017154033.1| PREDICTED: protein transport protein Sec61 subunit alpha [Di
>gi|1036910271|ref|XP_017154035.1| PREDICTED: transmembrane protein 115 [Drosophila miranda]
>gi|1036900996|ref|XP_017153532.1| PREDICTED: UDP-N-acetylhexosamine pyrophosphorylase-like pro
>gi|1036901013|ref|XP_017153533.1| PREDICTED: UDP-N-acetylhexosamine pyrophosphorylase isoform
>gi|1036900976|ref|XP_017153531.1| PREDICTED: UDP-N-acetylhexosamine pyrophosphorylase isoform
>gi|1036901102|ref|XP_017153538.1| PREDICTED: hemK methyltransferase family member 1 [Drosophil
>gi|1036901051|ref|XP_017153535.1| PREDICTED: RING finger and CHY zinc finger domain-containing
>gi|1036901067|ref|XP_017153536.1| PREDICTED: RING finger and CHY zinc finger domain-containing
>gi|1036901084|ref|XP_017153537.1| PREDICTED: RING finger and CHY zinc finger domain-containing
>gi|1036901033|ref|XP_017153534.1| PREDICTED: lysophospholipid acyltransferase 7 [Drosophila mi
>gi|1036928781|ref|XP_017155081.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036928799|ref|XP_017155082.1| PREDICTED: phospholipase A1 2 [Drosophila miranda]
>gi|1036928853|ref|XP_017155086.1| PREDICTED: dynein light chain roadblock-type 1 [Drosophila m
>gi|1036928835|ref|XP_017155085.1| PREDICTED: metacaspase-1 [Drosophila miranda]
>gi|1036928817|ref|XP_017155083.1| PREDICTED: uncharacterized protein LOC108164019 [Drosophila
>gi|1036932475|ref|XP_017155278.1| PREDICTED: RNA polymerase-associated protein Rtf1 isoform X
>gi|1036932493|ref|XP_017155279.1| PREDICTED: RNA polymerase-associated protein Rtf1 isoform X2
>gi|1036932510|ref|XP_017155281.1| PREDICTED: uncharacterized protein LOC108164175 [Drosophila
>gi|1036931873|ref|XP_017155246.1| PREDICTED: BTB/POZ domain-containing protein 9 isoform X1 [I
>gi|1036931891|ref|XP_017155247.1| PREDICTED: uncharacterized protein LOC108164154 isoform X2
>gi|1036931908|ref|XP_017155249.1| PREDICTED: uncharacterized protein LOC108164154 isoform X2
>gi|1036931925|ref|XP_017155250.1| PREDICTED: uncharacterized protein LOC108164154 isoform X2
>gi|1036927238|ref|XP_017154999.1| PREDICTED: transcription initiation factor TFIID subunit 5
>gi|1036927182|ref|XP_017154996.1| PREDICTED: uncharacterized protein LOC108163958 isoform X1
>gi|1036927218|ref|XP_017154998.1| PREDICTED: uncharacterized protein LOC108163958 isoform X3
>gi|1036927200|ref|XP_017154997.1| PREDICTED: uncharacterized protein LOC108163958 isoform X2
>gi|1036927258|ref|XP_017155000.1| PREDICTED: proteasome subunit alpha type-7-1-like isoform X
>gi|1036927276|ref|XP_017155001.1| PREDICTED: proteasome subunit alpha type-7-1-like isoform X2
>gi|1036853894|ref|XP_017150913.1| PREDICTED: uncharacterized protein LOC108161202 [Drosophila
>gi|1036853913|ref|XP_017150914.1| PREDICTED: uncharacterized protein LOC108161203 [Drosophila
>gi|1036927383|ref|XP_017155009.1| PREDICTED: protein tiptop [Drosophila miranda]
>gi|1036927401|ref|XP_017155010.1| PREDICTED: protein tiptop [Drosophila miranda]
>gi|1036881875|ref|XP_017152461.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036853932|ref|XP_017150915.1| PREDICTED: uncharacterized protein LOC108161205 [Drosophila
>gi|1036853949|ref|XP_017150916.1| PREDICTED: uncharacterized protein LOC108161206 [Drosophila
>gi|1036881827|ref|XP_017152457.1| PREDICTED: uncharacterized protein LOC108162302 [Drosophila
>gi|1036881892|ref|XP_017152462.1| PREDICTED: uncharacterized protein LOC108162305 [Drosophila
>gi|1036881817|ref|XP_017152456.1| PREDICTED: uncharacterized protein LOC108162301 [Drosophila
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>gi|1036881930|ref|XP_017152464.1| PREDICTED: uncharacterized protein LOC108162307 [Drosophila
>gi|1036881947|ref|XP_017152465.1| PREDICTED: uncharacterized protein LOC108162307 [Drosophila
>gi|1036881911|ref|XP_017152463.1| PREDICTED: uncharacterized protein LOC108162306 [Drosophila
>gi|1036853968|ref|XP_017150917.1| PREDICTED: ER membrane protein complex subunit 7 homolog [D
>gi|1036881802|ref|XP_017152455.1| PREDICTED: protein teashirt [Drosophila miranda]
>gi|1036853987|ref|XP_017150919.1| PREDICTED: uncharacterized protein LOC108161208 [Drosophila
>gi|1036881783|ref|XP_017152454.1| PREDICTED: zinc finger protein 700 [Drosophila miranda]
>gi|1036881838|ref|XP_017152459.1| PREDICTED: activator of 90 kDa heat shock protein ATPase hom
>gi|1036881856|ref|XP_017152460.1| PREDICTED: activator of 90 kDa heat shock protein ATPase hom
>gi|1036893331|ref|XP_017153101.1| PREDICTED: cytohesin-1 isoform X2 [Drosophila miranda]
>gi|1036893349|ref|XP_017153102.1| PREDICTED: cytohesin-1 isoform X3 [Drosophila miranda]
>gi|1036893314|ref|XP_017153100.1| PREDICTED: ankyrin repeat, PH and SEC7 domain containing pro
>gi|1036892851|ref|XP_017153070.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase al
>gi|1036892870|ref|XP_017153072.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase al
>gi|1036892811|ref|XP_017153068.1| PREDICTED: zinc finger protein 708 [Drosophila miranda]
>gi|1036892689|ref|XP_017153064.1| PREDICTED: myosin-G heavy chain isoform X1 [Drosophila miran
>gi|1036892711|ref|XP_017153065.1| PREDICTED: unconventional myosin-IXb isoform X2 [Drosophila
>gi|1036892730|ref|XP_017153066.1| PREDICTED: unconventional myosin-IXb isoform X2 [Drosophila
>gi|1036892792|ref|XP_017153067.1| PREDICTED: unconventional myosin-IXb isoform X2 [Drosophila
>gi|1036892889|ref|XP_017153073.1| PREDICTED: uncharacterized protein LOC108162719 [Drosophila
>gi|1036892907|ref|XP_017153074.1| PREDICTED: uncharacterized protein LOC108162720 [Drosophila
>gi|1036892963|ref|XP_017153077.1| PREDICTED: diuretic hormone class 2 isoform X2 [Drosophila m
>gi|1036892926|ref|XP_017153075.1| PREDICTED: diuretic hormone class 2 isoform X1 [Drosophila m
>gi|1036892945|ref|XP_017153076.1| PREDICTED: diuretic hormone class 2 isoform X1 [Drosophila m
>gi|1036892832|ref|XP_017153069.1| PREDICTED: lysosomal aspartic protease [Drosophila miranda]
>gi|1036916436|ref|XP_017154391.1| PREDICTED: ribosomal L1 domain-containing protein CG13096 [D
>gi|1036875902|ref|XP_017152141.1| PREDICTED: U3 small nucleolar ribonucleoprotein protein MPP1
>gi|1036875973|ref|XP_017152144.1| PREDICTED: phosphatidylinositol glycan anchor biosynthesis c
>gi|1036876011|ref|XP_017152146.1| PREDICTED: 39S ribosomal protein L51, mitochondrial [Drosoph
>gi|1036875952|ref|XP_017152143.1| PREDICTED: adenylyltransferase and sulfurtransferase MOCS3 [
>gi|1036875933|ref|XP_017152142.1| PREDICTED: angiotensin-converting enzyme-related protein [D
>gi|1036875992|ref|XP_017152145.1| PREDICTED: homeobox protein Hmx [Drosophila miranda]
>gi|1036876028|ref|XP_017152147.1| PREDICTED: anaphase-promoting complex subunit 11 [Drosophila
>gi|1036902641|ref|XP_017153622.1| PREDICTED: uncharacterized protein LOC108163062 [Drosophila
>gi|1036902659|ref|XP_017153623.1| PREDICTED: uncharacterized protein LOC108163062 [Drosophila
>gi|1036902677|ref|XP_017153625.1| PREDICTED: probable phosphomevalonate kinase [Drosophila mir
>gi|1036902623|ref|XP_017153621.1| PREDICTED: E3 ubiquitin-protein ligase Hakai [Drosophila mir
>gi|1036856441|ref|XP_017151067.1| PREDICTED: proliferating cell nuclear antigen [Drosophila mi
>gi|1036907230|ref|XP_017153881.1| PREDICTED: probable RNA helicase armi [Drosophila miranda]
>gi|1036923716|ref|XP_017154803.1| PREDICTED: cyclin-dependent kinases regulatory subunit [Dros
>gi|1036927788|ref|XP_017155033.1| PREDICTED: excitatory amino acid transporter 1 [Drosophila m
>gi|1036927806|ref|XP_017155034.1| PREDICTED: excitatory amino acid transporter 1 [Drosophila m
>gi|1036927824|ref|XP_017155035.1| PREDICTED: excitatory amino acid transporter 1 [Drosophila m
>gi|1036934561|ref|XP_017155394.1| PREDICTED: uncharacterized protein LOC108164263 [Drosophila
>gi|1036854005|ref|XP_017150920.1| PREDICTED: uncharacterized protein LOC108161209 [Drosophila
>gi|1036854022|ref|XP_017150921.1| PREDICTED: uncharacterized protein LOC108161210 [Drosophila
>gi|1036863311|ref|XP_017151435.1| PREDICTED: trypsin inhibitor [Drosophila miranda]
>gi|1036860155|ref|XP_017151261.1| PREDICTED: uncharacterized protein LOC108161497 [Drosophila
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>gi|1036917277|ref|XP_017154436.1| PREDICTED: potassium voltage-gated channel protein Shaw [Dro
>gi|1036917314|ref|XP_017154438.1| PREDICTED: aldehyde dehydrogenase, mitochondrial-like [Dros
>gi|1036917294|ref|XP_017154437.1| PREDICTED: aldehyde dehydrogenase, mitochondrial-like [Dros
>gi|1036921613|ref|XP_017154686.1| PREDICTED: laccase-2 [Drosophila miranda]
>gi|1036917108|ref|XP_017154425.1| PREDICTED: fat body protein 2 [Drosophila miranda]
>gi|1036917072|ref|XP_017154423.1| PREDICTED: alpha-(1,3)-fucosyltransferase B-like [Drosophila
>gi|1036917090|ref|XP_017154424.1| PREDICTED: uncharacterized protein LOC108163568 [Drosophila
>gi|1036854039|ref|XP_017150922.1| PREDICTED: alpha-(1,3)-fucosyltransferase B-like [Drosophila
>gi|1036919650|ref|XP_017154567.1| PREDICTED: alpha-(1,3)-fucosyltransferase B-like [Drosophila
>gi|1036888487|ref|XP_017152821.1| PREDICTED: vacuolar protein sorting-associated protein 54 [
>gi|1036888520|ref|XP_017152823.1| PREDICTED: cytoplasmic dynein 2 light intermediate chain 1
>gi|1036888537|ref|XP_017152825.1| PREDICTED: 40S ribosomal protein S28 [Drosophila miranda]
>gi|1036888502|ref|XP_017152822.1| PREDICTED: protein numb [Drosophila miranda]
>gi|1036909519|ref|XP_017154002.1| PREDICTED: rab GDP dissociation inhibitor alpha [Drosophila
>gi|1036909537|ref|XP_017154003.1| PREDICTED: putative nuclease HARBI1 [Drosophila miranda]
>gi|1036901307|ref|XP_017153551.1| PREDICTED: probable phospholipid-transporting ATPase VA iso
>gi|1036901325|ref|XP_017153552.1| PREDICTED: probable phospholipid-transporting ATPase VA iso
>gi|1036901347|ref|XP_017153553.1| PREDICTED: probable phospholipid-transporting ATPase VA iso
>gi|1036903963|ref|XP_017153695.1| PREDICTED: solute carrier organic anion transporter family
>gi|1036903983|ref|XP_017153696.1| PREDICTED: solute carrier organic anion transporter family
>gi|1036904001|ref|XP_017153697.1| PREDICTED: solute carrier organic anion transporter family
>gi|1036904019|ref|XP_017153698.1| PREDICTED: solute carrier organic anion transporter family
>gi|1036904037|ref|XP_017153699.1| PREDICTED: solute carrier organic anion transporter family
>gi|1036921989|ref|XP_017154709.1| PREDICTED: transmembrane and coiled-coil domain-containing
>gi|1036922007|ref|XP_017154710.1| PREDICTED: transmembrane and coiled-coil domain-containing
>gi|1036921973|ref|XP_017154708.1| PREDICTED: FAD-dependent oxidoreductase domain-containing pr
>gi|1036875534|ref|XP_017152119.1| PREDICTED: probable 26S proteasome non-ATPase regulatory su
>gi|1036875475|ref|XP_017152115.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036875493|ref|XP_017152116.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036875517|ref|XP_017152118.1| PREDICTED: trithorax group protein osa isoform X2 [Drosophil
>gi|1036875455|ref|XP_017152114.1| PREDICTED: protein phosphatase PHLPP-like protein [Drosophil
>gi|1036875585|ref|XP_017152122.1| PREDICTED: tRNA-dihydrouridine(20a/20b) synthase [NAD(P)+]-l
>gi|1036875548|ref|XP_017152120.1| PREDICTED: protein CNPPD1 isoform X1 [Drosophila miranda]
>gi|1036875567|ref|XP_017152121.1| PREDICTED: protein CNPPD1 isoform X2 [Drosophila miranda]
>gi|1036875621|ref|XP_017152124.1| PREDICTED: transmembrane protein 65 isoform X2 [Drosophila
>gi|1036875602|ref|XP_017152123.1| PREDICTED: uncharacterized protein C03B8.3 isoform X1 [Dros
>gi|1036875640|ref|XP_017152126.1| PREDICTED: general odorant-binding protein 28a [Drosophila
>gi|1036854057|ref|XP_017150923.1| PREDICTED: LOW QUALITY PROTEIN: CD209 antigen-like protein
>gi|1036925303|ref|XP_017154888.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036901685|ref|XP_017153570.1| PREDICTED: caltractin [Drosophila miranda]
>gi|1036901667|ref|XP_017153569.1| PREDICTED: uncharacterized protein LOC108163032 isoform X2
>gi|1036901612|ref|XP_017153566.1| PREDICTED: uncharacterized protein LOC108163032 isoform X1
>gi|1036901629|ref|XP_017153567.1| PREDICTED: uncharacterized protein LOC108163032 isoform X1
>gi|1036901647|ref|XP_017153568.1| PREDICTED: uncharacterized protein LOC108163032 isoform X1
>gi|1036861242|ref|XP_017151321.1| PREDICTED: uncharacterized protein LOC108161550 [Drosophila
>gi|1036854075|ref|XP_017150924.1| PREDICTED: uncharacterized protein LOC108161213 [Drosophila
>gi|1036854094|ref|XP_017150925.1| PREDICTED: uncharacterized protein LOC108161214 [Drosophila
>gi|1036923678|ref|XP_017154801.1| PREDICTED: lipase 3 [Drosophila miranda]
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>gi|1036923696|ref|XP_017154802.1| PREDICTED: C-type lectin domain family 4 member M-like [Dros
>gi|1036915082|ref|XP_017154306.1| PREDICTED: zinc finger protein 239 isoform X1 [Drosophila m:
>gi|1036915098|ref|XP_017154307.1| PREDICTED: zinc finger protein 239 isoform X1 [Drosophila m:
>gi|1036915113|ref|XP_017154308.1| PREDICTED: zinc finger protein 2 homolog isoform X2 [Drosop]
>gi|1036916177|ref|XP_017154375.1| PREDICTED: uncharacterized protein LOC108163533 [Drosophila
>gi|1036916157|ref|XP_017154374.1| PREDICTED: uncharacterized protein LOC108163532 [Drosophila
>gi|1036887794|ref|XP_017152786.1| PREDICTED: nuclear transcription factor Y subunit B-1 [Dros(
>gi|1036887776|ref|XP_017152785.1| PREDICTED: uncharacterized protein LOC108162515 [Drosophila
>gi|1036906363|ref|XP_017153835.1| PREDICTED: GTP-binding protein 10 homolog [Drosophila mirand
>gi|1036906345|ref|XP_017153834.1| PREDICTED: tRNA-dihydrouridine(47) synthase [NAD(P)(+)]-like
>gi|1036906399|ref|XP_017153837.1| PREDICTED: fructose-1,6-bisphosphatase 1 isoform X2 [Drosop]
>gi|1036906381|ref|XP_017153836.1| PREDICTED: fructose-1,6-bisphosphatase 1 isoform X1 [Drosop]
>gi|1036906417|ref|XP_017153838.1| PREDICTED: fructose-1,6-bisphosphatase 1 isoform X3 [Drosop]
>gi|1036860524|ref|XP_017151280.1| PREDICTED: brain-specific homeobox protein [Drosophila mirar
>gi|1036912229|ref|XP_017154139.1| PREDICTED: coronin-6-like isoform X1 [Drosophila miranda]
>gi|1036912247|ref|XP_017154140.1| PREDICTED: coronin-6-like isoform X2 [Drosophila miranda]
>gi|1036931354|ref|XP_017155218.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1036931372|ref|XP_017155219.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1036931390|ref|XP_017155220.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1036931408|ref|XP_017155221.1| PREDICTED: myb-like protein Q [Drosophila miranda]
>gi|1036907806|ref|XP_017153909.1| PREDICTED: uncharacterized protein LOC108163253 isoform X1
>gi|1036907824|ref|XP_017153910.1| PREDICTED: uncharacterized protein LOC108163253 isoform X2
>gi|1036907842|ref|XP_017153911.1| PREDICTED: uncharacterized protein LOC108163253 isoform X3
>gi|1036907860|ref|XP_017153912.1| PREDICTED: uncharacterized protein LOC108163253 isoform X4
>gi|1036927770|ref|XP_017155032.1| PREDICTED: uncharacterized protein LOC108163982 [Drosophila
>gi|1036854110|ref|XP_017150927.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1036854129|ref|XP_017150928.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1036854146|ref|XP_017150929.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1036913257|ref|XP_017154201.1| PREDICTED: RNA-binding motif protein, X-linked 2 [Drosophila
>gi|1036913223|ref|XP_017154199.1| PREDICTED: probable RNA polymerase II nuclear localization p
>gi|1036913239|ref|XP_017154200.1| PREDICTED: uncharacterized protein LOC108163429 isoform X2
>gi|1036921431|ref|XP_017154675.1| PREDICTED: UV radiation resistance-associated gene protein
>gi|1036901367|ref|XP_017153554.1| PREDICTED: probable phospholipid-transporting ATPase IIA isc
>gi|1036901384|ref|XP_017153555.1| PREDICTED: probable phospholipid-transporting ATPase IIB isc
>gi|1036901402|ref|XP_017153556.1| PREDICTED: uncharacterized protein LOC108163026 [Drosophila
>gi|1036854162|ref|XP_017150930.1| PREDICTED: uncharacterized protein LOC108161218 [Drosophila
>gi|1036912938|ref|XP_017154181.1| PREDICTED: uncharacterized protein LOC108163423 [Drosophila
>gi|1036912920|ref|XP_017154180.1| PREDICTED: uncharacterized protein LOC108163422 [Drosophila
>gi|1036912956|ref|XP_017154182.1| PREDICTED: GTPase-activating protein CdGAPr-like [Drosophila
>gi|1036934858|ref|XP_017155410.1| PREDICTED: uncharacterized protein LOC108164275 [Drosophila
>gi|1036854177|ref|XP_017150931.1| PREDICTED: protein toll-like [Drosophila miranda]
>gi|1036854196|ref|XP_017150932.1| PREDICTED: protein toll-like [Drosophila miranda]
>gi|1036890838|ref|XP_017152954.1| PREDICTED: elongation factor 2 [Drosophila miranda]
>gi|1036890798|ref|XP_017152952.1| PREDICTED: acyl-CoA synthetase family member 4 homolog isofc
>gi|1036890817|ref|XP_017152953.1| PREDICTED: acyl-CoA synthetase family member 4 homolog isofc
>gi|1036932992|ref|XP_017155309.1| PREDICTED: keratin, ultra high-sulfur matrix protein [Drosop
>gi|1036933240|ref|XP_017155323.1| PREDICTED: D-aspartate oxidase [Drosophila miranda]
>gi|1036933260|ref|XP_017155324.1| PREDICTED: uncharacterized protein LOC108164205 [Drosophila
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>gi|1036931165|ref|XP_017155208.1| PREDICTED: zinc finger protein sens isoform X1 [Drosophila r
>gi|1036931180|ref|XP_017155209.1| PREDICTED: zinc finger protein sens isoform X2 [Drosophila r
>gi|1036923326|ref|XP_017154792.1| PREDICTED: uncharacterized protein LOC108163813 [Drosophila
>gi|1036869809|ref|XP_017151793.1| PREDICTED: HEAT repeat-containing protein 1 homolog [Drosoph
>gi|1036869881|ref|XP_017151797.1| PREDICTED: mitochondrial sodium/hydrogen exchanger 9B2 isofo
>gi|1036869900|ref|XP_017151799.1| PREDICTED: mitochondrial sodium/hydrogen exchanger 9B2 isofo
>gi|1036869918|ref|XP_017151800.1| PREDICTED: mitochondrial sodium/hydrogen exchanger 9B2 isofo
>gi|1036869826|ref|XP_017151794.1| PREDICTED: uncharacterized protein LOC108161906 [Drosophila
>gi|1036869845|ref|XP_017151795.1| PREDICTED: uncharacterized protein LOC108161906 [Drosophila
>gi|1036869937|ref|XP_017151801.1| PREDICTED: DNA-directed RNA polymerase I subunit RPA43 [Dros
>gi|1036869864|ref|XP_017151796.1| PREDICTED: 5'-3' exoribonuclease 2 homolog [Drosophila miran
>gi|1036873427|ref|XP_017152001.1| PREDICTED: wee1-like protein kinase [Drosophila miranda]
>gi|1036873501|ref|XP_017152005.1| PREDICTED: probable splicing factor, arginine/serine-rich 6
>gi|1036873484|ref|XP_017152004.1| PREDICTED: probable splicing factor, arginine/serine-rich 6
>gi|1036873446|ref|XP_017152002.1| PREDICTED: nucleolar protein 58 [Drosophila miranda]
>gi|1036873465|ref|XP_017152003.1| PREDICTED: heterogeneous nuclear ribonucleoprotein 27C [Dros
>gi|1036900334|ref|XP_017153494.1| PREDICTED: uncharacterized protein LOC108162992 [Drosophila
>gi|1036900263|ref|XP_017153489.1| PREDICTED: myosin-4 isoform X2 [Drosophila miranda]
>gi|1036900281|ref|XP_017153490.1| PREDICTED: interaptin isoform X3 [Drosophila miranda]
>gi|1036900227|ref|XP_017153487.1| PREDICTED: myosin-4 isoform X1 [Drosophila miranda]
>gi|1036900246|ref|XP_017153488.1| PREDICTED: myosin-4 isoform X1 [Drosophila miranda]
>gi|1036900297|ref|XP_017153492.1| PREDICTED: circulating cathodic antigen homolog [Drosophila
>gi|1036900316|ref|XP_017153493.1| PREDICTED: circulating cathodic antigen homolog [Drosophila
>gi|1036913994|ref|XP_017154246.1| PREDICTED: LOW QUALITY PROTEIN: interference hedgehog [Droso
>gi|1036876695|ref|XP_017152182.1| PREDICTED: LOW QUALITY PROTEIN: YEATS domain-containing prot
>gi|1036876548|ref|XP_017152173.1| PREDICTED: cohesin subunit SA-1 [Drosophila miranda]
>gi|1036876640|ref|XP_017152179.1| PREDICTED: zinc finger BED domain-containing protein 1 isofo
>gi|1036876621|ref|XP_017152178.1| PREDICTED: zinc finger BED domain-containing protein 1 isofo
>gi|1036876566|ref|XP_017152175.1| PREDICTED: mini-chromosome maintenance complex-binding prote
>gi|1036876659|ref|XP_017152180.1| PREDICTED: ZZ-type zinc finger-containing protein 3 [Drosoph
>gi|1036876676|ref|XP_017152181.1| PREDICTED: hydroxymethylglutaryl-CoA lyase, mitochondrial [D
>gi|1036876584|ref|XP_017152176.1| PREDICTED: uncharacterized protein LOC108162115 isoform X1
>gi|1036876602|ref|XP_017152177.1| PREDICTED: uncharacterized protein LOC108162115 isoform X2
>gi|1036924344|ref|XP_017154839.1| PREDICTED: oxygen-dependent coproporphyrinogen-III oxidase
>gi|1036877571|ref|XP_017152225.1| PREDICTED: small ubiquitin-related modifier 2 [Drosophila mi
>gi|1036877497|ref|XP_017152221.1| PREDICTED: U1 small nuclear ribonucleoprotein 70 kDa [Drosop
>gi|1036877461|ref|XP_017152218.1| PREDICTED: protein cup [Drosophila miranda]
>gi|1036877534|ref|XP_017152223.1| PREDICTED: probable galactose-1-phosphate uridylyltransferas
>gi|1036877516|ref|XP_017152222.1| PREDICTED: serpin B3 [Drosophila miranda]
>gi|1036877590|ref|XP_017152226.1| PREDICTED: uncharacterized protein LOC108162159 [Drosophila
>gi|1036877479|ref|XP_017152220.1| PREDICTED: protein cortex [Drosophila miranda]
>gi|1036877553|ref|XP_017152224.1| PREDICTED: non-structural maintenance of chromosomes element
>gi|1036909945|ref|XP_017154016.1| PREDICTED: sodium/hydrogen exchanger 9 isoform X3 [Drosophil
>gi|1036910038|ref|XP_017154021.1| PREDICTED: sodium/hydrogen exchanger 6 isoform X8 [Drosophil
>gi|1036909983|ref|XP_017154018.1| PREDICTED: sodium/hydrogen exchanger 6 isoform X5 [Drosophil
>gi|1036909965|ref|XP_017154017.1| PREDICTED: sodium/hydrogen exchanger 7 isoform X4 [Drosophil
>gi|1036909728|ref|XP_017154014.1| PREDICTED: sodium/hydrogen exchanger 9 isoform X1 [Drosophil
>gi|1036910001|ref|XP_017154019.1| PREDICTED: sodium/hydrogen exchanger 9 isoform X6 [Drosophil
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>gi|1036909746|ref|XP_017154015.1| PREDICTED: sodium/hydrogen exchanger 9 isoform X2 [Drosophil
>gi|1036910019|ref|XP_017154020.1| PREDICTED: sodium/hydrogen exchanger 6 isoform X7 [Drosophil
>gi|1036910060|ref|XP_017154023.1| PREDICTED: uncharacterized protein LOC108163311 [Drosophila
>gi|1036910080|ref|XP_017154024.1| PREDICTED: WD repeat-containing protein 24-like [Drosophila
>gi|1036910098|ref|XP_017154025.1| PREDICTED: WD repeat-containing protein 24-like [Drosophila
>gi|1036854212|ref|XP_017150934.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036917545|ref|XP_017154453.1| PREDICTED: acyl-coenzyme A thioesterase 9, mitochondrial-lil
>gi|1036917491|ref|XP_017154449.1| PREDICTED: ubiquitin conjugation factor E4 A [Drosophila mil
>gi|1036917509|ref|XP_017154450.1| PREDICTED: ubiquitin conjugation factor E4 A [Drosophila mil
>gi|1036917527|ref|XP_017154452.1| PREDICTED: ubiquitin conjugation factor E4 A [Drosophila mil
>gi|1036934800|ref|XP_017155407.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036934818|ref|XP_017155408.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036934836|ref|XP_017155409.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036908876|ref|XP_017153964.1| PREDICTED: uncharacterized protein LOC108163275 isoform X1
>gi|1036908894|ref|XP_017153965.1| PREDICTED: uncharacterized protein LOC108163275 isoform X2
>gi|1036908833|ref|XP_017153963.1| PREDICTED: transport and Golgi organization protein 1 [Drosc
>gi|1036906453|ref|XP_017153841.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036906479|ref|XP_017153842.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036906499|ref|XP_017153843.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036906517|ref|XP_017153844.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036929792|ref|XP_017155138.1| PREDICTED: solute carrier organic anion transporter family r
>gi|1036929810|ref|XP_017155139.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1036883158|ref|XP_017152529.1| PREDICTED: heme oxygenase 2-like [Drosophila miranda]
>gi|1036854230|ref|XP_017150935.1| PREDICTED: uncharacterized protein LOC108161222 [Drosophila
>gi|1036883046|ref|XP_017152523.1| PREDICTED: neurotrimin isoform X1 [Drosophila miranda]
>gi|1036883063|ref|XP_017152524.1| PREDICTED: neurotrimin isoform X2 [Drosophila miranda]
>gi|1036883082|ref|XP_017152525.1| PREDICTED: uncharacterized protein LOC108162341 [Drosophila
>gi|1036883101|ref|XP_017152526.1| PREDICTED: uncharacterized protein LOC108162341 [Drosophila
>gi|1036883214|ref|XP_017152533.1| PREDICTED: maf-like protein aq_1718 [Drosophila miranda]
>gi|1036883027|ref|XP_017152521.1| PREDICTED: argininosuccinate lyase [Drosophila miranda]
>gi|1036883119|ref|XP_017152527.1| PREDICTED: tetraspanin-6 [Drosophila miranda]
>gi|1036883139|ref|XP_017152528.1| PREDICTED: tetraspanin-6 [Drosophila miranda]
>gi|1036882991|ref|XP_017152519.1| PREDICTED: tubulin glycylase 3B-like [Drosophila miranda]
>gi|1036883010|ref|XP_017152520.1| PREDICTED: tubulin glycylase 3B-like [Drosophila miranda]
>gi|1036883177|ref|XP_017152530.1| PREDICTED: CD63 antigen [Drosophila miranda]
>gi|1036883195|ref|XP_017152532.1| PREDICTED: CD63 antigen [Drosophila miranda]
>gi|1036923898|ref|XP_017154814.1| PREDICTED: uncharacterized protein LOC108163827 isoform X7
>gi|1036923808|ref|XP_017154809.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036923862|ref|XP_017154812.1| PREDICTED: serine-rich adhesin for platelets isoform X5 [Dro
>gi|1036923826|ref|XP_017154810.1| PREDICTED: uncharacterized protein LOC108163827 isoform X3
>gi|1036923844|ref|XP_017154811.1| PREDICTED: serine-rich adhesin for platelets isoform X4 [Dro
>gi|1036923736|ref|XP_017154804.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036923754|ref|XP_017154805.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036923772|ref|XP_017154807.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036923790|ref|XP_017154808.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036923916|ref|XP_017154815.1| PREDICTED: serine-rich adhesin for platelets isoform X8 [Dro
>gi|1036923880|ref|XP_017154813.1| PREDICTED: serine-rich adhesin for platelets isoform X6 [Dro
>gi|1036863495|ref|XP_017151445.1| PREDICTED: uncharacterized protein LOC108161654 [Drosophila
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>gi|1036935754|ref|XP_017155463.1| PREDICTED: uncharacterized protein LOC108164320 [Drosophila
>gi|1036935771|ref|XP_017155464.1| PREDICTED: uncharacterized protein LOC108164321 [Drosophila
>gi|1036854248|ref|XP_017150936.1| PREDICTED: ATP-binding cassette sub-family A member 3 [Droso
>gi|1036930521|ref|XP_017155173.1| PREDICTED: uncharacterized protein LOC108164096 [Drosophila
>gi|1036859683|ref|XP_017151235.1| PREDICTED: uncharacterized protein LOC108161477 [Drosophila
>gi|1036920157|ref|XP_017154597.1| PREDICTED: LOW QUALITY PROTEIN: GATA zinc finger domain-cont
>gi|1036929259|ref|XP_017155108.1| PREDICTED: lysosomal alpha-mannosidase [Drosophila miranda]
>gi|1036854267|ref|XP_017150937.1| PREDICTED: lysosomal alpha-mannosidase-like, partial [Drosop
>gi|1036928941|ref|XP_017155091.1| PREDICTED: lysosomal alpha-mannosidase [Drosophila miranda]
>gi|1036928887|ref|XP_017155088.1| PREDICTED: lysosomal alpha-mannosidase-like isoform X1 [Dros
>gi|1036928923|ref|XP_017155090.1| PREDICTED: lysosomal alpha-mannosidase-like isoform X3 [Dros
>gi|1036928905|ref|XP_017155089.1| PREDICTED: lysosomal alpha-mannosidase-like isoform X2 [Dros
>gi|1036859850|ref|XP_017151244.1| PREDICTED: uncharacterized protein LOC108161484 [Drosophila
>gi|1036859050|ref|XP_017151199.1| PREDICTED: uncharacterized protein LOC108161448 [Drosophila
>gi|1036906202|ref|XP_017153826.1| PREDICTED: uncharacterized protein LOC108163193 isoform X4
>gi|1036906148|ref|XP_017153823.1| PREDICTED: uncharacterized protein LOC108163193 isoform X1
>gi|1036906182|ref|XP_017153825.1| PREDICTED: uncharacterized protein LOC108163193 isoform X3
>gi|1036906164|ref|XP_017153824.1| PREDICTED: uncharacterized protein LOC108163193 isoform X2
>gi|1036906220|ref|XP_017153827.1| PREDICTED: uncharacterized protein LOC108163193 isoform X5
>gi|1036913782|ref|XP_017154233.1| PREDICTED: catalase [Drosophila miranda]
>gi|1036913764|ref|XP_017154232.1| PREDICTED: transcription factor HNF-4 homolog isoform X3 [D:
>gi|1036913746|ref|XP_017154231.1| PREDICTED: transcription factor HNF-4 homolog isoform X2 [D:
>gi|1036913726|ref|XP_017154230.1| PREDICTED: transcription factor HNF-4 homolog isoform X1 [D:
>gi|1036928869|ref|XP_017155087.1| PREDICTED: trafficking protein particle complex subunit 4 [I
>gi|1036924736|ref|XP_017154861.1| PREDICTED: probable cGMP 3',5'-cyclic phosphodiesterase subu
>gi|1036924718|ref|XP_017154860.1| PREDICTED: glutactin [Drosophila miranda]
>gi|1036854286|ref|XP_017150938.1| PREDICTED: accessory gland protein Acp29AB-like [Drosophila
>gi|1036854304|ref|XP_017150939.1| PREDICTED: glutactin [Drosophila miranda]
>gi|1036885830|ref|XP_017152677.1| PREDICTED: glutactin [Drosophila miranda]
>gi|1036885941|ref|XP_017152682.1| PREDICTED: putative uncharacterized protein DDB_G0290521 [Di
>gi|1036885903|ref|XP_017152680.1| PREDICTED: semaphorin-1A isoform X3 [Drosophila miranda]
>gi|1036885868|ref|XP_017152679.1| PREDICTED: semaphorin-1A isoform X2 [Drosophila miranda]
>gi|1036885849|ref|XP_017152678.1| PREDICTED: semaphorin-1A isoform X1 [Drosophila miranda]
>gi|1036885922|ref|XP_017152681.1| PREDICTED: larval serum protein 1 gamma chain [Drosophila mi
>gi|1036885960|ref|XP_017152683.1| PREDICTED: immune-induced peptide 3 [Drosophila miranda]
>gi|1036910811|ref|XP_017154063.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036910829|ref|XP_017154064.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036935429|ref|XP_017155442.1| PREDICTED: cytochrome b561 domain-containing protein 2 [Dros
>gi|1036862220|ref|XP_017151376.1| PREDICTED: uncharacterized protein LOC108161600 [Drosophila
>gi|1036926724|ref|XP_017154971.1| PREDICTED: uncharacterized protein LOC108163943 [Drosophila
>gi|1036925445|ref|XP_017154895.1| PREDICTED: uncharacterized protein LOC108163886 isoform X4
>gi|1036925427|ref|XP_017154894.1| PREDICTED: uncharacterized protein LOC108163886 isoform X3
>gi|1036925410|ref|XP_017154893.1| PREDICTED: uncharacterized protein LOC108163886 isoform X2
>gi|1036925374|ref|XP_017154891.1| PREDICTED: uncharacterized protein LOC108163886 isoform X1
>gi|1036925392|ref|XP_017154892.1| PREDICTED: uncharacterized protein LOC108163886 isoform X1
>gi|1036921194|ref|XP_017154662.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036921212|ref|XP_017154663.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036921232|ref|XP_017154664.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
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>gi|1036920608|ref|XP_017154625.1| PREDICTED: uncharacterized protein LOC108163701 isoform X1
>gi|1036920626|ref|XP_017154626.1| PREDICTED: uncharacterized protein LOC108163701 isoform X2
>gi|1036920644|ref|XP_017154627.1| PREDICTED: uncharacterized protein LOC108163701 isoform X3
>gi|1036930737|ref|XP_017155186.1| PREDICTED: uncharacterized protein LOC108164106 [Drosophila
>gi|1036854322|ref|XP_017150941.1| PREDICTED: uncharacterized protein LOC108161227 [Drosophila
>gi|1036928683|ref|XP_017155076.1| PREDICTED: uncharacterized protein LOC108164012 [Drosophila
>gi|1036928666|ref|XP_017155075.1| PREDICTED: uncharacterized protein LOC108164011 isoform X2
>gi|1036928650|ref|XP_017155074.1| PREDICTED: uncharacterized protein LOC108164011 isoform X1
>gi|1036933841|ref|XP_017155353.1| PREDICTED: uncharacterized protein LOC108164233 [Drosophila
>gi|1036854340|ref|XP_017150942.1| PREDICTED: proteasome subunit beta type-6-like [Drosophila r
>gi|1036933638|ref|XP_017155344.1| PREDICTED: uncharacterized protein LOC108164222 [Drosophila
>gi|1036934946|ref|XP_017155415.1| PREDICTED: adenylate kinase isoenzyme 5 isoform X2 [Drosophi
>gi|1036934930|ref|XP_017155414.1| PREDICTED: adenylate kinase isoenzyme 5 isoform X1 [Drosophi
>gi|1036934964|ref|XP_017155416.1| PREDICTED: adenylate kinase isoenzyme 5 isoform X3 [Drosophi
>gi|1036929438|ref|XP_017155119.1| PREDICTED: protein rolling stone [Drosophila miranda]
>gi|1036929456|ref|XP_017155120.1| PREDICTED: protein rolling stone [Drosophila miranda]
>gi|1036929474|ref|XP_017155121.1| PREDICTED: sex-regulated protein janus-A [Drosophila miranda
>gi|1036911269|ref|XP_017154088.1| PREDICTED: protein rolling stone [Drosophila miranda]
>gi|1036911347|ref|XP_017154091.1| PREDICTED: protein rolling stone isoform X2 [Drosophila mira
>gi|1036911329|ref|XP_017154090.1| PREDICTED: protein rolling stone isoform X1 [Drosophila mira
>gi|1036911287|ref|XP_017154089.1| PREDICTED: protein rolling stone-like [Drosophila miranda]
>gi|1036911250|ref|XP_017154087.1| PREDICTED: COP9 signalosome complex subunit 2 isoform X2 [Dr
>gi|1036911232|ref|XP_017154086.1| PREDICTED: COP9 signalosome complex subunit 2 isoform X1 [Dr
>gi|1036922879|ref|XP_017154765.1| PREDICTED: UPF0585 protein CG18661 isoform X1 [Drosophila mi
>gi|1036922896|ref|XP_017154766.1| PREDICTED: UPF0585 protein CG18661 isoform X2 [Drosophila mi
>gi|1036922916|ref|XP_017154767.1| PREDICTED: UPF0585 protein CG18661 isoform X3 [Drosophila mi
>gi|1036922841|ref|XP_017154763.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036922859|ref|XP_017154764.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036922934|ref|XP_017154768.1| PREDICTED: uncharacterized protein LOC108163792 [Drosophila
>gi|1036922952|ref|XP_017154769.1| PREDICTED: uncharacterized protein LOC108163793 [Drosophila
>gi|1036854358|ref|XP_017150943.1| PREDICTED: mitochondrial 2-oxodicarboxylate carrier [Drosoph
>gi|1036899793|ref|XP_017153463.1| PREDICTED: lachesin [Drosophila miranda]
>gi|1036858959|ref|XP_017151194.1| PREDICTED: uncharacterized protein LOC108161443 [Drosophila
>gi|1036864541|ref|XP_017151496.1| PREDICTED: odorant receptor 30a [Drosophila miranda]
>gi|1036862814|ref|XP_017151408.1| PREDICTED: uncharacterized protein LOC108161622 isoform X1
>gi|1036862833|ref|XP_017151409.1| PREDICTED: uncharacterized protein LOC108161622 isoform X2
>gi|1036902911|ref|XP_017153637.1| PREDICTED: uncharacterized protein LOC108163073 isoform X1
>gi|1036902975|ref|XP_017153640.1| PREDICTED: uncharacterized protein LOC108163073 isoform X3
>gi|1036902933|ref|XP_017153638.1| PREDICTED: uncharacterized protein LOC108163073 isoform X2
>gi|1036902955|ref|XP_017153639.1| PREDICTED: uncharacterized protein LOC108163073 isoform X2
>gi|1036902994|ref|XP_017153641.1| PREDICTED: uncharacterized protein LOC108163073 isoform X4
>gi|1036903012|ref|XP_017153642.1| PREDICTED: uncharacterized protein LOC108163073 isoform X5
>gi|1036903030|ref|XP_017153643.1| PREDICTED: glutamic acid-rich protein [Drosophila miranda]
>gi|1036903047|ref|XP_017153644.1| PREDICTED: coiled-coil domain-containing protein 43 [Drosoph
>gi|1036888555|ref|XP_017152826.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036888574|ref|XP_017152827.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036888619|ref|XP_017152829.1| PREDICTED: nuclear receptor coactivator 3 isoform X3 [Drosop
>gi|1036888600|ref|XP_017152828.1| PREDICTED: nuclear receptor coactivator 3 isoform X2 [Drosop
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>gi|1036854377|ref|XP_017150944.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036907284|ref|XP_017153884.1| PREDICTED: borealin isoform X2 [Drosophila miranda]
>gi|1036907248|ref|XP_017153882.1| PREDICTED: borealin isoform X1 [Drosophila miranda]
>gi|1036907266|ref|XP_017153883.1| PREDICTED: borealin isoform X1 [Drosophila miranda]
>gi|1036907303|ref|XP_017153885.1| PREDICTED: guanine nucleotide-binding protein subunit gamma-
>gi|1036907321|ref|XP_017153886.1| PREDICTED: guanine nucleotide-binding protein subunit gamma-
>gi|1036932741|ref|XP_017155293.1| PREDICTED: uncharacterized protein LOC108164186 [Drosophila
>gi|1036925356|ref|XP_017154890.1| PREDICTED: uncharacterized protein LOC108163885 isoform X2
>gi|1036925321|ref|XP_017154889.1| PREDICTED: uncharacterized protein LOC108163885 isoform X1
>gi|1036923042|ref|XP_017154775.1| PREDICTED: zinc finger protein weckle [Drosophila miranda]
>gi|1036876414|ref|XP_017152168.1| PREDICTED: tyrosine-protein phosphatase Lar isoform X2 [Dros
>gi|1036876454|ref|XP_017152170.1| PREDICTED: tyrosine-protein phosphatase Lar isoform X4 [Dros
>gi|1036876395|ref|XP_017152167.1| PREDICTED: tyrosine-protein phosphatase Lar isoform X1 [Dros
>gi|1036876435|ref|XP_017152169.1| PREDICTED: tyrosine-protein phosphatase Lar isoform X3 [Dros
>gi|1036876492|ref|XP_017152172.1| PREDICTED: uncharacterized protein LOC108162111 [Drosophila
>gi|1036876473|ref|XP_017152171.1| PREDICTED: protein screw [Drosophila miranda]
>gi|1036856676|ref|XP_017151080.1| PREDICTED: uncharacterized protein LOC108161349 isoform X4
>gi|1036856585|ref|XP_017151076.1| PREDICTED: mucin-17 isoform X1 [Drosophila miranda]
>gi|1036856602|ref|XP_017151077.1| PREDICTED: mucin-17 isoform X1 [Drosophila miranda]
>gi|1036856621|ref|XP_017151078.1| PREDICTED: mucin-17 isoform X2 [Drosophila miranda]
>gi|1036856658|ref|XP_017151079.1| PREDICTED: uncharacterized protein LOC108161349 isoform X3
>gi|1036856987|ref|XP_017151100.1| PREDICTED: uncharacterized protein LOC108161367 [Drosophila
>gi|1036856877|ref|XP_017151092.1| PREDICTED: cyclin-K [Drosophila miranda]
>gi|1036856710|ref|XP_017151083.1| PREDICTED: uncharacterized protein LOC108161352 [Drosophila
>gi|1036857096|ref|XP_017151107.1| PREDICTED: uncharacterized protein LOC108161371 [Drosophila
>gi|1036856783|ref|XP_017151087.1| PREDICTED: protein real-time [Drosophila miranda]
>gi|1036856747|ref|XP_017151085.1| PREDICTED: peroxisomal acyl-coenzyme A oxidase 3 isoform X2
>gi|1036856728|ref|XP_017151084.1| PREDICTED: peroxisomal acyl-coenzyme A oxidase 3 isoform X1
>gi|1036856823|ref|XP_017151089.1| PREDICTED: adenosine monophosphate-protein transferase FICD
>gi|1036856568|ref|XP_017151075.1| PREDICTED: tiggrin [Drosophila miranda]
>gi|1036856949|ref|XP_017151098.1| PREDICTED: uncharacterized protein LOC108161366 isoform X1
>gi|1036856968|ref|XP_017151099.1| PREDICTED: uncharacterized protein LOC108161366 isoform X2
>gi|1036857006|ref|XP_017151101.1| PREDICTED: GTP-binding nuclear protein Ran-like isoform X1
>gi|1036857024|ref|XP_017151102.1| PREDICTED: GTP-binding nuclear protein Ran-like isoform X2
>gi|1036857041|ref|XP_017151103.1| PREDICTED: GTP-binding nuclear protein Ran-like isoform X3
>gi|1036856841|ref|XP_017151090.1| PREDICTED: neurotrimin [Drosophila miranda]
>gi|1036857058|ref|XP_017151104.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036857077|ref|XP_017151105.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036856931|ref|XP_017151096.1| PREDICTED: uncharacterized protein LOC108161364 [Drosophila
>gi|1036856764|ref|XP_017151086.1| PREDICTED: membrane metallo-endopeptidase-like 1 [Drosophila
>gi|1036856695|ref|XP_017151081.1| PREDICTED: uncharacterized protein LOC108161351 [Drosophila
>gi|1036856802|ref|XP_017151088.1| PREDICTED: endothelin-converting enzyme 1 [Drosophila miranc
>gi|1036854396|ref|XP_017150945.1| PREDICTED: techylectin-5B [Drosophila miranda]
>gi|1036854415|ref|XP_017150946.1| PREDICTED: pickpocket protein 19 [Drosophila miranda]
>gi|1036854434|ref|XP_017150948.1| PREDICTED: pickpocket protein 19 [Drosophila miranda]
>gi|1036856860|ref|XP_017151091.1| PREDICTED: uncharacterized protein LOC108161360 [Drosophila
>gi|1036856913|ref|XP_017151095.1| PREDICTED: uncharacterized protein LOC108161363 [Drosophila
>gi|1036856896|ref|XP_017151093.1| PREDICTED: uncharacterized protein LOC108161362 [Drosophila
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>gi|1036902695|ref|XP_017153626.1| PREDICTED: NADPH--cytochrome P450 reductase isoform X1 [Dros
>gi|1036902713|ref|XP_017153627.1| PREDICTED: NADPH--cytochrome P450 reductase isoform X2 [Dros
>gi|1036902731|ref|XP_017153628.1| PREDICTED: zinc finger protein 180-like [Drosophila miranda]
>gi|1036868326|ref|XP_017151712.1| PREDICTED: gastrula zinc finger protein XlCGF57.1-like [Dros
>gi|1036868344|ref|XP_017151713.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036868308|ref|XP_017151711.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 2
>gi|1036868289|ref|XP_017151709.1| PREDICTED: rap guanine nucleotide exchange factor 2 isoform
>gi|1036868270|ref|XP_017151708.1| PREDICTED: rap guanine nucleotide exchange factor 2 isoform
>gi|1036854453|ref|XP_017150949.1| PREDICTED: RAB6A-GEF complex partner protein 2-like [Drosoph
>gi|1036924700|ref|XP_017154859.1| PREDICTED: programmed cell death 6-interacting protein-like
>gi|1036854472|ref|XP_017150950.1| PREDICTED: limonin dehydrogenase-like [Drosophila miranda]
>gi|1036861298|ref|XP_017151325.1| PREDICTED: uncharacterized protein LOC108161553 [Drosophila
>gi|1036862096|ref|XP_017151369.1| PREDICTED: uncharacterized protein LOC108161594 [Drosophila
>gi|1036876842|ref|XP_017152192.1| PREDICTED: discoidin domain-containing receptor 2-like isofo
>gi|1036876861|ref|XP_017152193.1| PREDICTED: discoidin domain-containing receptor 2-like isofo
>gi|1036876880|ref|XP_017152194.1| PREDICTED: discoidin domain-containing receptor 2-like isofo
>gi|1036877159|ref|XP_017152211.1| PREDICTED: LOW QUALITY PROTEIN: dopamine N-acetyltransferase
>gi|1036877065|ref|XP_017152205.1| PREDICTED: uncharacterized protein CG1339 [Drosophila miran
>gi|1036877248|ref|XP_017152216.1| PREDICTED: uncharacterized protein LOC108162146 [Drosophila
>gi|1036877193|ref|XP_017152213.1| PREDICTED: uncharacterized protein LOC108162142 [Drosophila
>gi|1036876899|ref|XP_017152195.1| PREDICTED: discoidin domain-containing receptor 2-like isofo
>gi|1036877231|ref|XP_017152215.1| PREDICTED: uncharacterized protein LOC108162145 [Drosophila
>gi|1036877103|ref|XP_017152208.1| PREDICTED: uncharacterized protein LOC108162137 [Drosophila
>gi|1036877122|ref|XP_017152209.1| PREDICTED: uncharacterized protein LOC108162138 [Drosophila
>gi|1036877141|ref|XP_017152210.1| PREDICTED: uncharacterized protein LOC108162139 [Drosophila
>gi|1036876973|ref|XP_017152200.1| PREDICTED: UDP-glucuronosyltransferase 1-7C-like [Drosophila
>gi|1036877009|ref|XP_017152202.1| PREDICTED: UDP-glucuronosyltransferase 2B1-like isoform X2
>gi|1036876992|ref|XP_017152201.1| PREDICTED: UDP-glucuronosyltransferase 2B15-like isoform X1
>gi|1036877028|ref|XP_017152203.1| PREDICTED: UDP-glucuronosyltransferase 2B1-like [Drosophila
>gi|1036876937|ref|XP_017152197.1| PREDICTED: uncharacterized protein LOC108162129 [Drosophila
>gi|1036854490|ref|XP_017150951.1| PREDICTED: ficolin-1-like [Drosophila miranda]
>gi|1036854509|ref|XP_017150952.1| PREDICTED: LOW QUALITY PROTEIN: ficolin-1-like [Drosophila
>gi|1036877174|ref|XP_017152212.1| PREDICTED: uncharacterized protein LOC108162141 [Drosophila
>gi|1036877047|ref|XP_017152204.1| PREDICTED: LOW QUALITY PROTEIN: salivary glue protein Sgs-3
>gi|1036876806|ref|XP_017152190.1| PREDICTED: uncharacterized protein LOC108162125 [Drosophila
>gi|1036876825|ref|XP_017152191.1| PREDICTED: uncharacterized protein LOC108162125 [Drosophila
>gi|1036854528|ref|XP_017150953.1| PREDICTED: uncharacterized protein LOC108161239 [Drosophila
>gi|1036877267|ref|XP_017152217.1| PREDICTED: protein kish-A-like [Drosophila miranda]
>gi|1036877084|ref|XP_017152207.1| PREDICTED: testis-specific serine/threonine-protein kinase
>gi|1036877212|ref|XP_017152214.1| PREDICTED: B9 domain-containing protein 2 [Drosophila miran
>gi|1036876956|ref|XP_017152199.1| PREDICTED: tectonic [Drosophila miranda]
>gi|1036876918|ref|XP_017152196.1| PREDICTED: dynein heavy chain 1, axonemal [Drosophila miran
>gi|1036916775|ref|XP_017154410.1| PREDICTED: telomerase Cajal body protein 1 [Drosophila mira
>gi|1036916793|ref|XP_017154412.1| PREDICTED: actin-related protein 2/3 complex subunit 4 [Dros
>gi|1036894545|ref|XP_017153166.1| PREDICTED: stathmin isoform X4 [Drosophila miranda]
>gi|1036894564|ref|XP_017153167.1| PREDICTED: stathmin isoform X5 [Drosophila miranda]
>gi|1036894527|ref|XP_017153165.1| PREDICTED: golgin subfamily A member 4 isoform X3 [Drosophil
>gi|1036894504|ref|XP_017153164.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X2 [Dro
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>gi|1036894485|ref|XP_017153163.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1036894466|ref|XP_017153162.1| PREDICTED: uncharacterized protein LOC108162777 [Drosophila
>gi|1036920316|ref|XP_017154607.1| PREDICTED: prolactin regulatory element-binding protein [Dro
>gi|1036854547|ref|XP_017150954.1| PREDICTED: uncharacterized protein LOC108161240 [Drosophila
>gi|1036854566|ref|XP_017150955.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036919954|ref|XP_017154586.1| PREDICTED: Krueppel homolog 1-like isoform X3 [Drosophila mi
>gi|1036919918|ref|XP_017154583.1| PREDICTED: Krueppel homolog 1-like isoform X1 [Drosophila mi
>gi|1036919936|ref|XP_017154585.1| PREDICTED: Krueppel homolog 1-like isoform X2 [Drosophila mi
>gi|1036899468|ref|XP_017153444.1| PREDICTED: Krueppel homolog 2 [Drosophila miranda]
>gi|1036899508|ref|XP_017153446.1| PREDICTED: LOW QUALITY PROTEIN: protein-lysine N-methyltrans
>gi|1036899545|ref|XP_017153448.1| PREDICTED: prefoldin subunit 1 [Drosophila miranda]
>gi|1036899487|ref|XP_017153445.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036899428|ref|XP_017153442.1| PREDICTED: uncharacterized protein LOC108162955 isoform X1
>gi|1036899449|ref|XP_017153443.1| PREDICTED: uncharacterized protein LOC108162955 isoform X2
>gi|1036899409|ref|XP_017153441.1| PREDICTED: F-box/WD repeat-containing protein 5 [Drosophila
>gi|1036899527|ref|XP_017153447.1| PREDICTED: protein phosphatase 1 regulatory subunit 11 [Dros
>gi|1036922463|ref|XP_017154737.1| PREDICTED: NADH dehydrogenase [ubiquinone] flavoprotein 1, m
>gi|1036922477|ref|XP_017154738.1| PREDICTED: uncharacterized protein LOC108163773 [Drosophila
>gi|1036932936|ref|XP_017155304.1| PREDICTED: kiSS-1 receptor [Drosophila miranda]
>gi|1036932954|ref|XP_017155305.1| PREDICTED: kiSS-1 receptor [Drosophila miranda]
>gi|1036875073|ref|XP_017152095.1| PREDICTED: protein RCC2 homolog [Drosophila miranda]
>gi|1036875092|ref|XP_017152096.1| PREDICTED: uncharacterized protein LOC108162063 [Drosophila
>gi|1036875111|ref|XP_017152097.1| PREDICTED: sesquipedalian-1 [Drosophila miranda]
>gi|1036875174|ref|XP_017152099.1| PREDICTED: uncharacterized protein LOC108162066 [Drosophila
>gi|1036875130|ref|XP_017152098.1| PREDICTED: protein slowmo [Drosophila miranda]
>gi|1036874980|ref|XP_017152089.1| PREDICTED: phosphoribosylformylglycinamidine synthase [Dros
>gi|1036874999|ref|XP_017152090.1| PREDICTED: phosphoribosylformylglycinamidine synthase [Dros
>gi|1036875018|ref|XP_017152091.1| PREDICTED: phosphoribosylformylglycinamidine synthase [Dros
>gi|1036875035|ref|XP_017152093.1| PREDICTED: uncharacterized protein LOC108162061 isoform X1
>gi|1036875054|ref|XP_017152094.1| PREDICTED: uncharacterized protein LOC108162061 isoform X2
>gi|1036914376|ref|XP_017154261.1| PREDICTED: metallo-beta-lactamase domain-containing protein
>gi|1036914307|ref|XP_017154257.1| PREDICTED: myotubularin-related protein 2 [Drosophila miran
>gi|1036914325|ref|XP_017154258.1| PREDICTED: beta-1,3-glucosyltransferase isoform X1 [Drosophi
>gi|1036914339|ref|XP_017154259.1| PREDICTED: beta-1,3-glucosyltransferase isoform X1 [Drosophi
>gi|1036914357|ref|XP_017154260.1| PREDICTED: beta-1,3-glucosyltransferase isoform X2 [Drosophi
>gi|1036928761|ref|XP_017155080.1| PREDICTED: ribosomal RNA-processing protein 7 homolog A [Dro
>gi|1036857615|ref|XP_017151133.1| PREDICTED: uncharacterized protein LOC108161396 [Drosophila
>gi|1036857189|ref|XP_017151112.1| PREDICTED: homeobox protein H2.0 [Drosophila miranda]
>gi|1036929972|ref|XP_017155145.1| PREDICTED: breast cancer anti-estrogen resistance protein 3
>gi|1036929990|ref|XP_017155146.1| PREDICTED: breast cancer anti-estrogen resistance protein 3
>gi|1036915635|ref|XP_017154342.1| PREDICTED: beta-galactosidase [Drosophila miranda]
>gi|1036915713|ref|XP_017154346.1| PREDICTED: tetraspanin-17 isoform X4 [Drosophila miranda]
>gi|1036915655|ref|XP_017154343.1| PREDICTED: tetraspanin-17 isoform X1 [Drosophila miranda]
>gi|1036915673|ref|XP_017154344.1| PREDICTED: tetraspanin-17 isoform X2 [Drosophila miranda]
>gi|1036915731|ref|XP_017154349.1| PREDICTED: tetraspanin-17 isoform X5 [Drosophila miranda]
>gi|1036915693|ref|XP_017154345.1| PREDICTED: tetraspanin-17 isoform X3 [Drosophila miranda]
>gi|1036893368|ref|XP_017153103.1| PREDICTED: lysine-specific demethylase lid [Drosophila miran
>gi|1036893387|ref|XP_017153104.1| PREDICTED: lysine-specific demethylase lid [Drosophila miran
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>gi|1036896366|ref|XP_017153269.1| PREDICTED: sphingolipid delta(4)-desaturase DES1 [Drosophila
>gi|1036896385|ref|XP_017153270.1| PREDICTED: eukaryotic initiation factor 4A isoform X1 [Dros
>gi|1036896404|ref|XP_017153271.1| PREDICTED: eukaryotic initiation factor 4A isoform X2 [Dros
>gi|1036907770|ref|XP_017153907.1| PREDICTED: profilin [Drosophila miranda]
>gi|1036907789|ref|XP_017153908.1| PREDICTED: profilin [Drosophila miranda]
>gi|1036932631|ref|XP_017155286.1| PREDICTED: mitochondrial uncoupling protein 3-like [Drosoph
>gi|1036932587|ref|XP_017155284.1| PREDICTED: mitochondrial uncoupling protein 4-like isoform
>gi|1036932611|ref|XP_017155285.1| PREDICTED: mitochondrial uncoupling protein 4-like isoform
>gi|1036932847|ref|XP_017155299.1| PREDICTED: trithorax group protein osa [Drosophila miranda]
>gi|1036932865|ref|XP_017155300.1| PREDICTED: trithorax group protein osa [Drosophila miranda]
>gi|1036935789|ref|XP_017155465.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036933414|ref|XP_017155333.1| PREDICTED: vitelline membrane protein Vm26Aa [Drosophila mi
>gi|1036935824|ref|XP_017155467.1| PREDICTED: myosin heavy chain IB [Drosophila miranda]
>gi|1036857707|ref|XP_017151139.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mi
>gi|1036904171|ref|XP_017153707.1| PREDICTED: transmembrane protein 138 [Drosophila miranda]
>gi|1036904150|ref|XP_017153706.1| PREDICTED: serine/threonine-protein kinase 11-interacting p
>gi|1036904130|ref|XP_017153705.1| PREDICTED: serine/threonine-protein kinase 11-interacting p
>gi|1036904189|ref|XP_017153708.1| PREDICTED: pulmonary surfactant-associated protein D-like [
>gi|1036921108|ref|XP_017154656.1| PREDICTED: glycerol-3-phosphate dehydrogenase [NAD(+)], cyto
>gi|1036921090|ref|XP_017154655.1| PREDICTED: glycerol-3-phosphate dehydrogenase [NAD(+)], cyto
>gi|1036927293|ref|XP_017155002.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop
>gi|1036859932|ref|XP_017151249.1| PREDICTED: uncharacterized protein LOC108161487 [Drosophila
>gi|1036862132|ref|XP_017151372.1| PREDICTED: mitochondrial import receptor subunit TOM7 homol
>gi|1036854585|ref|XP_017150956.1| PREDICTED: uncharacterized protein LOC108161243 [Drosophila
>gi|1036934405|ref|XP_017155384.1| PREDICTED: macrophage mannose receptor 1-like isoform X1 [D
>gi|1036934422|ref|XP_017155385.1| PREDICTED: uncharacterized protein LOC108164256 isoform X2
>gi|1036857308|ref|XP_017151117.1| PREDICTED: eggshell protein [Drosophila miranda]
>gi|1036857325|ref|XP_017151118.1| PREDICTED: eggshell protein [Drosophila miranda]
>gi|1036893795|ref|XP_017153123.1| PREDICTED: WD repeat and FYVE domain-containing protein 3 [
>gi|1036852628|ref|XP_017150833.1| PREDICTED: uncharacterized protein LOC108161121 [Drosophila
>gi|1036860136|ref|XP_017151260.1| PREDICTED: uncharacterized protein LOC108161496 [Drosophila
>gi|1036860712|ref|XP_017151290.1| PREDICTED: uncharacterized protein LOC108161523 [Drosophila
>gi|1036930454|ref|XP_017155170.1| PREDICTED: uncharacterized protein LOC108164092 [Drosophila
>gi|1036930471|ref|XP_017155171.1| PREDICTED: uncharacterized protein LOC108164093 [Drosophila
>gi|1036859984|ref|XP_017151251.1| PREDICTED: accessory gland-specific peptide 26Ab-like [Dros
>gi|1036854604|ref|XP_017150957.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop
>gi|1036860210|ref|XP_017151264.1| PREDICTED: protein TsetseEP [Drosophila miranda]
>gi|1036859106|ref|XP_017151202.1| PREDICTED: leucine-rich repeat extensin-like protein 1 [Dro
>gi|1036917765|ref|XP_017154466.1| PREDICTED: transmembrane protein 209-like [Drosophila miran
>gi|1036917784|ref|XP_017154467.1| PREDICTED: uncharacterized protein LOC108163595 [Drosophila
>gi|1036926917|ref|XP_017154982.1| PREDICTED: neuropeptide Y receptor type 5 isoform X2 [Droso
>gi|1036926884|ref|XP_017154981.1| PREDICTED: alpha-2C adrenergic receptor isoform X1 [Drosoph
>gi|1036926935|ref|XP_017154983.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosoph
>gi|1036926953|ref|XP_017154984.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosoph
>gi|1036926971|ref|XP_017154985.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosoph
>gi|1036926989|ref|XP_017154986.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosoph
>gi|1036927007|ref|XP_017154987.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosoph
>gi|1036927025|ref|XP_017154988.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosoph
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>gi|1036927042|ref|XP_017154989.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosophi
>gi|1036927058|ref|XP_017154990.1| PREDICTED: alpha-2C adrenergic receptor isoform X3 [Drosophi
>gi|1036927076|ref|XP_017154991.1| PREDICTED: uncharacterized protein LOC108163953 [Drosophila
>gi|1036927094|ref|XP_017154992.1| PREDICTED: uncharacterized protein LOC108163954 [Drosophila
>gi|1036934643|ref|XP_017155400.1| PREDICTED: uncharacterized protein LOC108164268 [Drosophila
>gi|1036860562|ref|XP_017151282.1| PREDICTED: uncharacterized protein LOC108161515 isoform X2
>gi|1036860543|ref|XP_017151281.1| PREDICTED: vasotab-like isoform X1 [Drosophila miranda]
>gi|1036861727|ref|XP_017151351.1| PREDICTED: uncharacterized protein LOC108161574 [Drosophila
>gi|1036934175|ref|XP_017155371.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036907658|ref|XP_017153901.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X2 [Drosophi
>gi|1036907640|ref|XP_017153900.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X1 [Drosophi
>gi|1036907677|ref|XP_017153902.1| PREDICTED: GATA zinc finger domain-containing protein 10 [D
>gi|1036907696|ref|XP_017153903.1| PREDICTED: uncharacterized protein LOC108163248 [Drosophila
>gi|1036906738|ref|XP_017153855.1| PREDICTED: uncharacterized protein LOC108163211 [Drosophila
>gi|1036906625|ref|XP_017153848.1| PREDICTED: beta-1,4-glucuronyltransferase 1 isoform X1 [Dros
>gi|1036906643|ref|XP_017153849.1| PREDICTED: beta-1,4-glucuronyltransferase 1 isoform X2 [Dros
>gi|1036906663|ref|XP_017153850.1| PREDICTED: beta-1,4-glucuronyltransferase 1 isoform X2 [Dros
>gi|1036906681|ref|XP_017153851.1| PREDICTED: beta-1,4-glucuronyltransferase 1 isoform X2 [Dros
>gi|1036906719|ref|XP_017153854.1| PREDICTED: peritrophin-1 isoform X2 [Drosophila miranda]
>gi|1036906699|ref|XP_017153853.1| PREDICTED: uncharacterized protein LOC108163210 isoform X1
>gi|1036910178|ref|XP_017154029.1| PREDICTED: neuroguidin [Drosophila miranda]
>gi|1036910140|ref|XP_017154027.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036910119|ref|XP_017154026.1| PREDICTED: ABC transporter G family member 20 [Drosophila m
>gi|1036910160|ref|XP_017154028.1| PREDICTED: uncharacterized protein LOC108163315 [Drosophila
>gi|1036917637|ref|XP_017154458.1| PREDICTED: septin-interacting protein 1 [Drosophila miranda]
>gi|1036917655|ref|XP_017154460.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917673|ref|XP_017154461.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917691|ref|XP_017154462.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917709|ref|XP_017154463.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036917729|ref|XP_017154464.1| PREDICTED: beta-1,4-glucuronyltransferase 1-like [Drosophila
>gi|1036854623|ref|XP_017150959.1| PREDICTED: uncharacterized protein LOC108161246 [Drosophila
>gi|1036865549|ref|XP_017151553.1| PREDICTED: uncharacterized protein LOC108161742 [Drosophila
>gi|1036905035|ref|XP_017153753.1| PREDICTED: LOW QUALITY PROTEIN: protein msta [Drosophila mi
>gi|1036905073|ref|XP_017153755.1| PREDICTED: LOW QUALITY PROTEIN: palmitoyltransferase ZDHHC3
>gi|1036905053|ref|XP_017153754.1| PREDICTED: aurora kinase B [Drosophila miranda]
>gi|1036906792|ref|XP_017153856.1| PREDICTED: LOW QUALITY PROTEIN: signal transducing adapter
>gi|1036906812|ref|XP_017153857.1| PREDICTED: uncharacterized protein LOC108163214 isoform X1
>gi|1036906830|ref|XP_017153858.1| PREDICTED: uncharacterized protein LOC108163214 isoform X2
>gi|1036906863|ref|XP_017153860.1| PREDICTED: uncharacterized protein LOC108163216 [Drosophila
>gi|1036906848|ref|XP_017153859.1| PREDICTED: uncharacterized protein LOC108163214 isoform X3
>gi|1036860043|ref|XP_017151255.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036858924|ref|XP_017151192.1| PREDICTED: ribose-phosphate pyrophosphokinase 2-like [Drosop
>gi|1036933739|ref|XP_017155348.1| PREDICTED: LOW QUALITY PROTEIN: protein I'm not dead yet [D
>gi|1036933722|ref|XP_017155346.1| PREDICTED: uncharacterized protein LOC108164227 [Drosophila
>gi|1036933278|ref|XP_017155325.1| PREDICTED: uncharacterized protein LOC108164206 isoform X1
>gi|1036933296|ref|XP_017155326.1| PREDICTED: uncharacterized protein LOC108164206 isoform X1
>gi|1036933314|ref|XP_017155327.1| PREDICTED: hemicentin-1 isoform X2 [Drosophila miranda]
>gi|1036933344|ref|XP_017155328.1| PREDICTED: uncharacterized protein LOC108164208 [Drosophila
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>gi|1036911543|ref|XP_017154101.1| PREDICTED: transmembrane protein 214 isoform X1 [Drosophila
>gi|1036911574|ref|XP_017154102.1| PREDICTED: uncharacterized protein LOC108163363 isoform X2
>gi|1036897699|ref|XP_017153344.1| PREDICTED: DET1 homolog [Drosophila miranda]
>gi|1036897717|ref|XP_017153345.1| PREDICTED: LOW QUALITY PROTEIN: ATP synthase subunit g, mit
>gi|1036897680|ref|XP_017153343.1| PREDICTED: wiskott-Aldrich syndrome protein family member 2
>gi|1036897621|ref|XP_017153338.1| PREDICTED: wiskott-Aldrich syndrome protein family member 3
>gi|1036897640|ref|XP_017153339.1| PREDICTED: wiskott-Aldrich syndrome protein family member 3
>gi|1036897659|ref|XP_017153342.1| PREDICTED: wiskott-Aldrich syndrome protein family member 3
>gi|1036913531|ref|XP_017154217.1| PREDICTED: LOW QUALITY PROTEIN: protein piwi [Drosophila mi
>gi|1036930773|ref|XP_017155188.1| PREDICTED: lysosomal alpha-mannosidase isoform X1 [Drosophil
>gi|1036930791|ref|XP_017155189.1| PREDICTED: lysosomal alpha-mannosidase isoform X1 [Drosophil
>gi|1036930813|ref|XP_017155191.1| PREDICTED: lysosomal alpha-mannosidase isoform X1 [Drosophil
>gi|1036930834|ref|XP_017155192.1| PREDICTED: lysosomal alpha-mannosidase isoform X2 [Drosophil
>gi|1036900543|ref|XP_017153506.1| PREDICTED: LOW QUALITY PROTEIN: lysosomal alpha-mannosidase
>gi|1036900562|ref|XP_017153507.1| PREDICTED: LOW QUALITY PROTEIN: RNA pseudouridylate synthase
>gi|1036912010|ref|XP_017154125.1| PREDICTED: RNA pseudouridylate synthase domain-containing pr
>gi|1036893422|ref|XP_017153106.1| PREDICTED: gamma-tubulin complex component 4 homolog isofor
>gi|1036893441|ref|XP_017153107.1| PREDICTED: gamma-tubulin complex component 4 homolog isofor
>gi|1036893460|ref|XP_017153108.1| PREDICTED: gamma-tubulin complex component 4 homolog isofor
>gi|1036893403|ref|XP_017153105.1| PREDICTED: conserved oligomeric Golgi complex subunit 4 [Dr
>gi|1036893479|ref|XP_017153109.1| PREDICTED: 60S ribosomal protein L9 [Drosophila miranda]
>gi|1036900194|ref|XP_017153485.1| PREDICTED: nuclear pore complex protein Nup155 [Drosophila m
>gi|1036900210|ref|XP_017153486.1| PREDICTED: protein arginine N-methyltransferase 6 [Drosophil
>gi|1036889040|ref|XP_017152852.1| PREDICTED: protein aubergine [Drosophila miranda]
>gi|1036888985|ref|XP_017152850.1| PREDICTED: tubulin polyglutamylase TTLL4 isoform X1 [Drosopl
>gi|1036889022|ref|XP_017152851.1| PREDICTED: uncharacterized protein LOC108162564 isoform X2
>gi|1036895114|ref|XP_017153199.1| PREDICTED: putative nuclease HARBI1 [Drosophila miranda]
>gi|1036895076|ref|XP_017153197.1| PREDICTED: uncharacterized protein LOC108162801 isoform X1
>gi|1036895095|ref|XP_017153198.1| PREDICTED: uncharacterized protein LOC108162801 isoform X2
>gi|1036894984|ref|XP_017153191.1| PREDICTED: thrombospondin type-1 domain-containing protein 4
>gi|1036894965|ref|XP_017153190.1| PREDICTED: thrombospondin type-1 domain-containing protein 4
>gi|1036895057|ref|XP_017153195.1| PREDICTED: thrombospondin type-1 domain-containing protein 4
>gi|1036895002|ref|XP_017153192.1| PREDICTED: thrombospondin type-1 domain-containing protein 4
>gi|1036895021|ref|XP_017153193.1| PREDICTED: thrombospondin type-1 domain-containing protein 4
>gi|1036895040|ref|XP_017153194.1| PREDICTED: thrombospondin type-1 domain-containing protein 4
>gi|1036895133|ref|XP_017153200.1| PREDICTED: uncharacterized protein LOC108162803 [Drosophila
>gi|1036900865|ref|XP_017153524.1| PREDICTED: STIP1 homology and U box-containing protein 1 iso
>gi|1036900847|ref|XP_017153523.1| PREDICTED: STIP1 homology and U box-containing protein 1 iso
>gi|1036900885|ref|XP_017153525.1| PREDICTED: STIP1 homology and U box-containing protein 1 iso
>gi|1036900903|ref|XP_017153526.1| PREDICTED: STIP1 homology and U box-containing protein 1 iso
>gi|1036900811|ref|XP_017153520.1| PREDICTED: replication factor C subunit 5 [Drosophila miran
>gi|1036900829|ref|XP_017153522.1| PREDICTED: CD2 antigen cytoplasmic tail-binding protein 2 h
>gi|1036900921|ref|XP_017153527.1| PREDICTED: cAMP-responsive element-binding protein-like 2 is
>gi|1036900939|ref|XP_017153528.1| PREDICTED: cAMP-responsive element-binding protein-like 2 is
>gi|1036900956|ref|XP_017153529.1| PREDICTED: cAMP-responsive element-binding protein-like 2 is
>gi|1036908111|ref|XP_017153924.1| PREDICTED: kinesin-like protein KIF21A isoform X2 [Drosophil
>gi|1036908091|ref|XP_017153923.1| PREDICTED: kinesin-like protein KIF21A isoform X1 [Drosophil
>gi|1036924876|ref|XP_017154869.1| PREDICTED: ATPase family AAA domain-containing protein 1-A
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>gi|1036924503|ref|XP_017154849.1| PREDICTED: LIX1-like protein isoform X1 [Drosophila miranda]
>gi|1036924521|ref|XP_017154850.1| PREDICTED: LIX1-like protein isoform X2 [Drosophila miranda]
>gi|1036921595|ref|XP_017154685.1| PREDICTED: LOW QUALITY PROTEIN: probable trafficking protei
>gi|1036934035|ref|XP_017155366.1| PREDICTED: cathepsin L1 isoform X1 [Drosophila miranda]
>gi|1036934055|ref|XP_017155367.1| PREDICTED: cathepsin L1 isoform X2 [Drosophila miranda]
>gi|1036922358|ref|XP_017154730.1| PREDICTED: protein MEF2BNB homolog [Drosophila miranda]
>gi|1036911724|ref|XP_017154111.1| PREDICTED: LOW QUALITY PROTEIN: probable dolichyl pyrophosph
>gi|1036854640|ref|XP_017150960.1| PREDICTED: importin-4-like [Drosophila miranda]
>gi|1036926869|ref|XP_017154980.1| PREDICTED: ER lumen protein-retaining receptor [Drosophila m
>gi|1036927717|ref|XP_017155029.1| PREDICTED: small nuclear ribonucleoprotein-associated protei
>gi|1036921322|ref|XP_017154669.1| PREDICTED: NAD-dependent protein deacetylase Sirt6 isoform X
>gi|1036921356|ref|XP_017154671.1| PREDICTED: NAD-dependent protein deacetylase Sirt6 isoform X
>gi|1036921339|ref|XP_017154670.1| PREDICTED: NAD-dependent protein deacetylase Sirt6 isoform X
>gi|1036896476|ref|XP_017153274.1| PREDICTED: LOW QUALITY PROTEIN: transcription initiation fac
>gi|1036896457|ref|XP_017153273.1| PREDICTED: LOW QUALITY PROTEIN: TBC1 domain family member 16
>gi|1036896494|ref|XP_017153275.1| PREDICTED: COMM domain-containing protein 5 [Drosophila mira
>gi|1036896513|ref|XP_017153276.1| PREDICTED: COMM domain-containing protein 5 [Drosophila mira
>gi|1036906062|ref|XP_017153817.1| PREDICTED: LOW QUALITY PROTEIN: lumican [Drosophila miranda]
>gi|1036906098|ref|XP_017153819.1| PREDICTED: methionine aminopeptidase 1D, mitochondrial [Dros
>gi|1036906116|ref|XP_017153821.1| PREDICTED: cilia- and flagella-associated protein 20 [Drosop
>gi|1036906080|ref|XP_017153818.1| PREDICTED: protoheme IX farnesyltransferase, mitochondrial
>gi|1036926084|ref|XP_017154936.1| PREDICTED: myotrophin [Drosophila miranda]
>gi|1036902515|ref|XP_017153615.1| PREDICTED: LOW QUALITY PROTEIN: ATP-dependent Clp protease p
>gi|1036902494|ref|XP_017153614.1| PREDICTED: cullin-associated NEDD8-dissociated protein 1 [Dr
>gi|1036916723|ref|XP_017154407.1| PREDICTED: uncharacterized protein LOC108163555 isoform X1
>gi|1036916758|ref|XP_017154409.1| PREDICTED: uncharacterized protein LOC108163555 isoform X3
>gi|1036916741|ref|XP_017154408.1| PREDICTED: uncharacterized protein LOC108163555 isoform X2
>gi|1036916705|ref|XP_017154406.1| PREDICTED: LOW QUALITY PROTEIN: MOB kinase activator-like 3
>gi|1036904803|ref|XP_017153743.1| PREDICTED: cyclin-dependent kinase 1 isoform X1 [Drosophila
>gi|1036904821|ref|XP_017153744.1| PREDICTED: cyclin-dependent kinase 1 isoform X2 [Drosophila
>gi|1036904840|ref|XP_017153745.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036887608|ref|XP_017152775.1| PREDICTED: mitochondrial chaperone BCS1 [Drosophila miranda]
>gi|1036887627|ref|XP_017152776.1| PREDICTED: mitochondrial chaperone BCS1 [Drosophila miranda]
>gi|1036887643|ref|XP_017152777.1| PREDICTED: mitochondrial chaperone BCS1 [Drosophila miranda]
>gi|1036887757|ref|XP_017152784.1| PREDICTED: proteasome subunit alpha type-1 [Drosophila miran
>gi|1036887702|ref|XP_017152780.1| PREDICTED: LIM/homeobox protein Lhx4 isoform X1 [Drosophila
>gi|1036887721|ref|XP_017152782.1| PREDICTED: LIM/homeobox protein Lhx4 isoform X2 [Drosophila
>gi|1036887740|ref|XP_017152783.1| PREDICTED: protein apterous isoform X3 [Drosophila miranda]
>gi|1036887664|ref|XP_017152778.1| PREDICTED: uncharacterized protein LOC108162512 [Drosophila
>gi|1036887683|ref|XP_017152779.1| PREDICTED: uncharacterized protein LOC108162512 [Drosophila
>gi|1036854659|ref|XP_017150961.1| PREDICTED: nipped-B protein [Drosophila miranda]
>gi|1036904486|ref|XP_017153727.1| PREDICTED: putative ATP-dependent RNA helicase DHX33 isoform
>gi|1036904507|ref|XP_017153728.1| PREDICTED: putative ATP-dependent RNA helicase DHX33 isoform
>gi|1036904525|ref|XP_017153729.1| PREDICTED: sodium-dependent phosphate transport protein 3 is
>gi|1036904581|ref|XP_017153733.1| PREDICTED: uncharacterized protein LOC108163129 isoform X4
>gi|1036904563|ref|XP_017153731.1| PREDICTED: vesicular glutamate transporter 1 isoform X3 [Dro
>gi|1036904544|ref|XP_017153730.1| PREDICTED: vesicular glutamate transporter 1 isoform X2 [Dro
>gi|1036913688|ref|XP_017154228.1| PREDICTED: LOW QUALITY PROTEIN: ribonucleoside-diphosphate r
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>gi|1036930663|ref|XP_017155182.1| PREDICTED: LOW QUALITY PROTEIN: inactive rhomboid protein 1
>gi|1036931546|ref|XP_017155229.1| PREDICTED: uncharacterized protein LOC108164138 [Drosophila
>gi|1036857433|ref|XP_017151124.1| PREDICTED: uncharacterized protein LOC108161386 [Drosophila
>gi|1036856497|ref|XP_017151070.1| PREDICTED: uncharacterized protein LOC108161345 [Drosophila
>gi|1036854678|ref|XP_017150962.1| PREDICTED: uncharacterized protein LOC108161250 [Drosophila
>gi|1036921578|ref|XP_017154684.1| PREDICTED: WD repeat and FYVE domain-containing protein 2 [
>gi|1036929224|ref|XP_017155105.1| PREDICTED: sm-like protein LSM4 [Drosophila miranda]
>gi|1036935635|ref|XP_017155457.1| PREDICTED: retinal homeobox protein Rx-B isoform X2 [Drosoph
>gi|1036935617|ref|XP_017155456.1| PREDICTED: short stature homeobox protein 2 isoform X1 [Dros
>gi|1036935686|ref|XP_017155461.1| PREDICTED: paired mesoderm homeobox protein 1 isoform X5 [Di
>gi|1036935671|ref|XP_017155460.1| PREDICTED: aristaless homeobox protein isoform X4 [Drosophil
>gi|1036935653|ref|XP_017155459.1| PREDICTED: paired mesoderm homeobox protein 1 isoform X3 [Di
>gi|1036932545|ref|XP_017155282.1| PREDICTED: alpha-N-acetylgalactosaminidase [Drosophila miran
>gi|1036932565|ref|XP_017155283.1| PREDICTED: alpha-N-acetylgalactosaminidase [Drosophila miran
>gi|1036856390|ref|XP_017151066.1| PREDICTED: uncharacterized protein LOC108161338 [Drosophila
>gi|1036856253|ref|XP_017151058.1| PREDICTED: repressor of RNA polymerase III transcription MAP
>gi|1036855842|ref|XP_017151033.1| PREDICTED: LOW QUALITY PROTEIN: dipeptidyl peptidase 4 [Dros
>gi|1036855695|ref|XP_017151025.1| PREDICTED: histone demethylase UTY isoform X1 [Drosophila mi
>gi|1036855715|ref|XP_017151026.1| PREDICTED: histone demethylase UTY isoform X2 [Drosophila mi
>gi|1036855804|ref|XP_017151031.1| PREDICTED: lysine-specific demethylase 6A isoform X5 [Drosop
>gi|1036855823|ref|XP_017151032.1| PREDICTED: lysine-specific demethylase 6A isoform X6 [Drosop
>gi|1036855731|ref|XP_017151027.1| PREDICTED: histone demethylase UTY isoform X3 [Drosophila mi
>gi|1036855750|ref|XP_017151028.1| PREDICTED: histone demethylase UTY isoform X3 [Drosophila mi
>gi|1036855769|ref|XP_017151029.1| PREDICTED: lysine-specific demethylase 6A isoform X4 [Drosop
>gi|1036855785|ref|XP_017151030.1| PREDICTED: lysine-specific demethylase 6A isoform X4 [Drosop
>gi|1036855937|ref|XP_017151038.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase CYLD [Drosc
>gi|1036855956|ref|XP_017151040.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase CYLD [Drosc
>gi|1036855975|ref|XP_017151041.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase CYLD [Drosc
>gi|1036856161|ref|XP_017151052.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036856013|ref|XP_017151043.1| PREDICTED: LOW QUALITY PROTEIN: ubiquitin carboxyl-terminal
>gi|1036856310|ref|XP_017151061.1| PREDICTED: phospholipid phosphatase 6 [Drosophila miranda]
>gi|1036856180|ref|XP_017151053.1| PREDICTED: uncharacterized protein LOC108161329 isoform X1
>gi|1036856199|ref|XP_017151054.1| PREDICTED: uncharacterized protein LOC108161329 isoform X2
>gi|1036856217|ref|XP_017151055.1| PREDICTED: uncharacterized protein LOC108161329 isoform X3
>gi|1036856143|ref|XP_017151051.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036855676|ref|XP_017151024.1| PREDICTED: LOW QUALITY PROTEIN: E3 ubiquitin-protein ligase
>gi|1036856272|ref|XP_017151059.1| PREDICTED: 28S ribosomal protein S7, mitochondrial [Drosoph:
>gi|1036855861|ref|XP_017151034.1| PREDICTED: TNF receptor-associated factor family protein DDI
>gi|1036855880|ref|XP_017151035.1| PREDICTED: TNF receptor-associated factor family protein DDI
>gi|1036855899|ref|XP_017151036.1| PREDICTED: hybrid signal transduction histidine kinase A iso
>gi|1036855916|ref|XP_017151037.1| PREDICTED: uncharacterized protein LOC108161319 isoform X3
>gi|1036855994|ref|XP_017151042.1| PREDICTED: nicalin [Drosophila miranda]
>gi|1036856329|ref|XP_017151062.1| PREDICTED: RNA-binding protein Rsf1 [Drosophila miranda]
>gi|1036856236|ref|XP_017151056.1| PREDICTED: LOW QUALITY PROTEIN: protein trunk [Drosophila mi
>gi|1036856032|ref|XP_017151044.1| PREDICTED: acylglycerol kinase, mitochondrial [Drosophila mi
>gi|1036856069|ref|XP_017151046.1| PREDICTED: stress-activated protein kinase JNK isoform X2 [I
>gi|1036856050|ref|XP_017151045.1| PREDICTED: stress-activated protein kinase JNK isoform X1 [I
>gi|1036856291|ref|XP_017151060.1| PREDICTED: uncharacterized protein LOC108161333 [Drosophila
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>gi|1036856371|ref|XP_017151064.1| PREDICTED: heart- and neural crest derivatives-expressed pro
>gi|1036856353|ref|XP_017151063.1| PREDICTED: heart- and neural crest derivatives-expressed pro
>gi|1036856088|ref|XP_017151047.1| PREDICTED: S-adenosylmethionine decarboxylase proenzyme iso
>gi|1036856105|ref|XP_017151049.1| PREDICTED: S-adenosylmethionine decarboxylase proenzyme iso
>gi|1036856124|ref|XP_017151050.1| PREDICTED: S-adenosylmethionine decarboxylase proenzyme iso
>gi|1036886555|ref|XP_017152714.1| PREDICTED: protein daughterless [Drosophila miranda]
>gi|1036886573|ref|XP_017152715.1| PREDICTED: uncharacterized protein LOC108162478 isoform X1
>gi|1036886592|ref|XP_017152716.1| PREDICTED: cyclin-dependent kinase 5 activator 1 isoform X2
>gi|1036886611|ref|XP_017152717.1| PREDICTED: cyclin-dependent kinase 5 activator 1 isoform X3
>gi|1036886627|ref|XP_017152718.1| PREDICTED: cyclin-dependent kinase 5 activator 1 isoform X3
>gi|1036886646|ref|XP_017152719.1| PREDICTED: putative ATP-dependent RNA helicase me31b [Drosop
>gi|1036878947|ref|XP_017152301.1| PREDICTED: LOW QUALITY PROTEIN: insulin receptor substrate
>gi|1036854697|ref|XP_017150963.1| PREDICTED: putative serine/threonine-protein kinase STE20-l
>gi|1036878985|ref|XP_017152303.1| PREDICTED: schwannomin-interacting protein 1 homolog isofor
>gi|1036879004|ref|XP_017152304.1| PREDICTED: schwannomin-interacting protein 1 homolog isofor
>gi|1036878966|ref|XP_017152302.1| PREDICTED: LOW QUALITY PROTEIN: tyrosine-protein kinase tra
>gi|1036879023|ref|XP_017152305.1| PREDICTED: probable elongation factor 1-delta [Drosophila m
>gi|1036862408|ref|XP_017151385.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862429|ref|XP_017151387.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862372|ref|XP_017151383.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862391|ref|XP_017151384.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862297|ref|XP_017151380.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862334|ref|XP_017151381.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862353|ref|XP_017151382.1| PREDICTED: phosphatidylinositol 3,4,5-trisphosphate 3-phosp
>gi|1036862606|ref|XP_017151399.1| PREDICTED: protein CEPU-1 [Drosophila miranda]
>gi|1036862519|ref|XP_017151393.1| PREDICTED: FERM domain-containing protein 5 isoform X1 [Dros
>gi|1036862548|ref|XP_017151395.1| PREDICTED: FERM domain-containing protein 5 isoform X3 [Dros
>gi|1036862535|ref|XP_017151394.1| PREDICTED: FERM domain-containing protein 3 isoform X2 [Dros
>gi|1036862447|ref|XP_017151388.1| PREDICTED: uncharacterized protein LOC108161605 isoform X1
>gi|1036862465|ref|XP_017151389.1| PREDICTED: uncharacterized protein LOC108161605 isoform X2
>gi|1036862660|ref|XP_017151402.1| PREDICTED: FUN14 domain-containing protein 2 [Drosophila mi
>gi|1036862623|ref|XP_017151400.1| PREDICTED: probable transcriptional regulatory protein Dtur
>gi|1036862567|ref|XP_017151397.1| PREDICTED: LOW QUALITY PROTEIN: dolichyl-diphosphooligosacc
>gi|1036862586|ref|XP_017151398.1| PREDICTED: tryptase-2 [Drosophila miranda]
>gi|1036862681|ref|XP_017151404.1| PREDICTED: ubiquitin-40S ribosomal protein S27a [Drosophila
>gi|1036862642|ref|XP_017151401.1| PREDICTED: LOW QUALITY PROTEIN: ribosome biogenesis protein
>gi|1036862501|ref|XP_017151392.1| PREDICTED: prolyl endopeptidase [Drosophila miranda]
>gi|1036862482|ref|XP_017151390.1| PREDICTED: coiled-coil domain-containing protein 40 [Drosop
>gi|1036874437|ref|XP_017152058.1| PREDICTED: uncharacterized protein LOC108162043 isoform X1
>gi|1036874455|ref|XP_017152059.1| PREDICTED: uncharacterized protein LOC108162043 isoform X2
>gi|1036874491|ref|XP_017152060.1| PREDICTED: transcription factor MafB [Drosophila miranda]
>gi|1036874399|ref|XP_017152055.1| PREDICTED: 60S ribosomal protein L7 [Drosophila miranda]
>gi|1036874545|ref|XP_017152063.1| PREDICTED: RIP-like protein [Drosophila miranda]
>gi|1036854716|ref|XP_017150964.1| PREDICTED: LOW QUALITY PROTEIN: ATP-binding cassette sub-fa
>gi|1036874294|ref|XP_017152050.1| PREDICTED: LOW QUALITY PROTEIN: fumarate hydratase, mitochor
>gi|1036874526|ref|XP_017152062.1| PREDICTED: LOW QUALITY PROTEIN: cGMP-dependent protein kinas
>gi|1036874507|ref|XP_017152061.1| PREDICTED: uncharacterized protein LOC108162045 [Drosophila
>gi|1036874258|ref|XP_017152048.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
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>gi|1036874418|ref|XP_017152056.1| PREDICTED: heat shock protein 23 [Drosophila miranda]
>gi|1036874313|ref|XP_017152051.1| PREDICTED: leukocyte elastase inhibitor isoform X1 [Drosoph
>gi|1036874349|ref|XP_017152052.1| PREDICTED: leukocyte elastase inhibitor isoform X2 [Drosoph
>gi|1036874368|ref|XP_017152053.1| PREDICTED: leukocyte elastase inhibitor isoform X3 [Drosoph
>gi|1036874383|ref|XP_017152054.1| PREDICTED: leukocyte elastase inhibitor isoform X3 [Drosoph
>gi|1036874276|ref|XP_017152049.1| PREDICTED: importin subunit alpha [Drosophila miranda]
>gi|1036857854|ref|XP_017151147.1| PREDICTED: uncharacterized protein LOC108161407 [Drosophila
>gi|1036933775|ref|XP_017155350.1| PREDICTED: uncharacterized protein LOC108164230 [Drosophila
>gi|1036914250|ref|XP_017154253.1| PREDICTED: uncharacterized protein LOC108163467 [Drosophila
>gi|1036914232|ref|XP_017154252.1| PREDICTED: uncharacterized protein LOC108163466 isoform X3
>gi|1036914194|ref|XP_017154250.1| PREDICTED: uncharacterized protein LOC108163466 isoform X1
>gi|1036914214|ref|XP_017154251.1| PREDICTED: uncharacterized protein LOC108163466 isoform X2
>gi|1036914286|ref|XP_017154256.1| PREDICTED: translocation protein SEC62 isoform X2 [Drosophil
>gi|1036914268|ref|XP_017154254.1| PREDICTED: translocation protein SEC62 isoform X1 [Drosophil
>gi|1036912137|ref|XP_017154133.1| PREDICTED: putative oxidoreductase GLYR1 homolog isoform X2
>gi|1036912117|ref|XP_017154132.1| PREDICTED: putative oxidoreductase GLYR1 homolog isoform X1
>gi|1036902304|ref|XP_017153604.1| PREDICTED: putative ATP-dependent RNA helicase SoYb [Drosoph
>gi|1036902326|ref|XP_017153605.1| PREDICTED: neurogenic protein big brain isoform X1 [Drosoph
>gi|1036902361|ref|XP_017153607.1| PREDICTED: neurogenic protein big brain isoform X3 [Drosoph
>gi|1036902343|ref|XP_017153606.1| PREDICTED: neurogenic protein big brain isoform X2 [Drosoph
>gi|1036905873|ref|XP_017153805.1| PREDICTED: 60S ribosomal protein L13 [Drosophila miranda]
>gi|1036905859|ref|XP_017153804.1| PREDICTED: uncharacterized protein LOC108163178 [Drosophila
>gi|1036889710|ref|XP_017152894.1| PREDICTED: transmembrane protein 184C isoform X1 [Drosophila
>gi|1036889727|ref|XP_017152895.1| PREDICTED: transmembrane protein 184C isoform X2 [Drosophila
>gi|1036889746|ref|XP_017152896.1| PREDICTED: transmembrane protein 184C isoform X3 [Drosophila
>gi|1036889764|ref|XP_017152897.1| PREDICTED: uncharacterized protein LOC108162591 [Drosophila
>gi|1036889797|ref|XP_017152899.1| PREDICTED: DNA-binding protein RFXANK [Drosophila miranda]
>gi|1036889780|ref|XP_017152898.1| PREDICTED: zinc finger CCCH-type with G patch domain-contain
>gi|1036889981|ref|XP_017152910.1| PREDICTED: methyltransferase-like protein 17, mitochondrial
>gi|1036889925|ref|XP_017152907.1| PREDICTED: ATP-binding cassette sub-family G member 4 [Dros
>gi|1036889943|ref|XP_017152908.1| PREDICTED: ATP-binding cassette sub-family G member 4 [Dros
>gi|1036889961|ref|XP_017152909.1| PREDICTED: uncharacterized protein LOC108162601 [Drosophila
>gi|1036890018|ref|XP_017152912.1| PREDICTED: bis(5'-nucleosyl)-tetraphosphatase [asymmetrical]
>gi|1036890037|ref|XP_017152913.1| PREDICTED: bis(5'-nucleosyl)-tetraphosphatase [asymmetrical]
>gi|1036889999|ref|XP_017152911.1| PREDICTED: UPF0430 protein CG31712 [Drosophila miranda]
>gi|1036857671|ref|XP_017151137.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1036874944|ref|XP_017152087.1| PREDICTED: pupal cuticle protein Edg-84A [Drosophila miranda
>gi|1036874687|ref|XP_017152071.1| PREDICTED: cyclin-dependent kinase F-4 isoform X2 [Drosophil
>gi|1036874608|ref|XP_017152067.1| PREDICTED: cyclin-dependent kinase F-4 isoform X1 [Drosophil
>gi|1036874627|ref|XP_017152068.1| PREDICTED: cyclin-dependent kinase F-4 isoform X1 [Drosophil
>gi|1036874646|ref|XP_017152069.1| PREDICTED: cyclin-dependent kinase F-4 isoform X1 [Drosophil
>gi|1036874668|ref|XP_017152070.1| PREDICTED: cyclin-dependent kinase F-4 isoform X1 [Drosophil
>gi|1036874706|ref|XP_017152072.1| PREDICTED: dynein regulatory complex subunit 3 isoform X1 [D
>gi|1036874726|ref|XP_017152073.1| PREDICTED: dynein regulatory complex subunit 3 isoform X2 [D
>gi|1036874835|ref|XP_017152080.1| PREDICTED: enoyl-CoA delta isomerase 1, mitochondrial [Dros
>gi|1036874889|ref|XP_017152083.1| PREDICTED: enoyl-CoA delta isomerase 1, mitochondrial-like
>gi|1036874908|ref|XP_017152085.1| PREDICTED: enoyl-CoA delta isomerase 1, mitochondrial-like
>gi|1036874852|ref|XP_017152081.1| PREDICTED: enoyl-CoA delta isomerase 1, mitochondrial [Dros
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>gi|1036874870|ref|XP_017152082.1| PREDICTED: enoyl-CoA delta isomerase 1, mitochondrial [Dros
>gi|1036874927|ref|XP_017152086.1| PREDICTED: translocon-associated protein subunit gamma [Dros
>gi|1036874816|ref|XP_017152079.1| PREDICTED: 3-ketoacyl-CoA thiolase, mitochondrial [Drosophil
>gi|1036874745|ref|XP_017152074.1| PREDICTED: probable splicing factor, arginine/serine-rich 7
>gi|1036874582|ref|XP_017152064.1| PREDICTED: SWI/SNF-related matrix-associated actin-dependent
>gi|1036874763|ref|XP_017152076.1| PREDICTED: uncharacterized protein LOC108162052 isoform X1
>gi|1036874780|ref|XP_017152077.1| PREDICTED: uncharacterized protein LOC108162052 isoform X1
>gi|1036874798|ref|XP_017152078.1| PREDICTED: uncharacterized protein LOC108162052 isoform X2
>gi|1036922127|ref|XP_017154715.1| PREDICTED: sorting nexin-17 [Drosophila miranda]
>gi|1036899655|ref|XP_017153454.1| PREDICTED: LOW QUALITY PROTEIN: 40S ribosomal protein S2 [Di
>gi|1036899678|ref|XP_017153455.1| PREDICTED: rRNA-processing protein FCF1 homolog [Drosophila
>gi|1036899716|ref|XP_017153457.1| PREDICTED: uncharacterized protein LOC108162970 [Drosophila
>gi|1036899617|ref|XP_017153452.1| PREDICTED: twinkle protein, mitochondrial [Drosophila miran
>gi|1036899735|ref|XP_017153458.1| PREDICTED: cysteine-rich PDZ-binding protein [Drosophila mi
>gi|1036899697|ref|XP_017153456.1| PREDICTED: putative RNA polymerase II subunit B1 CTD phospha
>gi|1036899636|ref|XP_017153453.1| PREDICTED: FK506-binding protein 59 [Drosophila miranda]
>gi|1036931060|ref|XP_017155205.1| PREDICTED: uncharacterized protein LOC108164118 [Drosophila
>gi|1036920260|ref|XP_017154604.1| PREDICTED: acetylcholine receptor subunit alpha-type acr-16
>gi|1036920224|ref|XP_017154602.1| PREDICTED: acetylcholine receptor subunit alpha-type acr-16
>gi|1036920206|ref|XP_017154601.1| PREDICTED: acetylcholine receptor subunit alpha-type acr-16
>gi|1036920191|ref|XP_017154600.1| PREDICTED: acetylcholine receptor subunit alpha-type acr-16
>gi|1036920242|ref|XP_017154603.1| PREDICTED: acetylcholine receptor subunit alpha-type acr-16
>gi|1036880656|ref|XP_017152390.1| PREDICTED: zinc finger protein 521 [Drosophila miranda]
>gi|1036880675|ref|XP_017152391.1| PREDICTED: alpha-ketoglutarate-dependent dioxygenase alkB h
>gi|1036880584|ref|XP_017152386.1| PREDICTED: inositol-trisphosphate 3-kinase A isoform X2 [Dr
>gi|1036880565|ref|XP_017152385.1| PREDICTED: inositol-trisphosphate 3-kinase A isoform X1 [Dr
>gi|1036854734|ref|XP_017150965.1| PREDICTED: pickpocket protein 11 [Drosophila miranda]
>gi|1036880529|ref|XP_017152383.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036880751|ref|XP_017152396.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036880639|ref|XP_017152389.1| PREDICTED: zinc carboxypeptidase A 1 [Drosophila miranda]
>gi|1036880622|ref|XP_017152388.1| PREDICTED: zinc carboxypeptidase A 1 [Drosophila miranda]
>gi|1036880603|ref|XP_017152387.1| PREDICTED: methionine aminopeptidase 2 [Drosophila miranda]
>gi|1036880714|ref|XP_017152394.1| PREDICTED: transcription initiation factor TFIID subunit 11
>gi|1036880478|ref|XP_017152379.1| PREDICTED: zinc finger protein 91 isoform X1 [Drosophila mi
>gi|1036880494|ref|XP_017152380.1| PREDICTED: zinc finger protein 91 isoform X1 [Drosophila mi
>gi|1036880511|ref|XP_017152382.1| PREDICTED: zinc finger protein 91 isoform X2 [Drosophila mi
>gi|1036880548|ref|XP_017152384.1| PREDICTED: cytochrome P450 4e3 [Drosophila miranda]
>gi|1036880317|ref|XP_017152369.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880336|ref|XP_017152370.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880393|ref|XP_017152374.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880355|ref|XP_017152372.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880412|ref|XP_017152375.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880428|ref|XP_017152376.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880461|ref|XP_017152378.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880376|ref|XP_017152373.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880445|ref|XP_017152377.1| PREDICTED: sodium/potassium/calcium exchanger Nckx30C isofor
>gi|1036880733|ref|XP_017152395.1| PREDICTED: uncharacterized protein LOC108162258 [Drosophila
>gi|1036880694|ref|XP_017152392.1| PREDICTED: uncharacterized protein LOC108162256 [Drosophila
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>gi|1036930163|ref|XP_017155154.1| PREDICTED: apoptosis regulatory protein Siva-like [Drosophil
>gi|1036900684|ref|XP_017153513.1| PREDICTED: pescadillo homolog isoform X1 [Drosophila miranda
>gi|1036900702|ref|XP_017153514.1| PREDICTED: pescadillo homolog isoform X2 [Drosophila miranda
>gi|1036900719|ref|XP_017153515.1| PREDICTED: protein pelota [Drosophila miranda]
>gi|1036900755|ref|XP_017153517.1| PREDICTED: uncharacterized protein LOC108163006 [Drosophila
>gi|1036900773|ref|XP_017153518.1| PREDICTED: uncharacterized protein LOC108163006 [Drosophila
>gi|1036900791|ref|XP_017153519.1| PREDICTED: uncharacterized protein LOC108163006 [Drosophila
>gi|1036900737|ref|XP_017153516.1| PREDICTED: NHP2-like protein 1 homolog [Drosophila miranda]
>gi|1036903468|ref|XP_017153666.1| PREDICTED: cAMP-dependent protein kinase catalytic subunit
>gi|1036854751|ref|XP_017150967.1| PREDICTED: uncharacterized protein LOC108161256 [Drosophila
>gi|1036866633|ref|XP_017151615.1| PREDICTED: uncharacterized protein LOC108161784 [Drosophila
>gi|1036866904|ref|XP_017151631.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucuro
>gi|1036866921|ref|XP_017151632.1| PREDICTED: thioredoxin-2 [Drosophila miranda]
>gi|1036866777|ref|XP_017151624.1| PREDICTED: transcription factor glial cells missing 2 isofo
>gi|1036866762|ref|XP_017151623.1| PREDICTED: transcription factor glial cells missing 2 isofo
>gi|1036866795|ref|XP_017151625.1| PREDICTED: transcription factor glial cells missing 2 isofo
>gi|1036866812|ref|XP_017151626.1| PREDICTED: venom carboxylesterase-6 [Drosophila miranda]
>gi|1036866831|ref|XP_017151627.1| PREDICTED: esterase E4 [Drosophila miranda]
>gi|1036866850|ref|XP_017151628.1| PREDICTED: transcription factor glial cells missing [Drosop
>gi|1036866743|ref|XP_017151622.1| PREDICTED: trifunctional enzyme subunit alpha, mitochondrial
>gi|1036866868|ref|XP_017151629.1| PREDICTED: uncharacterized protein LOC108161791 [Drosophila
>gi|1036866885|ref|XP_017151630.1| PREDICTED: uncharacterized protein LOC108161791 [Drosophila
>gi|1036866651|ref|XP_017151616.1| PREDICTED: junctophilin-1 isoform X1 [Drosophila miranda]
>gi|1036866669|ref|XP_017151617.1| PREDICTED: junctophilin-1 isoform X1 [Drosophila miranda]
>gi|1036866688|ref|XP_017151618.1| PREDICTED: junctophilin-1 isoform X1 [Drosophila miranda]
>gi|1036866707|ref|XP_017151619.1| PREDICTED: junctophilin-1 isoform X1 [Drosophila miranda]
>gi|1036866725|ref|XP_017151620.1| PREDICTED: uncharacterized protein LOC108161785 isoform X2
>gi|1036929313|ref|XP_017155111.1| PREDICTED: leucine-rich repeat extensin-like protein 1 [Dros
>gi|1036929351|ref|XP_017155113.1| PREDICTED: uncharacterized protein LOC108164046 [Drosophila
>gi|1036929369|ref|XP_017155114.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1036929333|ref|XP_017155112.1| PREDICTED: uncharacterized protein LOC108164045 [Drosophila
>gi|1036913941|ref|XP_017154243.1| PREDICTED: apoptotic chromatin condensation inducer in the n
>gi|1036913959|ref|XP_017154244.1| PREDICTED: apoptotic chromatin condensation inducer in the n
>gi|1036913976|ref|XP_017154245.1| PREDICTED: apoptotic chromatin condensation inducer in the n
>gi|1036893871|ref|XP_017153127.1| PREDICTED: tetratricopeptide repeat protein 19 homolog, mito
>gi|1036893909|ref|XP_017153129.1| PREDICTED: tetratricopeptide repeat protein 19 homolog, mito
>gi|1036893890|ref|XP_017153128.1| PREDICTED: tetratricopeptide repeat protein 19 homolog, mito
>gi|1036893852|ref|XP_017153126.1| PREDICTED: protein catecholamines up [Drosophila miranda]
>gi|1036893814|ref|XP_017153124.1| PREDICTED: E3 ubiquitin-protein ligase MIB2 [Drosophila mira
>gi|1036893833|ref|XP_017153125.1| PREDICTED: E3 ubiquitin-protein ligase MIB2 [Drosophila mira
>gi|1036893927|ref|XP_017153130.1| PREDICTED: uncharacterized protein C1orf50 homolog [Drosophi
>gi|1036918456|ref|XP_017154506.1| PREDICTED: protein hook [Drosophila miranda]
>gi|1036918476|ref|XP_017154507.1| PREDICTED: uncharacterized protein LOC108163620 [Drosophila
>gi|1036854770|ref|XP_017150968.1| PREDICTED: uncharacterized protein LOC108161257 [Drosophila
>gi|1036919186|ref|XP_017154542.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1036919972|ref|XP_017154587.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1036923400|ref|XP_017154797.1| PREDICTED: 40S ribosomal protein S13 [Drosophila miranda]
>gi|1036923381|ref|XP_017154796.1| PREDICTED: COP9 signalosome complex subunit 8 [Drosophila mi
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>gi|1036911632|ref|XP_017154105.1| PREDICTED: serine/threonine-protein phosphatase PP2A 65 kDa
>gi|1036911650|ref|XP_017154106.1| PREDICTED: serine/threonine-protein phosphatase PP2A 65 kDa
>gi|1036911594|ref|XP_017154103.1| PREDICTED: serine/threonine-protein phosphatase PP2A 65 kDa
>gi|1036911612|ref|XP_017154104.1| PREDICTED: serine/threonine-protein phosphatase PP2A 65 kDa
>gi|1036911668|ref|XP_017154107.1| PREDICTED: inactive pancreatic lipase-related protein 1 [Dro
>gi|1036916418|ref|XP_017154390.1| PREDICTED: TLD domain-containing protein 1 [Drosophila miran
>gi|1036916400|ref|XP_017154389.1| PREDICTED: neither inactivation nor afterpotential protein C
>gi|1036861279|ref|XP_017151324.1| PREDICTED: protein Wnt-10b [Drosophila miranda]
>gi|1036895207|ref|XP_017153205.1| PREDICTED: protein Wnt-6 [Drosophila miranda]
>gi|1036895189|ref|XP_017153203.1| PREDICTED: protein wingless [Drosophila miranda]
>gi|1036895152|ref|XP_017153201.1| PREDICTED: protein Wnt-4 isoform X1 [Drosophila miranda]
>gi|1036895170|ref|XP_017153202.1| PREDICTED: protein Wnt-4 isoform X2 [Drosophila miranda]
>gi|1036877907|ref|XP_017152244.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1036877926|ref|XP_017152245.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1036877739|ref|XP_017152235.1| PREDICTED: sodium bicarbonate cotransporter 3 isoform X8 [D
>gi|1036877609|ref|XP_017152227.1| PREDICTED: sodium-driven chloride bicarbonate exchanger iso
>gi|1036877647|ref|XP_017152229.1| PREDICTED: electroneutral sodium bicarbonate exchanger 1 iso
>gi|1036877684|ref|XP_017152231.1| PREDICTED: sodium bicarbonate cotransporter 3 isoform X5 [D
>gi|1036877720|ref|XP_017152234.1| PREDICTED: sodium bicarbonate cotransporter 3 isoform X7 [D
>gi|1036877757|ref|XP_017152236.1| PREDICTED: electroneutral sodium bicarbonate exchanger 1 iso
>gi|1036877776|ref|XP_017152237.1| PREDICTED: sodium bicarbonate cotransporter 3 isoform X10 [
>gi|1036877851|ref|XP_017152241.1| PREDICTED: sodium-driven chloride bicarbonate exchanger iso
>gi|1036877628|ref|XP_017152228.1| PREDICTED: sodium-driven chloride bicarbonate exchanger iso
>gi|1036877701|ref|XP_017152233.1| PREDICTED: sodium bicarbonate cotransporter 3 isoform X6 [D
>gi|1036877813|ref|XP_017152239.1| PREDICTED: electroneutral sodium bicarbonate exchanger 1 iso
>gi|1036877832|ref|XP_017152240.1| PREDICTED: electroneutral sodium bicarbonate exchanger 1 iso
>gi|1036877666|ref|XP_017152230.1| PREDICTED: sodium-driven chloride bicarbonate exchanger iso
>gi|1036877795|ref|XP_017152238.1| PREDICTED: electroneutral sodium bicarbonate exchanger 1 iso
>gi|1036877888|ref|XP_017152243.1| PREDICTED: elongation factor G, mitochondrial [Drosophila m
>gi|1036877869|ref|XP_017152242.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036863748|ref|XP_017151454.1| PREDICTED: uncharacterized protein LOC108161669 [Drosophila
>gi|1036867258|ref|XP_017151653.1| PREDICTED: engulfment and cell motility protein 1 isoform X
>gi|1036867277|ref|XP_017151654.1| PREDICTED: engulfment and cell motility protein 1 isoform X
>gi|1036854789|ref|XP_017150969.1| PREDICTED: probable protein S-acyltransferase 23 [Drosophila
>gi|1036867241|ref|XP_017151652.1| PREDICTED: threonine--tRNA ligase, cytoplasmic isoform X2 [
>gi|1036867222|ref|XP_017151651.1| PREDICTED: threonine--tRNA ligase, cytoplasmic isoform X1 [
>gi|1036867388|ref|XP_017151660.1| PREDICTED: ras-related protein Rab6 [Drosophila miranda]
>gi|1036867350|ref|XP_017151658.1| PREDICTED: cationic trypsin [Drosophila miranda]
>gi|1036867296|ref|XP_017151655.1| PREDICTED: sorting nexin-2 [Drosophila miranda]
>gi|1036867369|ref|XP_017151659.1| PREDICTED: AN1-type zinc finger protein 2A [Drosophila miran
>gi|1036867183|ref|XP_017151648.1| PREDICTED: conserved oligomeric Golgi complex subunit 3 [Dro
>gi|1036867201|ref|XP_017151649.1| PREDICTED: exocyst complex component 2 [Drosophila miranda]
>gi|1036867333|ref|XP_017151657.1| PREDICTED: uncharacterized protein LOC108161808 isoform X2
>gi|1036867315|ref|XP_017151656.1| PREDICTED: uncharacterized protein LOC108161808 isoform X1
>gi|1036907750|ref|XP_017153906.1| PREDICTED: E3 ubiquitin-protein ligase msl-2 isoform X2 [Dro
>gi|1036907714|ref|XP_017153904.1| PREDICTED: E3 ubiquitin-protein ligase msl-2 isoform X1 [Dro
>gi|1036907732|ref|XP_017153905.1| PREDICTED: E3 ubiquitin-protein ligase msl-2 isoform X1 [Dro
>gi|1036892623|ref|XP_017153060.1| PREDICTED: protein FAM151B isoform X1 [Drosophila miranda]
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>gi|1036892638|ref|XP_017153061.1| PREDICTED: protein FAM151B isoform X2 [Drosophila miranda]
>gi|1036892498|ref|XP_017153053.1| PREDICTED: protein croquemort isoform X1 [Drosophila miranda
>gi|1036892517|ref|XP_017153054.1| PREDICTED: protein croquemort isoform X1 [Drosophila miranda
>gi|1036892536|ref|XP_017153055.1| PREDICTED: protein croquemort isoform X2 [Drosophila miranda
>gi|1036892552|ref|XP_017153056.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036892569|ref|XP_017153057.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036892588|ref|XP_017153059.1| PREDICTED: uncharacterized protein LOC108162710 [Drosophila
>gi|1036902247|ref|XP_017153601.1| PREDICTED: zinc finger protein 3 isoform X10 [Drosophila mi
>gi|1036902225|ref|XP_017153599.1| PREDICTED: zinc finger protein 3 isoform X9 [Drosophila mira
>gi|1036902265|ref|XP_017153602.1| PREDICTED: zinc finger protein 3 isoform X11 [Drosophila mi
>gi|1036902205|ref|XP_017153598.1| PREDICTED: zinc finger protein 3 isoform X8 [Drosophila mira
>gi|1036902143|ref|XP_017153595.1| PREDICTED: zinc finger protein 497 isoform X5 [Drosophila m
>gi|1036902105|ref|XP_017153593.1| PREDICTED: zinc finger protein 271 isoform X3 [Drosophila m
>gi|1036902183|ref|XP_017153597.1| PREDICTED: zinc finger protein 497 isoform X7 [Drosophila m
>gi|1036902083|ref|XP_017153592.1| PREDICTED: zinc finger protein 271 isoform X2 [Drosophila m
>gi|1036902125|ref|XP_017153594.1| PREDICTED: zinc finger protein 271 isoform X4 [Drosophila m
>gi|1036902163|ref|XP_017153596.1| PREDICTED: zinc finger protein 497 isoform X6 [Drosophila m
>gi|1036902026|ref|XP_017153589.1| PREDICTED: zinc finger protein 271 isoform X1 [Drosophila m
>gi|1036902046|ref|XP_017153590.1| PREDICTED: zinc finger protein 271 isoform X1 [Drosophila m
>gi|1036902065|ref|XP_017153591.1| PREDICTED: zinc finger protein 271 isoform X1 [Drosophila m
>gi|1036902286|ref|XP_017153603.1| PREDICTED: gastrula zinc finger protein XlCGF57.1 isoform X
>gi|1036925588|ref|XP_017154904.1| PREDICTED: uncharacterized protein LOC108163896 [Drosophila
>gi|1036925610|ref|XP_017154905.1| PREDICTED: uncharacterized protein LOC108163897 [Drosophila
>gi|1036933791|ref|XP_017155351.1| PREDICTED: homeobox protein goosecoid [Drosophila miranda]
>gi|1036894090|ref|XP_017153140.1| PREDICTED: homeobox protein orthopedia [Drosophila miranda]
>gi|1036894128|ref|XP_017153142.1| PREDICTED: uncharacterized protein LOC108162766 isoform X1
>gi|1036894149|ref|XP_017153143.1| PREDICTED: uncharacterized protein LOC108162766 isoform X2
>gi|1036894168|ref|XP_017153144.1| PREDICTED: uncharacterized protein LOC108162768 [Drosophila
>gi|1036894109|ref|XP_017153141.1| PREDICTED: inositol polyphosphate multikinase [Drosophila m
>gi|1036894071|ref|XP_017153139.1| PREDICTED: splicing factor 3B subunit 1 [Drosophila miranda]
>gi|1036916962|ref|XP_017154416.1| PREDICTED: protein Notchless [Drosophila miranda]
>gi|1036916944|ref|XP_017154415.1| PREDICTED: pseudouridine-metabolizing bifunctional protein (
>gi|1036912491|ref|XP_017154156.1| PREDICTED: uncharacterized protein LOC108163403 [Drosophila
>gi|1036912440|ref|XP_017154152.1| PREDICTED: vegetative cell wall protein gp1 [Drosophila mira
>gi|1036912458|ref|XP_017154153.1| PREDICTED: vegetative cell wall protein gp1 [Drosophila mira
>gi|1036912473|ref|XP_017154155.1| PREDICTED: translocator protein [Drosophila miranda]
>gi|1036912424|ref|XP_017154151.1| PREDICTED: intraflagellar transport protein 140 homolog [Dro
>gi|1036927893|ref|XP_017155039.1| PREDICTED: DNA-binding protein D-ETS-6 isoform X1 [Drosophil
>gi|1036927911|ref|XP_017155041.1| PREDICTED: DNA-binding protein D-ETS-6 isoform X2 [Drosophil
>gi|1036927947|ref|XP_017155043.1| PREDICTED: neuronal acetylcholine receptor subunit alpha-2
>gi|1036927929|ref|XP_017155042.1| PREDICTED: neuronal acetylcholine receptor subunit alpha-2
>gi|1036923060|ref|XP_017154776.1| PREDICTED: minor histocompatibility antigen H13 [Drosophila
>gi|1036927592|ref|XP_017155022.1| PREDICTED: SUMO-conjugating enzyme UBC9-B [Drosophila miranc
>gi|1036883252|ref|XP_017152535.1| PREDICTED: zinc finger protein ush isoform X2 [Drosophila m
>gi|1036883233|ref|XP_017152534.1| PREDICTED: zinc finger protein ush isoform X1 [Drosophila m
>gi|1036883271|ref|XP_017152536.1| PREDICTED: zinc finger protein ush isoform X3 [Drosophila m
>gi|1036883327|ref|XP_017152539.1| PREDICTED: protein arginine N-methyltransferase 8-B-like [D:
>gi|1036883308|ref|XP_017152538.1| PREDICTED: Krueppel-like factor 10 [Drosophila miranda]
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>gi|1036883289|ref|XP_017152537.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036875658|ref|XP_017152127.1| PREDICTED: uncharacterized protein LOC108162084 [Drosophila
>gi|1036875713|ref|XP_017152130.1| PREDICTED: uncharacterized protein LOC108162086 isoform X2
>gi|1036875696|ref|XP_017152129.1| PREDICTED: uncharacterized protein LOC108162086 isoform X1
>gi|1036875773|ref|XP_017152133.1| PREDICTED: uncharacterized protein LOC108162086 isoform X5
>gi|1036875750|ref|XP_017152132.1| PREDICTED: uncharacterized protein LOC108162086 isoform X4
>gi|1036875731|ref|XP_017152131.1| PREDICTED: uncharacterized protein LOC108162086 isoform X3
>gi|1036875885|ref|XP_017152140.1| PREDICTED: dnaJ homolog subfamily B member 11 [Drosophila mi
>gi|1036875792|ref|XP_017152135.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036875811|ref|XP_017152136.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036875830|ref|XP_017152137.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036875849|ref|XP_017152138.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036875866|ref|XP_017152139.1| PREDICTED: protein croquemort [Drosophila miranda]
>gi|1036875677|ref|XP_017152128.1| PREDICTED: protein expanded [Drosophila miranda]
>gi|1036867484|ref|XP_017151666.1| PREDICTED: tetratricopeptide repeat protein 8 [Drosophila mi
>gi|1036867522|ref|XP_017151668.1| PREDICTED: ADP-ribosylation factor-like protein 16 [Drosoph
>gi|1036867541|ref|XP_017151669.1| PREDICTED: uncharacterized protein LOC108161822 [Drosophila
>gi|1036867560|ref|XP_017151670.1| PREDICTED: 60S acidic ribosomal protein P1 [Drosophila mira
>gi|1036867503|ref|XP_017151667.1| PREDICTED: ribonuclease H2 subunit A [Drosophila miranda]
>gi|1036867461|ref|XP_017151665.1| PREDICTED: F-box-like/WD repeat-containing protein ebi [Dros
>gi|1036867442|ref|XP_017151664.1| PREDICTED: AP-2 complex subunit alpha [Drosophila miranda]
>gi|1036867425|ref|XP_017151663.1| PREDICTED: DNA-directed RNA polymerase I subunit RPA2 [Droso
>gi|1036867407|ref|XP_017151661.1| PREDICTED: protein lava lamp [Drosophila miranda]
>gi|1036926760|ref|XP_017154973.1| PREDICTED: homeobox protein aristaless isoform X1 [Drosophil
>gi|1036926775|ref|XP_017154974.1| PREDICTED: homeobox protein aristaless isoform X2 [Drosophil
>gi|1036926742|ref|XP_017154972.1| PREDICTED: 60 kDa heat shock protein homolog 1, mitochondria
>gi|1036888875|ref|XP_017152844.1| PREDICTED: equilibrative nucleoside transporter 1 [Drosophil
>gi|1036888841|ref|XP_017152842.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036888807|ref|XP_017152840.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036888824|ref|XP_017152841.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036888779|ref|XP_017152838.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036888792|ref|XP_017152839.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036888893|ref|XP_017152846.1| PREDICTED: uncharacterized protein LOC108162560 [Drosophila
>gi|1036888858|ref|XP_017152843.1| PREDICTED: uncharacterized protein LOC108162558 [Drosophila
>gi|1036888910|ref|XP_017152847.1| PREDICTED: lectizyme [Drosophila miranda]
>gi|1036888929|ref|XP_017152848.1| PREDICTED: lectizyme [Drosophila miranda]
>gi|1036888948|ref|XP_017152849.1| PREDICTED: lectizyme-like [Drosophila miranda]
>gi|1036873244|ref|XP_017151993.1| PREDICTED: rap1 GTPase-activating protein 1 isoform X1 [Dros
>gi|1036873260|ref|XP_017151994.1| PREDICTED: rap1 GTPase-activating protein 1 isoform X2 [Dros
>gi|1036873277|ref|XP_017151995.1| PREDICTED: rap1 GTPase-activating protein 1 isoform X3 [Dros
>gi|1036873317|ref|XP_017151998.1| PREDICTED: lectizyme-like [Drosophila miranda]
>gi|1036873355|ref|XP_017152000.1| PREDICTED: uncharacterized protein LOC108162017 isoform X2
>gi|1036873336|ref|XP_017151999.1| PREDICTED: uncharacterized protein LOC108162017 isoform X1
>gi|1036873296|ref|XP_017151996.1| PREDICTED: uncharacterized protein LOC108162015 [Drosophila
>gi|1036880072|ref|XP_017152356.1| PREDICTED: serine-rich adhesin for platelets isoform X9 [Dro
>gi|1036879953|ref|XP_017152350.1| PREDICTED: serine-rich adhesin for platelets isoform X5 [Dro
>gi|1036879994|ref|XP_017152352.1| PREDICTED: serine-rich adhesin for platelets isoform X5 [Dro
>gi|1036879928|ref|XP_017152349.1| PREDICTED: uncharacterized protein LOC108162232 isoform X4
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>gi|1036879909|ref|XP_017152348.1| PREDICTED: uncharacterized protein LOC108162232 isoform X3
>gi|1036879816|ref|XP_017152347.1| PREDICTED: uncharacterized protein LOC108162232 isoform X2
>gi|1036879796|ref|XP_017152346.1| PREDICTED: uncharacterized protein LOC108162232 isoform X1
>gi|1036880013|ref|XP_017152353.1| PREDICTED: serine-rich adhesin for platelets isoform X6 [Dro
>gi|1036880051|ref|XP_017152355.1| PREDICTED: slowpoke-binding protein isoform X8 [Drosophila r
>gi|1036880032|ref|XP_017152354.1| PREDICTED: slowpoke-binding protein isoform X7 [Drosophila r
>gi|1036880129|ref|XP_017152359.1| PREDICTED: uncharacterized protein LOC108162235 [Drosophila
>gi|1036879646|ref|XP_017152338.1| PREDICTED: myosin-VIIa [Drosophila miranda]
>gi|1036880091|ref|XP_017152357.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1036880110|ref|XP_017152358.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1036880148|ref|XP_017152361.1| PREDICTED: uncharacterized protein LOC108162236 [Drosophila
>gi|1036879777|ref|XP_017152345.1| PREDICTED: CD109 antigen isoform X6 [Drosophila miranda]
>gi|1036879684|ref|XP_017152340.1| PREDICTED: CD109 antigen isoform X1 [Drosophila miranda]
>gi|1036879724|ref|XP_017152342.1| PREDICTED: CD109 antigen isoform X3 [Drosophila miranda]
>gi|1036879759|ref|XP_017152344.1| PREDICTED: CD109 antigen isoform X5 [Drosophila miranda]
>gi|1036879742|ref|XP_017152343.1| PREDICTED: CD109 antigen isoform X4 [Drosophila miranda]
>gi|1036879703|ref|XP_017152341.1| PREDICTED: CD109 antigen isoform X2 [Drosophila miranda]
>gi|1036879665|ref|XP_017152339.1| PREDICTED: CD109 antigen [Drosophila miranda]
>gi|1036907475|ref|XP_017153892.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036907493|ref|XP_017153893.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036907511|ref|XP_017153894.1| PREDICTED: sodium- and chloride-dependent neutral and basic
>gi|1036907439|ref|XP_017153890.1| PREDICTED: uncharacterized protein LOC108163240 [Drosophila
>gi|1036907457|ref|XP_017153891.1| PREDICTED: uncharacterized protein LOC108163240 [Drosophila
>gi|1036899105|ref|XP_017153423.1| PREDICTED: intersectin-2 isoform X3 [Drosophila miranda]
>gi|1036898965|ref|XP_017153415.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036898987|ref|XP_017153416.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036899006|ref|XP_017153417.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036899027|ref|XP_017153418.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036899046|ref|XP_017153419.1| PREDICTED: sodium- and chloride-dependent glycine transporte
>gi|1036899255|ref|XP_017153432.1| PREDICTED: adult cuticle protein 1-like [Drosophila miranda]
>gi|1036899239|ref|XP_017153431.1| PREDICTED: adult cuticle protein 1 [Drosophila miranda]
>gi|1036899294|ref|XP_017153434.1| PREDICTED: ATP synthase subunit g, mitochondrial [Drosophila
>gi|1036899275|ref|XP_017153433.1| PREDICTED: adult cuticle protein 1 [Drosophila miranda]
>gi|1036899086|ref|XP_017153422.1| PREDICTED: intersectin-2 isoform X2 [Drosophila miranda]
>gi|1036899065|ref|XP_017153420.1| PREDICTED: uncharacterized protein LOC108162942 isoform X1
>gi|1036899164|ref|XP_017153426.1| PREDICTED: PERQ amino acid-rich with GYF domain-containing p
>gi|1036899183|ref|XP_017153427.1| PREDICTED: PERQ amino acid-rich with GYF domain-containing p
>gi|1036899201|ref|XP_017153428.1| PREDICTED: PERQ amino acid-rich with GYF domain-containing p
>gi|1036899220|ref|XP_017153429.1| PREDICTED: PERQ amino acid-rich with GYF domain-containing p
>gi|1036898947|ref|XP_017153414.1| PREDICTED: uncharacterized protein LOC108162939 [Drosophila
>gi|1036899124|ref|XP_017153424.1| PREDICTED: uricase [Drosophila miranda]
>gi|1036899143|ref|XP_017153425.1| PREDICTED: uncharacterized protein LOC108162945 [Drosophila
>gi|1036903486|ref|XP_017153667.1| PREDICTED: bromodomain-containing protein 9 [Drosophila mira
>gi|1036903504|ref|XP_017153668.1| PREDICTED: ethanolaminephosphotransferase 1 [Drosophila mira
>gi|1036905151|ref|XP_017153760.1| PREDICTED: alpha-aminoadipic semialdehyde synthase, mitochor
>gi|1036905169|ref|XP_017153761.1| PREDICTED: homocysteine-responsive endoplasmic reticulum-res
>gi|1036905187|ref|XP_017153762.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036917134|ref|XP_017154427.1| PREDICTED: dual specificity protein phosphatase CDC14B [Dros
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>gi|1036892141|ref|XP_017153032.1| PREDICTED: protein transport protein SFT2 [Drosophila miran
>gi|1036892122|ref|XP_017153031.1| PREDICTED: U5 small nuclear ribonucleoprotein 40 kDa protei
>gi|1036892160|ref|XP_017153033.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prot
>gi|1036892103|ref|XP_017153030.1| PREDICTED: putative tRNA pseudouridine synthase Pus10 [Dros
>gi|1036892048|ref|XP_017153027.1| PREDICTED: nicotinate phosphoribosyltransferase isoform X1
>gi|1036892065|ref|XP_017153028.1| PREDICTED: nicotinate phosphoribosyltransferase isoform X1
>gi|1036892084|ref|XP_017153029.1| PREDICTED: nicotinate phosphoribosyltransferase isoform X2
>gi|1036926432|ref|XP_017154953.1| PREDICTED: tubulin-specific chaperone C [Drosophila miranda]
>gi|1036919014|ref|XP_017154536.1| PREDICTED: odorant receptor 24a isoform X1 [Drosophila mira
>gi|1036919031|ref|XP_017154537.1| PREDICTED: odorant receptor 24a isoform X2 [Drosophila mira
>gi|1036919049|ref|XP_017154538.1| PREDICTED: odorant receptor 24a isoform X3 [Drosophila mira
>gi|1036919084|ref|XP_017154540.1| PREDICTED: uncharacterized protein LOC108163640 [Drosophila
>gi|1036919067|ref|XP_017154539.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036918993|ref|XP_017154535.1| PREDICTED: uncharacterized protein LOC108163636 isoform X2
>gi|1036918975|ref|XP_017154534.1| PREDICTED: uncharacterized protein LOC108163636 isoform X1
>gi|1036876190|ref|XP_017152155.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036861695|ref|XP_017151349.1| PREDICTED: S-phase kinase-associated protein 1 [Drosophila
>gi|1036858905|ref|XP_017151191.1| PREDICTED: uncharacterized protein LOC108161440 [Drosophila
>gi|1036862258|ref|XP_017151379.1| PREDICTED: probable serine/threonine-protein kinase DDB_G02
>gi|1036860601|ref|XP_017151283.1| PREDICTED: uncharacterized protein LOC108161516 [Drosophila
>gi|1036857561|ref|XP_017151130.1| PREDICTED: uncharacterized protein LOC108161393 [Drosophila
>gi|1036862745|ref|XP_017151405.1| PREDICTED: leucine-rich repeat extensin-like protein 3 [Dros
>gi|1036934627|ref|XP_017155399.1| PREDICTED: LOW QUALITY PROTEIN: hsp90 co-chaperone Cdc37-li
>gi|1036891332|ref|XP_017152985.1| PREDICTED: uncharacterized protein LOC108162660 isoform X1
>gi|1036891368|ref|XP_017152988.1| PREDICTED: uncharacterized protein LOC108162660 isoform X3
>gi|1036891439|ref|XP_017152991.1| PREDICTED: E3 ubiquitin-protein ligase RFWD3-like [Drosophi
>gi|1036891351|ref|XP_017152987.1| PREDICTED: uncharacterized protein LOC108162660 isoform X2
>gi|1036891404|ref|XP_017152989.1| PREDICTED: hemicentin-1 isoform X4 [Drosophila miranda]
>gi|1036891279|ref|XP_017152982.1| PREDICTED: protein cappuccino isoform X1 [Drosophila miranda
>gi|1036891294|ref|XP_017152983.1| PREDICTED: protein cappuccino isoform X2 [Drosophila miranda
>gi|1036891313|ref|XP_017152984.1| PREDICTED: protein cappuccino isoform X3 [Drosophila miranda
>gi|1036891509|ref|XP_017152995.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 isof
>gi|1036891475|ref|XP_017152993.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 isof
>gi|1036891494|ref|XP_017152994.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 2 isof
>gi|1036891458|ref|XP_017152992.1| PREDICTED: probable RNA-binding protein EIF1AD [Drosophila
>gi|1036891421|ref|XP_017152990.1| PREDICTED: centrosomal protein of 97 kDa [Drosophila miranda
>gi|1036854808|ref|XP_017150970.1| PREDICTED: sterile alpha motif domain-containing protein 15-
>gi|1036933380|ref|XP_017155330.1| PREDICTED: uncharacterized protein LOC108164210 [Drosophila
>gi|1036916119|ref|XP_017154371.1| PREDICTED: uncharacterized protein LOC108163531 [Drosophila
>gi|1036916137|ref|XP_017154372.1| PREDICTED: uncharacterized protein LOC108163531 [Drosophila
>gi|1036916066|ref|XP_017154368.1| PREDICTED: leukotriene A-4 hydrolase isoform X1 [Drosophila
>gi|1036916084|ref|XP_017154369.1| PREDICTED: leukotriene A-4 hydrolase isoform X2 [Drosophila
>gi|1036916102|ref|XP_017154370.1| PREDICTED: 39S ribosomal protein L13, mitochondrial isoform
>gi|1036920871|ref|XP_017154642.1| PREDICTED: insulin gene enhancer protein isl-1 isoform X2 [
>gi|1036920853|ref|XP_017154641.1| PREDICTED: insulin gene enhancer protein isl-1 isoform X1 [
>gi|1036893277|ref|XP_017153098.1| PREDICTED: S phase cyclin A-associated protein in the endopl
>gi|1036893259|ref|XP_017153095.1| PREDICTED: S phase cyclin A-associated protein in the endopl
>gi|1036893295|ref|XP_017153099.1| PREDICTED: uncharacterized protein LOC108162735 [Drosophila
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>gi|1036935266|ref|XP_017155432.1| PREDICTED: uncharacterized protein LOC108164296 [Drosophila
>gi|1036859768|ref|XP_017151239.1| PREDICTED: coiled-coil domain-containing protein 170 isofor
>gi|1036859784|ref|XP_017151240.1| PREDICTED: coiled-coil domain-containing protein 170 isofor
>gi|1036935069|ref|XP_017155421.1| PREDICTED: LOW QUALITY PROTEIN: actin-related protein 2-like
>gi|1036892218|ref|XP_017153036.1| PREDICTED: uncharacterized protein LOC108162697 [Drosophila
>gi|1036892179|ref|XP_017153034.1| PREDICTED: probable cytochrome P450 310a1 [Drosophila miran
>gi|1036892197|ref|XP_017153035.1| PREDICTED: intraflagellar transport protein 46 homolog [Dros
>gi|1036923114|ref|XP_017154779.1| PREDICTED: LIM/homeobox protein Lhx2-like [Drosophila miran
>gi|1036923132|ref|XP_017154780.1| PREDICTED: uncharacterized protein LOC108163802 [Drosophila
>gi|1036923150|ref|XP_017154781.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036854827|ref|XP_017150971.1| PREDICTED: uncharacterized protein LOC108161261, partial [D
>gi|1036887865|ref|XP_017152791.1| PREDICTED: nascent polypeptide-associated complex subunit a
>gi|1036887903|ref|XP_017152793.1| PREDICTED: uncharacterized protein LOC108162523 [Drosophila
>gi|1036887922|ref|XP_017152794.1| PREDICTED: uncharacterized protein LOC108162523 [Drosophila
>gi|1036854846|ref|XP_017150972.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036887939|ref|XP_017152795.1| PREDICTED: importin subunit alpha-4-like [Drosophila miranda
>gi|1036887827|ref|XP_017152789.1| PREDICTED: uncharacterized protein LOC108162519 [Drosophila
>gi|1036887958|ref|XP_017152796.1| PREDICTED: importin subunit alpha-4-like [Drosophila miranda
>gi|1036888020|ref|XP_017152799.1| PREDICTED: acylphosphatase-1-like [Drosophila miranda]
>gi|1036887810|ref|XP_017152787.1| PREDICTED: uncharacterized protein LOC108162518 [Drosophila
>gi|1036887996|ref|XP_017152798.1| PREDICTED: putative uncharacterized protein DDB_G0294196 [D
>gi|1036887884|ref|XP_017152792.1| PREDICTED: mucin-22 [Drosophila miranda]
>gi|1036887977|ref|XP_017152797.1| PREDICTED: importin subunit alpha-4-like [Drosophila miranda
>gi|1036887846|ref|XP_017152790.1| PREDICTED: V-type proton ATPase 116 kDa subunit a [Drosophil
>gi|1036888093|ref|XP_017152800.1| PREDICTED: uncharacterized protein LOC108162530 [Drosophila
>gi|1036932194|ref|XP_017155263.1| PREDICTED: sialidase isoform X1 [Drosophila miranda]
>gi|1036932210|ref|XP_017155264.1| PREDICTED: uncharacterized protein LOC108164164 isoform X2
>gi|1036928426|ref|XP_017155061.1| PREDICTED: uncharacterized protein LOC108164000 [Drosophila
>gi|1036928409|ref|XP_017155060.1| PREDICTED: uncharacterized protein LOC108163999 [Drosophila
>gi|1036928358|ref|XP_017155057.1| PREDICTED: tetraspanin-15 isoform X1 [Drosophila miranda]
>gi|1036928375|ref|XP_017155058.1| PREDICTED: protein swallow isoform X2 [Drosophila miranda]
>gi|1036928393|ref|XP_017155059.1| PREDICTED: protein swallow isoform X2 [Drosophila miranda]
>gi|1036865836|ref|XP_017151570.1| PREDICTED: lysosomal Pro-X carboxypeptidase [Drosophila mira
>gi|1036865818|ref|XP_017151568.1| PREDICTED: nesprin-1 isoform X12 [Drosophila miranda]
>gi|1036865704|ref|XP_017151562.1| PREDICTED: nesprin-1 isoform X6 [Drosophila miranda]
>gi|1036865800|ref|XP_017151567.1| PREDICTED: nesprin-1 isoform X11 [Drosophila miranda]
>gi|1036865686|ref|XP_017151561.1| PREDICTED: nesprin-1 isoform X5 [Drosophila miranda]
>gi|1036865743|ref|XP_017151564.1| PREDICTED: nesprin-1 isoform X8 [Drosophila miranda]
>gi|1036865781|ref|XP_017151566.1| PREDICTED: nesprin-1 isoform X10 [Drosophila miranda]
>gi|1036865666|ref|XP_017151560.1| PREDICTED: nesprin-1 isoform X4 [Drosophila miranda]
>gi|1036865762|ref|XP_017151565.1| PREDICTED: nesprin-1 isoform X9 [Drosophila miranda]
>gi|1036865647|ref|XP_017151559.1| PREDICTED: nesprin-1 isoform X3 [Drosophila miranda]
>gi|1036865623|ref|XP_017151558.1| PREDICTED: nesprin-1 isoform X2 [Drosophila miranda]
>gi|1036865605|ref|XP_017151557.1| PREDICTED: nesprin-1 isoform X1 [Drosophila miranda]
>gi|1036865724|ref|XP_017151563.1| PREDICTED: nesprin-1 isoform X7 [Drosophila miranda]
>gi|1036879610|ref|XP_017152336.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prote
>gi|1036879593|ref|XP_017152335.1| PREDICTED: uncharacterized protein LOC108162225 [Drosophila
>gi|1036879574|ref|XP_017152334.1| PREDICTED: pantothenate kinase 4 [Drosophila miranda]
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>gi|1036879629|ref|XP_017152337.1| PREDICTED: 60S ribosomal protein L37a [Drosophila miranda]
>gi|1036879538|ref|XP_017152332.1| PREDICTED: polypeptide N-acetylgalactosaminyltransferase 5 :
>gi|1036879519|ref|XP_017152331.1| PREDICTED: polypeptide N-acetylgalactosaminyltransferase 5 :
>gi|1036879557|ref|XP_017152333.1| PREDICTED: polypeptide N-acetylgalactosaminyltransferase 5 :
>gi|1036876771|ref|XP_017152187.1| PREDICTED: acyl-CoA-binding protein [Drosophila miranda]
>gi|1036876752|ref|XP_017152185.1| PREDICTED: rabenosyn-5 [Drosophila miranda]
>gi|1036876714|ref|XP_017152183.1| PREDICTED: proline-rich protein 36 isoform X1 [Drosophila m:
>gi|1036876789|ref|XP_017152188.1| PREDICTED: uncharacterized protein LOC108162124 [Drosophila
>gi|1036876733|ref|XP_017152184.1| PREDICTED: proline-rich protein 36 isoform X2 [Drosophila m:
>gi|1036902379|ref|XP_017153608.1| PREDICTED: UPF0518 protein GA25918 isoform X1 [Drosophila m:
>gi|1036902417|ref|XP_017153610.1| PREDICTED: UPF0518 protein GL19323 isoform X3 [Drosophila m:
>gi|1036902397|ref|XP_017153609.1| PREDICTED: UPF0518 protein GA25918 isoform X2 [Drosophila m:
>gi|1036902439|ref|XP_017153611.1| PREDICTED: UPF0518 protein GL19323 isoform X4 [Drosophila m:
>gi|1036902457|ref|XP_017153612.1| PREDICTED: mitochondrial pyruvate carrier 2-like [Drosophil:
>gi|1036857527|ref|XP_017151129.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 1 [Dros
>gi|1036931302|ref|XP_017155215.1| PREDICTED: lipase 1-like isoform X1 [Drosophila miranda]
>gi|1036931319|ref|XP_017155216.1| PREDICTED: lipase 1-like isoform X2 [Drosophila miranda]
>gi|1036931633|ref|XP_017155236.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 1 isof
>gi|1036931653|ref|XP_017155237.1| PREDICTED: hydroxysteroid dehydrogenase-like protein 1 isof
>gi|1036870948|ref|XP_017151860.1| PREDICTED: uncharacterized protein LOC108161939 isoform X1
>gi|1036870967|ref|XP_017151861.1| PREDICTED: uncharacterized protein LOC108161939 isoform X2
>gi|1036870984|ref|XP_017151862.1| PREDICTED: phospholipase A1 member A isoform X1 [Drosophila
>gi|1036871003|ref|XP_017151863.1| PREDICTED: phospholipase A1 member A isoform X2 [Drosophila
>gi|1036870910|ref|XP_017151858.1| PREDICTED: aspartate--tRNA ligase, mitochondrial [Drosophil:
>gi|1036870929|ref|XP_017151859.1| PREDICTED: aspartate--tRNA ligase, mitochondrial [Drosophil:
>gi|1036871060|ref|XP_017151866.1| PREDICTED: diacylglycerol O-acyltransferase 1 [Drosophila m:
>gi|1036871085|ref|XP_017151867.1| PREDICTED: diacylglycerol O-acyltransferase 1 [Drosophila m:
>gi|1036871104|ref|XP_017151868.1| PREDICTED: diacylglycerol O-acyltransferase 1 [Drosophila m:
>gi|1036871142|ref|XP_017151871.1| PREDICTED: uncharacterized protein LOC108161946 [Drosophila
>gi|1036871022|ref|XP_017151864.1| PREDICTED: exportin-2 [Drosophila miranda]
>gi|1036871123|ref|XP_017151870.1| PREDICTED: trans-1,2-dihydrobenzene-1,2-diol dehydrogenase
>gi|1036871041|ref|XP_017151865.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036929866|ref|XP_017155143.1| PREDICTED: UDP-glucuronosyltransferase 2B9 [Drosophila miral
>gi|1036931942|ref|XP_017155251.1| PREDICTED: UDP-glucuronosyltransferase 2B15 [Drosophila mira
>gi|1036934746|ref|XP_017155403.1| PREDICTED: UDP-glucuronosyltransferase 1-1 [Drosophila mirar
>gi|1036908703|ref|XP_017153957.1| PREDICTED: cytochrome b5-related protein [Drosophila mirand:
>gi|1036908721|ref|XP_017153958.1| PREDICTED: cytochrome b5-related protein [Drosophila mirand:
>gi|1036908739|ref|XP_017153959.1| PREDICTED: cytochrome b5-related protein [Drosophila mirand:
>gi|1036908147|ref|XP_017153926.1| PREDICTED: myosin heavy chain, muscle isoform X2 [Drosophil:
>gi|1036908540|ref|XP_017153947.1| PREDICTED: myosin heavy chain, muscle isoform X23 [Drosophil
>gi|1036908683|ref|XP_017153956.1| PREDICTED: myosin heavy chain, muscle isoform X31 [Drosophil
>gi|1036908557|ref|XP_017153948.1| PREDICTED: myosin heavy chain, muscle isoform X24 [Drosophil
>gi|1036908500|ref|XP_017153945.1| PREDICTED: myosin heavy chain, muscle isoform X21 [Drosophil
>gi|1036908223|ref|XP_017153929.1| PREDICTED: myosin heavy chain, muscle isoform X5 [Drosophil:
>gi|1036908313|ref|XP_017153934.1| PREDICTED: myosin heavy chain, muscle isoform X10 [Drosophil
>gi|1036908375|ref|XP_017153938.1| PREDICTED: myosin heavy chain, muscle isoform X14 [Drosophil
>gi|1036908329|ref|XP_017153935.1| PREDICTED: myosin heavy chain, muscle isoform X11 [Drosophil
>gi|1036908410|ref|XP_017153940.1| PREDICTED: myosin heavy chain, muscle isoform X16 [Drosophil
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>gi|1036908239|ref|XP_017153930.1| PREDICTED: myosin heavy chain, muscle isoform X6 [Drosophila
>gi|1036908462|ref|XP_017153943.1| PREDICTED: myosin heavy chain, muscle isoform X19 [Drosophila
>gi|1036908129|ref|XP_017153925.1| PREDICTED: myosin heavy chain, muscle isoform X1 [Drosophila
>gi|1036908361|ref|XP_017153937.1| PREDICTED: myosin heavy chain, muscle isoform X13 [Drosophil
>gi|1036908257|ref|XP_017153931.1| PREDICTED: myosin heavy chain, muscle isoform X7 [Drosophila
>gi|1036908293|ref|XP_017153933.1| PREDICTED: myosin heavy chain, muscle isoform X9 [Drosophila
>gi|1036908480|ref|XP_017153944.1| PREDICTED: myosin heavy chain, muscle isoform X20 [Drosophil
>gi|1036908207|ref|XP_017153928.1| PREDICTED: myosin heavy chain, muscle isoform X4 [Drosophila
>gi|1036908428|ref|XP_017153941.1| PREDICTED: myosin heavy chain, muscle isoform X17 [Drosophil
>gi|1036908169|ref|XP_017153927.1| PREDICTED: myosin heavy chain, muscle isoform X3 [Drosophila
>gi|1036908393|ref|XP_017153939.1| PREDICTED: myosin heavy chain, muscle isoform X15 [Drosophil
>gi|1036908595|ref|XP_017153950.1| PREDICTED: myosin heavy chain, muscle isoform X26 [Drosophil
>gi|1036908446|ref|XP_017153942.1| PREDICTED: myosin heavy chain, muscle isoform X18 [Drosophil
>gi|1036908518|ref|XP_017153946.1| PREDICTED: myosin heavy chain, muscle isoform X22 [Drosophil
>gi|1036908347|ref|XP_017153936.1| PREDICTED: myosin heavy chain, muscle isoform X12 [Drosophil
>gi|1036908275|ref|XP_017153932.1| PREDICTED: myosin heavy chain, muscle isoform X8 [Drosophila
>gi|1036908575|ref|XP_017153949.1| PREDICTED: myosin heavy chain, muscle isoform X25 [Drosophil
>gi|1036908648|ref|XP_017153954.1| PREDICTED: myosin heavy chain, muscle isoform X29 [Drosophil
>gi|1036908630|ref|XP_017153953.1| PREDICTED: myosin heavy chain, muscle isoform X28 [Drosophil
>gi|1036908612|ref|XP_017153951.1| PREDICTED: myosin heavy chain, muscle isoform X27 [Drosophil
>gi|1036908665|ref|XP_017153955.1| PREDICTED: myosin heavy chain, muscle isoform X30 [Drosophil
>gi|1036930333|ref|XP_017155163.1| PREDICTED: degenerin del-1 [Drosophila miranda]
>gi|1036930297|ref|XP_017155161.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1036930315|ref|XP_017155162.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1036878186|ref|XP_017152261.1| PREDICTED: BUB3-interacting and GLEBS motif-containing prote
>gi|1036878226|ref|XP_017152264.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm7 [Drosoph
>gi|1036878169|ref|XP_017152260.1| PREDICTED: uncharacterized protein LOC108162180 [Drosophila
>gi|1036878150|ref|XP_017152259.1| PREDICTED: structural maintenance of chromosomes protein 4
>gi|1036878205|ref|XP_017152262.1| PREDICTED: cytosolic Fe-S cluster assembly factor NUBP1 homo
>gi|1036929403|ref|XP_017155116.1| PREDICTED: probable proteasome subunit beta type-2 [Drosophi
>gi|1036928978|ref|XP_017155093.1| PREDICTED: COX assembly mitochondrial protein homolog [Droso
>gi|1036928959|ref|XP_017155092.1| PREDICTED: tRNA-splicing endonuclease subunit Sen2 [Drosophi
>gi|1036909461|ref|XP_017153998.1| PREDICTED: V-type proton ATPase subunit H isoform X1 [Drosop
>gi|1036909479|ref|XP_017153999.1| PREDICTED: V-type proton ATPase subunit H isoform X2 [Drosop
>gi|1036909501|ref|XP_017154000.1| PREDICTED: cytochrome c-2 [Drosophila miranda]
>gi|1036916998|ref|XP_017154419.1| PREDICTED: cytochrome c-1 [Drosophila miranda]
>gi|1036917018|ref|XP_017154420.1| PREDICTED: cytochrome c-1 [Drosophila miranda]
>gi|1036917036|ref|XP_017154421.1| PREDICTED: cytochrome c-1 [Drosophila miranda]
>gi|1036917054|ref|XP_017154422.1| PREDICTED: uncharacterized protein LOC108163566 [Drosophila
>gi|1036862041|ref|XP_017151366.1| PREDICTED: CTD nuclear envelope phosphatase 1 [Drosophila m
>gi|1036918721|ref|XP_017154521.1| PREDICTED: solute carrier family 25 member 36 isoform X1 [Dr
>gi|1036918739|ref|XP_017154522.1| PREDICTED: solute carrier family 25 member 36 isoform X2 [Dr
>gi|1036891654|ref|XP_017153004.1| PREDICTED: WD repeat-containing and planar cell polarity ef
>gi|1036891711|ref|XP_017153007.1| PREDICTED: 39S ribosomal protein L48, mitochondrial [Drosoph
>gi|1036891673|ref|XP_017153005.1| PREDICTED: transmembrane protein 87A [Drosophila miranda]
>gi|1036891692|ref|XP_017153006.1| PREDICTED: F-actin-capping protein subunit beta [Drosophila
>gi|1036905886|ref|XP_017153806.1| PREDICTED: probable tRNA (guanine(26)-N(2))-dimethyltransfer
>gi|1036905899|ref|XP_017153807.1| PREDICTED: uncharacterized protein LOC108163181 [Drosophila
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>gi|1036905913|ref|XP_017153808.1| PREDICTED: serine/arginine-rich splicing factor 2 [Drosophil
>gi|1036914940|ref|XP_017154296.1| PREDICTED: eukaryotic peptide chain release factor GTP-bind
>gi|1036914958|ref|XP_017154297.1| PREDICTED: eukaryotic peptide chain release factor GTP-bind
>gi|1036914976|ref|XP_017154298.1| PREDICTED: eukaryotic peptide chain release factor GTP-bind
>gi|1036915012|ref|XP_017154300.1| PREDICTED: eukaryotic peptide chain release factor GTP-bind
>gi|1036914994|ref|XP_017154299.1| PREDICTED: eukaryotic peptide chain release factor GTP-bind
>gi|1036927149|ref|XP_017154995.1| PREDICTED: uncharacterized protein LOC108163957 [Drosophila
>gi|1036927111|ref|XP_017154993.1| PREDICTED: LOW QUALITY PROTEIN: solute carrier organic anio
>gi|1036927131|ref|XP_017154994.1| PREDICTED: uncharacterized protein LOC108163956 [Drosophila
>gi|1036872561|ref|XP_017151955.1| PREDICTED: solute carrier organic anion transporter family
>gi|1036872708|ref|XP_017151963.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X7
>gi|1036872689|ref|XP_017151961.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X6
>gi|1036872601|ref|XP_017151957.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X2
>gi|1036872580|ref|XP_017151956.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X1
>gi|1036872743|ref|XP_017151965.1| PREDICTED: protein bunched, class 1/class 3/D/E isoforms iso
>gi|1036872762|ref|XP_017151966.1| PREDICTED: protein bunched, class 1/class 3/D/E isoforms iso
>gi|1036872780|ref|XP_017151967.1| PREDICTED: protein bunched, class 1/class 3/D/E isoforms iso
>gi|1036872724|ref|XP_017151964.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X8
>gi|1036872648|ref|XP_017151959.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X4
>gi|1036872620|ref|XP_017151958.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X3
>gi|1036872669|ref|XP_017151960.1| PREDICTED: protein bunched, class 2/F/G isoform isoform X5
>gi|1036915263|ref|XP_017154318.1| PREDICTED: protein nubbin isoform X1 [Drosophila miranda]
>gi|1036915281|ref|XP_017154319.1| PREDICTED: protein nubbin isoform X2 [Drosophila miranda]
>gi|1036863418|ref|XP_017151442.1| PREDICTED: uncharacterized protein LOC108161650 [Drosophila
>gi|1036913549|ref|XP_017154218.1| PREDICTED: POU domain protein 2 isoform X1 [Drosophila mira
>gi|1036913585|ref|XP_017154220.1| PREDICTED: neprilysin [Drosophila miranda]
>gi|1036913567|ref|XP_017154219.1| PREDICTED: POU domain protein 2 isoform X2 [Drosophila mira
>gi|1036913601|ref|XP_017154221.1| PREDICTED: uncharacterized protein LOC108163447 [Drosophila
>gi|1036894033|ref|XP_017153137.1| PREDICTED: leucine-rich repeat and calponin homology domain
>gi|1036894052|ref|XP_017153138.1| PREDICTED: leucine-rich repeat and calponin homology domain
>gi|1036894014|ref|XP_017153136.1| PREDICTED: leucine-rich repeat and calponin homology domain
>gi|1036873520|ref|XP_017152006.1| PREDICTED: restin homolog isoform X1 [Drosophila miranda]
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>gi|1036873598|ref|XP_017152010.1| PREDICTED: restin homolog isoform X4 [Drosophila miranda]
>gi|1036873578|ref|XP_017152009.1| PREDICTED: restin homolog isoform X3 [Drosophila miranda]
>gi|1036873560|ref|XP_017152008.1| PREDICTED: restin homolog isoform X2 [Drosophila miranda]
>gi|1036873617|ref|XP_017152011.1| PREDICTED: restin homolog isoform X5 [Drosophila miranda]
>gi|1036873636|ref|XP_017152012.1| PREDICTED: restin homolog isoform X6 [Drosophila miranda]
>gi|1036873654|ref|XP_017152013.1| PREDICTED: restin homolog isoform X7 [Drosophila miranda]
>gi|1036873867|ref|XP_017152026.1| PREDICTED: restin homolog isoform X19 [Drosophila miranda]
>gi|1036873670|ref|XP_017152014.1| PREDICTED: restin homolog isoform X8 [Drosophila miranda]
>gi|1036873885|ref|XP_017152027.1| PREDICTED: restin homolog isoform X20 [Drosophila miranda]
>gi|1036873706|ref|XP_017152016.1| PREDICTED: restin homolog isoform X10 [Drosophila miranda]
>gi|1036873922|ref|XP_017152030.1| PREDICTED: restin homolog isoform X22 [Drosophila miranda]
>gi|1036873688|ref|XP_017152015.1| PREDICTED: restin homolog isoform X9 [Drosophila miranda]
>gi|1036873722|ref|XP_017152017.1| PREDICTED: restin homolog isoform X11 [Drosophila miranda]
>gi|1036873903|ref|XP_017152028.1| PREDICTED: restin homolog isoform X21 [Drosophila miranda]
>gi|1036873797|ref|XP_017152022.1| PREDICTED: restin homolog isoform X15 [Drosophila miranda]
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>gi|1036873834|ref|XP_017152024.1| PREDICTED: restin homolog isoform X17 [Drosophila miranda]
>gi|1036873815|ref|XP_017152023.1| PREDICTED: restin homolog isoform X16 [Drosophila miranda]
>gi|1036873849|ref|XP_017152025.1| PREDICTED: restin homolog isoform X18 [Drosophila miranda]
>gi|1036873740|ref|XP_017152019.1| PREDICTED: restin homolog isoform X12 [Drosophila miranda]
>gi|1036873757|ref|XP_017152020.1| PREDICTED: restin homolog isoform X13 [Drosophila miranda]
>gi|1036873774|ref|XP_017152021.1| PREDICTED: restin homolog isoform X14 [Drosophila miranda]
>gi|1036874031|ref|XP_017152036.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB11 [Dro
>gi|1036873995|ref|XP_017152034.1| PREDICTED: putative E3 ubiquitin-protein ligase UBR7 [Drosop
>gi|1036874012|ref|XP_017152035.1| PREDICTED: radial spoke head protein 9 homolog [Drosophila m
>gi|1036873976|ref|XP_017152033.1| PREDICTED: dorsal-related immunity factor Dif isoform X2 [D
>gi|1036873940|ref|XP_017152031.1| PREDICTED: embryonic polarity protein dorsal isoform X1 [Dro
>gi|1036873957|ref|XP_017152032.1| PREDICTED: embryonic polarity protein dorsal isoform X1 [Dro
>gi|1036929488|ref|XP_017155123.1| PREDICTED: uncharacterized protein LOC108164054 [Drosophila
>gi|1036854865|ref|XP_017150973.1| PREDICTED: uncharacterized protein LOC108161265 [Drosophila
>gi|1036895226|ref|XP_017153206.1| PREDICTED: embryonic polarity protein dorsal-like [Drosophil
>gi|1036895245|ref|XP_017153207.1| PREDICTED: embryonic polarity protein dorsal-like [Drosophil
>gi|1036852646|ref|XP_017150834.1| PREDICTED: embryonic polarity protein dorsal-like, partial
>gi|1036895283|ref|XP_017153209.1| PREDICTED: uncharacterized protein LOC108162809 [Drosophila
>gi|1036895321|ref|XP_017153211.1| PREDICTED: golgin subfamily B member 1-like [Drosophila mira
>gi|1036895302|ref|XP_017153210.1| PREDICTED: short-chain dehydrogenase/reductase family 16C me
>gi|1036895264|ref|XP_017153208.1| PREDICTED: probable ATP-dependent RNA helicase DHX35 [Drosop
>gi|1036928741|ref|XP_017155079.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036905711|ref|XP_017153793.1| PREDICTED: uncharacterized protein LOC108163172 [Drosophila
>gi|1036905729|ref|XP_017153794.1| PREDICTED: protein farnesyltransferase/geranylgeranyltransfe
>gi|1036933362|ref|XP_017155329.1| PREDICTED: glutamate receptor ionotropic, kainate 4 [Drosop
>gi|1036920552|ref|XP_017154622.1| PREDICTED: etoposide-induced protein 2.4 homolog isoform X2
>gi|1036920532|ref|XP_017154621.1| PREDICTED: etoposide-induced protein 2.4 isoform X1 [Drosop
>gi|1036857764|ref|XP_017151143.1| PREDICTED: uncharacterized protein LOC108161403 [Drosophila
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>gi|1036919380|ref|XP_017154550.1| PREDICTED: eukaryotic translation initiation factor 4E-bindi
>gi|1036919344|ref|XP_017154548.1| PREDICTED: eukaryotic translation initiation factor 4E-bindi
>gi|1036919362|ref|XP_017154549.1| PREDICTED: eukaryotic translation initiation factor 4E-bindi
>gi|1036916329|ref|XP_017154384.1| PREDICTED: uncharacterized protein LOC108163539 isoform X1
>gi|1036916346|ref|XP_017154385.1| PREDICTED: uncharacterized protein LOC108163539 isoform X2
>gi|1036924364|ref|XP_017154840.1| PREDICTED: protein grindelwald isoform X1 [Drosophila mirand
>gi|1036924381|ref|XP_017154841.1| PREDICTED: protein grindelwald isoform X2 [Drosophila mirand
>gi|1036900003|ref|XP_017153472.1| PREDICTED: chondroitin sulfate proteoglycan 4 [Drosophila mi
>gi|1036900018|ref|XP_017153473.1| PREDICTED: chondroitin sulfate proteoglycan 4 [Drosophila mi
>gi|1036900032|ref|XP_017153474.1| PREDICTED: chondroitin sulfate proteoglycan 4 [Drosophila mi
>gi|1036900062|ref|XP_017153476.1| PREDICTED: UDP-glucuronosyltransferase 2B13 [Drosophila mira
>gi|1036900047|ref|XP_017153475.1| PREDICTED: uncharacterized protein LOC108162980 [Drosophila
>gi|1036931198|ref|XP_017155210.1| PREDICTED: uncharacterized protein LOC108164124 [Drosophila
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>gi|1036931216|ref|XP_017155211.1| PREDICTED: uncharacterized protein LOC108164124 [Drosophila
>gi|1036898375|ref|XP_017153383.1| PREDICTED: dual 3',5'-cyclic-AMP and -GMP phosphodiesterase
>gi|1036898356|ref|XP_017153382.1| PREDICTED: dual 3',5'-cyclic-AMP and -GMP phosphodiesterase
>gi|1036898395|ref|XP_017153384.1| PREDICTED: dual 3',5'-cyclic-AMP and -GMP phosphodiesterase
>gi|1036854883|ref|XP_017150974.1| PREDICTED: mitochondrial import receptor subunit TOM20 homol
>gi|1036911367|ref|XP_017154092.1| PREDICTED: uncharacterized protein LOC108163359 isoform X1
>gi|1036911385|ref|XP_017154093.1| PREDICTED: uncharacterized protein LOC108163359 isoform X2
>gi|1036857113|ref|XP_017151108.1| PREDICTED: basic helix-loop-helix transcription factor amos
>gi|1036886680|ref|XP_017152721.1| PREDICTED: solute carrier family 12 member 9 [Drosophila mi
>gi|1036886697|ref|XP_017152722.1| PREDICTED: probable ATP-dependent RNA helicase DDX23 [Droso
>gi|1036886716|ref|XP_017152723.1| PREDICTED: cysteine-rich protein 2-binding protein [Drosophi
>gi|1036892359|ref|XP_017153044.1| PREDICTED: 60S ribosomal protein L44 [Drosophila miranda]
>gi|1036892343|ref|XP_017153043.1| PREDICTED: WASH complex subunit CCDC53 homolog [Drosophila m
>gi|1036892326|ref|XP_017153042.1| PREDICTED: striatin-3 isoform X4 [Drosophila miranda]
>gi|1036892290|ref|XP_017153040.1| PREDICTED: striatin isoform X2 [Drosophila miranda]
>gi|1036892307|ref|XP_017153041.1| PREDICTED: striatin-4 isoform X3 [Drosophila miranda]
>gi|1036892237|ref|XP_017153037.1| PREDICTED: striatin-4 isoform X1 [Drosophila miranda]
>gi|1036892256|ref|XP_017153038.1| PREDICTED: striatin-4 isoform X1 [Drosophila miranda]
>gi|1036892273|ref|XP_017153039.1| PREDICTED: striatin-4 isoform X1 [Drosophila miranda]
>gi|1036892371|ref|XP_017153046.1| PREDICTED: barrier-to-autointegration factor [Drosophila mi
>gi|1036920799|ref|XP_017154638.1| PREDICTED: uncharacterized protein LOC108163707 isoform X1
>gi|1036920817|ref|XP_017154639.1| PREDICTED: uncharacterized protein LOC108163707 isoform X2
>gi|1036920835|ref|XP_017154640.1| PREDICTED: LOW QUALITY PROTEIN: protein mitoshell [Drosophil
>gi|1036866448|ref|XP_017151603.1| PREDICTED: LOW QUALITY PROTEIN: multidrug resistance-associa
>gi|1036866562|ref|XP_017151610.1| PREDICTED: coiled-coil domain-containing protein 97 [Drosoph
>gi|1036866507|ref|XP_017151606.1| PREDICTED: methyltransferase-like protein 14 homolog [Drosop
>gi|1036866579|ref|XP_017151611.1| PREDICTED: tumor suppressor candidate 3 [Drosophila miranda]
>gi|1036866615|ref|XP_017151614.1| PREDICTED: polyprenol reductase [Drosophila miranda]
>gi|1036866544|ref|XP_017151609.1| PREDICTED: epsilon-sarcoglycan isoform X2 [Drosophila miran
>gi|1036866526|ref|XP_017151608.1| PREDICTED: epsilon-sarcoglycan isoform X1 [Drosophila miran
>gi|1036866597|ref|XP_017151613.1| PREDICTED: dolichyl-phosphate beta-glucosyltransferase [Dros
>gi|1036866467|ref|XP_017151604.1| PREDICTED: tyrosine-protein kinase Btk29A isoform X1 [Drosop
>gi|1036866488|ref|XP_017151605.1| PREDICTED: tyrosine-protein kinase Btk29A isoform X2 [Drosop
>gi|1036921649|ref|XP_017154688.1| PREDICTED: uncharacterized protein LOC108163740 isoform X1
>gi|1036921667|ref|XP_017154689.1| PREDICTED: uncharacterized protein LOC108163740 isoform X2
>gi|1036921737|ref|XP_017154694.1| PREDICTED: uncharacterized protein LOC108163740 isoform X6
>gi|1036921719|ref|XP_017154692.1| PREDICTED: extensin-2 isoform X5 [Drosophila miranda]
>gi|1036921703|ref|XP_017154691.1| PREDICTED: uncharacterized protein LOC108163740 isoform X4
>gi|1036921773|ref|XP_017154696.1| PREDICTED: uncharacterized protein LOC108163740 isoform X8
>gi|1036921755|ref|XP_017154695.1| PREDICTED: uncharacterized protein LOC108163740 isoform X7
>gi|1036921794|ref|XP_017154697.1| PREDICTED: uncharacterized protein LOC108163740 isoform X9
>gi|1036921685|ref|XP_017154690.1| PREDICTED: uncharacterized protein LOC108163740 isoform X3
>gi|1036921811|ref|XP_017154698.1| PREDICTED: uncharacterized protein LOC108163742 [Drosophila
>gi|1036854902|ref|XP_017150976.1| PREDICTED: outer dense fiber protein 3-like protein 2 [Dros
>gi|1036858849|ref|XP_017151188.1| PREDICTED: uncharacterized protein LOC108161436 [Drosophila
>gi|1036858830|ref|XP_017151187.1| PREDICTED: uncharacterized protein LOC108161435 [Drosophila
>gi|1036913153|ref|XP_017154194.1| PREDICTED: vascular endothelial growth factor receptor 3 iso
>gi|1036912974|ref|XP_017154183.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
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>gi|1036912993|ref|XP_017154184.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036913013|ref|XP_017154185.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036913100|ref|XP_017154191.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036913082|ref|XP_017154190.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036913049|ref|XP_017154188.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036913118|ref|XP_017154192.1| PREDICTED: vascular endothelial growth factor receptor 3 iso
>gi|1036913031|ref|XP_017154187.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036913136|ref|XP_017154193.1| PREDICTED: vascular endothelial growth factor receptor 3 iso
>gi|1036913064|ref|XP_017154189.1| PREDICTED: vascular endothelial growth factor receptor 1 iso
>gi|1036894619|ref|XP_017153170.1| PREDICTED: sorting nexin-32 isoform X1 [Drosophila miranda]
>gi|1036894637|ref|XP_017153172.1| PREDICTED: sorting nexin-32 isoform X2 [Drosophila miranda]
>gi|1036894656|ref|XP_017153173.1| PREDICTED: uncharacterized protein LOC108162784 [Drosophila
>gi|1036894677|ref|XP_017153174.1| PREDICTED: cytidine deaminase [Drosophila miranda]
>gi|1036894773|ref|XP_017153179.1| PREDICTED: cytidine deaminase [Drosophila miranda]
>gi|1036894794|ref|XP_017153180.1| PREDICTED: cytidine deaminase [Drosophila miranda]
>gi|1036894736|ref|XP_017153177.1| PREDICTED: transmembrane protein 222 isoform X1 [Drosophila
>gi|1036894754|ref|XP_017153178.1| PREDICTED: transmembrane protein 222 isoform X2 [Drosophila
>gi|1036894813|ref|XP_017153181.1| PREDICTED: RNA polymerase II transcriptional coactivator [Dr
>gi|1036894696|ref|XP_017153175.1| PREDICTED: heterochromatin protein 1 isoform X1 [Drosophila
>gi|1036894717|ref|XP_017153176.1| PREDICTED: heterochromatin protein 1 isoform X2 [Drosophila
>gi|1036894600|ref|XP_017153169.1| PREDICTED: tripartite motif-containing protein 45 [Drosophil
>gi|1036905091|ref|XP_017153756.1| PREDICTED: sodium-coupled monocarboxylate transporter 2-like
>gi|1036854924|ref|XP_017150977.1| PREDICTED: LOW QUALITY PROTEIN: sodium-coupled monocarboxyla
>gi|1036905129|ref|XP_017153759.1| PREDICTED: metallophosphoesterase 1 homolog [Drosophila mira
>gi|1036905109|ref|XP_017153758.1| PREDICTED: chitinase domain-containing protein 1 [Drosophila
>gi|1036891113|ref|XP_017152971.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036891130|ref|XP_017152972.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036891167|ref|XP_017152975.1| PREDICTED: uncharacterized protein LOC108162653 [Drosophila
>gi|1036891186|ref|XP_017152976.1| PREDICTED: uncharacterized protein LOC108162654 [Drosophila
>gi|1036891203|ref|XP_017152978.1| PREDICTED: uncharacterized protein LOC108162655 [Drosophila
>gi|1036891148|ref|XP_017152974.1| PREDICTED: trans-1,2-dihydrobenzene-1,2-diol dehydrogenase
>gi|1036915509|ref|XP_017154334.1| PREDICTED: trans-1,2-dihydrobenzene-1,2-diol dehydrogenase
>gi|1036915491|ref|XP_017154333.1| PREDICTED: uncharacterized protein LOC108163512 [Drosophila
>gi|1036913169|ref|XP_017154195.1| PREDICTED: ras-related protein Rab-5B [Drosophila miranda]
>gi|1036913187|ref|XP_017154197.1| PREDICTED: ras-related protein Rab-5B [Drosophila miranda]
>gi|1036913207|ref|XP_017154198.1| PREDICTED: ras-related protein RABF2a-like [Drosophila mirar
>gi|1036911011|ref|XP_017154073.1| PREDICTED: protein eyes shut [Drosophila miranda]
>gi|1036911027|ref|XP_017154074.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036911082|ref|XP_017154077.1| PREDICTED: MKRN2 opposite strand protein [Drosophila miranda
>gi|1036911100|ref|XP_017154078.1| PREDICTED: MKRN2 opposite strand protein [Drosophila miranda
>gi|1036911122|ref|XP_017154079.1| PREDICTED: MKRN2 opposite strand protein [Drosophila miranda
>gi|1036911196|ref|XP_017154084.1| PREDICTED: protein PRY1 [Drosophila miranda]
>gi|1036854938|ref|XP_017150978.1| PREDICTED: uncharacterized protein LOC108161269 [Drosophila
>gi|1036911064|ref|XP_017154076.1| PREDICTED: serine/threonine-protein phosphatase 4 catalytic
>gi|1036911160|ref|XP_017154081.1| PREDICTED: vesicle-associated membrane protein-associated pr
>gi|1036911214|ref|XP_017154085.1| PREDICTED: dynein light chain roadblock-type 2 [Drosophila r
>gi|1036854955|ref|XP_017150979.1| PREDICTED: myosin-2 essential light chain [Drosophila mirand
>gi|1036911046|ref|XP_017154075.1| PREDICTED: uncharacterized protein LOC108163346 [Drosophila
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>gi|1036911140|ref|XP_017154080.1| PREDICTED: trypsin alpha-like [Drosophila miranda]
>gi|1036911178|ref|XP_017154083.1| PREDICTED: Golgi-associated plant pathogenesis-related prote
>gi|1036859125|ref|XP_017151203.1| PREDICTED: longitudinals lacking protein-like [Drosophila mi
>gi|1036859146|ref|XP_017151205.1| PREDICTED: uncharacterized protein LOC108161454 [Drosophila
>gi|1036930681|ref|XP_017155183.1| PREDICTED: nardilysin-like [Drosophila miranda]
>gi|1036856532|ref|XP_017151073.1| PREDICTED: probable nuclear transport factor 2 isoform X2 [
>gi|1036856515|ref|XP_017151072.1| PREDICTED: probable nuclear transport factor 2 isoform X1 [
>gi|1036898885|ref|XP_017153411.1| PREDICTED: endochitinase A1-like [Drosophila miranda]
>gi|1036898903|ref|XP_017153412.1| PREDICTED: phospholipase A1 2 [Drosophila miranda]
>gi|1036898924|ref|XP_017153413.1| PREDICTED: endonuclease G, mitochondrial-like [Drosophila mi
>gi|1036898867|ref|XP_017153410.1| PREDICTED: protein transport protein Sec24C [Drosophila mira
>gi|1036875289|ref|XP_017152106.1| PREDICTED: Golgi phosphoprotein 3 homolog sauron [Drosophila
>gi|1036875371|ref|XP_017152111.1| PREDICTED: protein C10 [Drosophila miranda]
>gi|1036875235|ref|XP_017152103.1| PREDICTED: tudor and KH domain-containing protein [Drosophil
>gi|1036875254|ref|XP_017152104.1| PREDICTED: tudor and KH domain-containing protein [Drosophil
>gi|1036875270|ref|XP_017152105.1| PREDICTED: tudor and KH domain-containing protein [Drosophil
>gi|1036875191|ref|XP_017152101.1| PREDICTED: WD repeat-containing protein mio [Drosophila mira
>gi|1036875392|ref|XP_017152112.1| PREDICTED: protein CEBPZOS [Drosophila miranda]
>gi|1036875408|ref|XP_017152113.1| PREDICTED: uncharacterized protein LOC108162076 [Drosophila
>gi|1036875306|ref|XP_017152107.1| PREDICTED: uncharacterized protein LOC108162071 [Drosophila
>gi|1036875352|ref|XP_017152110.1| PREDICTED: uncharacterized protein LOC108162073 [Drosophila
>gi|1036875333|ref|XP_017152108.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase [Drosophila
>gi|1036875209|ref|XP_017152102.1| PREDICTED: 2-phosphoxylose phosphatase 1 [Drosophila miranda
>gi|1036932424|ref|XP_017155276.1| PREDICTED: cytochrome c oxidase subunit 7A1, mitochondrial
>gi|1036922653|ref|XP_017154751.1| PREDICTED: uncharacterized protein LOC108163783 isoform X1
>gi|1036922666|ref|XP_017154752.1| PREDICTED: uncharacterized protein LOC108163783 isoform X2
>gi|1036922680|ref|XP_017154753.1| PREDICTED: coiled-coil domain-containing protein 28B isoform
>gi|1036922695|ref|XP_017154754.1| PREDICTED: coiled-coil domain-containing protein 28B isoform
>gi|1036882549|ref|XP_017152495.1| PREDICTED: folylpolyglutamate synthase, mitochondrial-like
>gi|1036882512|ref|XP_017152493.1| PREDICTED: ets DNA-binding protein pokkuri [Drosophila miran
>gi|1036882531|ref|XP_017152494.1| PREDICTED: ets DNA-binding protein pokkuri [Drosophila miran
>gi|1036882568|ref|XP_017152496.1| PREDICTED: uncharacterized protein LOC108162323 [Drosophila
>gi|1036882494|ref|XP_017152492.1| PREDICTED: putative apoptosis-inducing factor 1, mitochondri
>gi|1036882477|ref|XP_017152491.1| PREDICTED: putative apoptosis-inducing factor 1, mitochondri
>gi|1036904916|ref|XP_017153746.1| PREDICTED: tubulin-specific chaperone D [Drosophila miranda]
>gi|1036904934|ref|XP_017153747.1| PREDICTED: uncharacterized SDCCAG3 family protein [Drosophil
>gi|1036885082|ref|XP_017152639.1| PREDICTED: THO complex subunit 2 [Drosophila miranda]
>gi|1036885100|ref|XP_017152640.1| PREDICTED: glycogen phosphorylase [Drosophila miranda]
>gi|1036914922|ref|XP_017154295.1| PREDICTED: leucine-rich PPR motif-containing protein, mitoch
>gi|1036910851|ref|XP_017154065.1| PREDICTED: pre-mRNA-splicing factor CWC22 homolog [Drosophil
>gi|1036920137|ref|XP_017154596.1| PREDICTED: 40S ribosomal protein S26 [Drosophila miranda]
>gi|1036920119|ref|XP_017154595.1| PREDICTED: pyruvate dehydrogenase E1 component subunit alpha
>gi|1036861429|ref|XP_017151332.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036861411|ref|XP_017151331.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036890949|ref|XP_017152961.1| PREDICTED: fasciclin-3 isoform X3 [Drosophila miranda]
>gi|1036890931|ref|XP_017152960.1| PREDICTED: fasciclin-3 isoform X2 [Drosophila miranda]
>gi|1036890985|ref|XP_017152963.1| PREDICTED: fasciclin-3 isoform X5 [Drosophila miranda]
>gi|1036890893|ref|XP_017152958.1| PREDICTED: fasciclin-3 isoform X1 [Drosophila miranda]
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>gi|1036890912|ref|XP_017152959.1| PREDICTED: fasciclin-3 isoform X1 [Drosophila miranda]
>gi|1036890967|ref|XP_017152962.1| PREDICTED: fasciclin-3 isoform X4 [Drosophila miranda]
>gi|1036891003|ref|XP_017152965.1| PREDICTED: factor V activator RVV-V alpha [Drosophila miran
>gi|1036932974|ref|XP_017155308.1| PREDICTED: cell division cycle 7-related protein kinase [Dro
>gi|1036860505|ref|XP_017151278.1| PREDICTED: transmembrane emp24 domain-containing protein 1
>gi|1036857928|ref|XP_017151151.1| PREDICTED: serine/threonine-protein kinase polo-like [Droso
>gi|1036907080|ref|XP_017153872.1| PREDICTED: uncharacterized protein LOC108163227 [Drosophila
>gi|1036907098|ref|XP_017153873.1| PREDICTED: G2/mitotic-specific cyclin-B-like [Drosophila mi
>gi|1036907154|ref|XP_017153877.1| PREDICTED: uncharacterized protein LOC108163230 isoform X2
>gi|1036907134|ref|XP_017153876.1| PREDICTED: uncharacterized protein LOC108163230 isoform X1
>gi|1036907172|ref|XP_017153878.1| PREDICTED: chymotrypsin inhibitor SCI-III-like [Drosophila
>gi|1036907045|ref|XP_017153870.1| PREDICTED: uncharacterized protein LOC108163224 [Drosophila
>gi|1036907117|ref|XP_017153875.1| PREDICTED: caltractin [Drosophila miranda]
>gi|1036907062|ref|XP_017153871.1| PREDICTED: receptor-type tyrosine-protein phosphatase kappa
>gi|1036918149|ref|XP_017154488.1| PREDICTED: jmjC domain-containing protein 4 homolog [Drosopl
>gi|1036918184|ref|XP_017154490.1| PREDICTED: uncharacterized protein LOC108163612 [Drosophila
>gi|1036918202|ref|XP_017154492.1| PREDICTED: uncharacterized protein LOC108163613 [Drosophila
>gi|1036918167|ref|XP_017154489.1| PREDICTED: uncharacterized protein LOC108163611 [Drosophila
>gi|1036857132|ref|XP_017151109.1| PREDICTED: uncharacterized protein LOC108161373 [Drosophila
>gi|1036857151|ref|XP_017151110.1| PREDICTED: uncharacterized protein LOC108161373 [Drosophila
>gi|1036857170|ref|XP_017151111.1| PREDICTED: uncharacterized protein LOC108161373 [Drosophila
>gi|1036934575|ref|XP_017155395.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036864724|ref|XP_017151507.1| PREDICTED: transcription factor hamlet isoform X1 [Drosophil
>gi|1036864743|ref|XP_017151509.1| PREDICTED: transcription factor hamlet isoform X2 [Drosophil
>gi|1036864947|ref|XP_017151521.1| PREDICTED: uncharacterized protein LOC108161712 [Drosophila
>gi|1036864966|ref|XP_017151522.1| PREDICTED: uncharacterized protein LOC108161713 [Drosophila
>gi|1036864985|ref|XP_017151523.1| PREDICTED: uncharacterized protein LOC108161714 [Drosophila
>gi|1036864928|ref|XP_017151519.1| PREDICTED: uncharacterized protein LOC108161711 [Drosophila
>gi|1036854974|ref|XP_017150980.1| PREDICTED: UDP-glucuronosyltransferase 1-3 [Drosophila mira
>gi|1036864873|ref|XP_017151516.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036864817|ref|XP_017151513.1| PREDICTED: UDP-glucuronosyltransferase 2B17 isoform X1 [Dros
>gi|1036864836|ref|XP_017151514.1| PREDICTED: UDP-glucuronosyltransferase 2B17 isoform X1 [Dros
>gi|1036864855|ref|XP_017151515.1| PREDICTED: UDP-glucuronosyltransferase 2B17 isoform X2 [Dros
>gi|1036864910|ref|XP_017151518.1| PREDICTED: non-specific lipid-transfer protein [Drosophila
>gi|1036864706|ref|XP_017151506.1| PREDICTED: condensin complex subunit 1 [Drosophila miranda]
>gi|1036864798|ref|XP_017151512.1| PREDICTED: non-specific lipid-transfer protein [Drosophila
>gi|1036864668|ref|XP_017151504.1| PREDICTED: uncharacterized protein LOC108161701 isoform X2
>gi|1036864687|ref|XP_017151505.1| PREDICTED: uncharacterized protein LOC108161701 isoform X3
>gi|1036864652|ref|XP_017151503.1| PREDICTED: uncharacterized protein LOC108161701 isoform X1
>gi|1036864781|ref|XP_017151511.1| PREDICTED: tubulin polyglutamylase TTLL4 isoform X2 [Drosopl
>gi|1036864762|ref|XP_017151510.1| PREDICTED: tubulin polyglutamylase TTLL4 isoform X1 [Drosopl
>gi|1036864891|ref|XP_017151517.1| PREDICTED: fork head domain transcription factor slp2 [Droso
>gi|1036929506|ref|XP_017155124.1| PREDICTED: fork head domain transcription factor slp1 [Droso
>gi|1036859184|ref|XP_017151207.1| PREDICTED: uncharacterized protein LOC108161456 [Drosophila
>gi|1036892390|ref|XP_017153047.1| PREDICTED: uncharacterized protein LOC108162703 [Drosophila
>gi|1036892444|ref|XP_017153050.1| PREDICTED: CCR4-NOT transcription complex subunit 3 isoform
>gi|1036892462|ref|XP_017153051.1| PREDICTED: CCR4-NOT transcription complex subunit 3 isoform
>gi|1036892481|ref|XP_017153052.1| PREDICTED: CCR4-NOT transcription complex subunit 3 isoform
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>gi|1036892425|ref|XP_017153049.1| PREDICTED: protein bark beetle isoform X2 [Drosophila miran
>gi|1036892408|ref|XP_017153048.1| PREDICTED: protein bark beetle isoform X1 [Drosophila miran
>gi|1036914393|ref|XP_017154262.1| PREDICTED: protein bowel [Drosophila miranda]
>gi|1036914411|ref|XP_017154263.1| PREDICTED: protein bowel [Drosophila miranda]
>gi|1036914430|ref|XP_017154264.1| PREDICTED: proteasome subunit alpha type-3-like [Drosophila
>gi|1036929122|ref|XP_017155101.1| PREDICTED: uncharacterized protein LOC108164033 isoform X4
>gi|1036929104|ref|XP_017155100.1| PREDICTED: uncharacterized protein LOC108164033 isoform X3
>gi|1036929087|ref|XP_017155099.1| PREDICTED: uncharacterized protein LOC108164033 isoform X2
>gi|1036929065|ref|XP_017155098.1| PREDICTED: uncharacterized protein LOC108164033 isoform X1
>gi|1036931786|ref|XP_017155244.1| PREDICTED: monocarboxylate transporter 8 [Drosophila miranda
>gi|1036931804|ref|XP_017155245.1| PREDICTED: uncharacterized protein LOC108164152 [Drosophila
>gi|1036929544|ref|XP_017155126.1| PREDICTED: potassium voltage-gated channel protein Shaw iso
>gi|1036929575|ref|XP_017155128.1| PREDICTED: potassium voltage-gated channel protein Shaw iso
>gi|1036929561|ref|XP_017155127.1| PREDICTED: potassium voltage-gated channel protein Shaw iso
>gi|1036929613|ref|XP_017155130.1| PREDICTED: potassium voltage-gated channel protein Shaw iso
>gi|1036929595|ref|XP_017155129.1| PREDICTED: potassium voltage-gated channel protein Shaw iso
>gi|1036900092|ref|XP_017153478.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [
>gi|1036900107|ref|XP_017153479.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [
>gi|1036900122|ref|XP_017153480.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [
>gi|1036900077|ref|XP_017153477.1| PREDICTED: chromosome transmission fidelity protein 18 homol
>gi|1036900139|ref|XP_017153481.1| PREDICTED: uncharacterized protein LOC108162985 [Drosophila
>gi|1036900158|ref|XP_017153483.1| PREDICTED: uncharacterized protein LOC108162986 [Drosophila
>gi|1036900177|ref|XP_017153484.1| PREDICTED: chymotrypsin inhibitor SCI-I [Drosophila miranda]
>gi|1036925164|ref|XP_017154879.1| PREDICTED: kunitz-type U1-aranetoxin-Av1a [Drosophila miran
>gi|1036925146|ref|XP_017154878.1| PREDICTED: kunitz-type protease inhibitor 3 [Drosophila mir
>gi|1036925192|ref|XP_017154881.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036925203|ref|XP_017154882.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036925231|ref|XP_017154883.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036925249|ref|XP_017154885.1| PREDICTED: kunitz-type serine protease inhibitor 2-like iso
>gi|1036925176|ref|XP_017154880.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036925267|ref|XP_017154886.1| PREDICTED: male accessory gland serine protease inhibitor-l
>gi|1036862762|ref|XP_017151406.1| PREDICTED: male accessory gland serine protease inhibitor [
>gi|1036860265|ref|XP_017151268.1| PREDICTED: chymotrypsin inhibitor SCI-II-like [Drosophila m
>gi|1036934438|ref|XP_017155386.1| PREDICTED: homeobox protein aristaless [Drosophila miranda]
>gi|1036884988|ref|XP_017152634.1| PREDICTED: cGMP-dependent protein kinase, isozyme 2 forms c
>gi|1036885007|ref|XP_017152636.1| PREDICTED: cGMP-dependent protein kinase, isozyme 2 forms c
>gi|1036885045|ref|XP_017152638.1| PREDICTED: PI-actitoxin-Aeq3a [Drosophila miranda]
>gi|1036885026|ref|XP_017152637.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1036923660|ref|XP_017154800.1| PREDICTED: protein odd-skipped [Drosophila miranda]
>gi|1036925463|ref|XP_017154897.1| PREDICTED: protein sister of odd and bowel [Drosophila mira
>gi|1036920889|ref|XP_017154643.1| PREDICTED: protein drumstick [Drosophila miranda]
>gi|1036920909|ref|XP_017154645.1| PREDICTED: protein drumstick [Drosophila miranda]
>gi|1036857726|ref|XP_017151140.1| PREDICTED: intraflagellar transport protein 57 homolog isof
>gi|1036857745|ref|XP_017151141.1| PREDICTED: intraflagellar transport protein 57 homolog isof
>gi|1036894870|ref|XP_017153185.1| PREDICTED: protein Spindly [Drosophila miranda]
>gi|1036894889|ref|XP_017153186.1| PREDICTED: uncharacterized protein LOC108162795 [Drosophila
>gi|1036894908|ref|XP_017153187.1| PREDICTED: centromere protein F [Drosophila miranda]
>gi|1036894927|ref|XP_017153188.1| PREDICTED: uncharacterized protein LOC108162797 [Drosophila
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>gi|1036894946|ref|XP_017153189.1| PREDICTED: coiled-coil domain-containing protein 47 [Drosoph
>gi|1036895571|ref|XP_017153225.1| PREDICTED: leucine--tRNA ligase, cytoplasmic [Drosophila mi
>gi|1036895590|ref|XP_017153226.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036895514|ref|XP_017153222.1| PREDICTED: protein timeless [Drosophila miranda]
>gi|1036895533|ref|XP_017153223.1| PREDICTED: protein timeless [Drosophila miranda]
>gi|1036895552|ref|XP_017153224.1| PREDICTED: protein timeless [Drosophila miranda]
>gi|1036933430|ref|XP_017155334.1| PREDICTED: uncharacterized protein LOC108164213 isoform X1
>gi|1036933450|ref|XP_017155335.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1036929385|ref|XP_017155115.1| PREDICTED: eukaryotic translation initiation factor 4E-bind
>gi|1036920928|ref|XP_017154646.1| PREDICTED: N-acetylgalactosaminyltransferase 4 [Drosophila r
>gi|1036920944|ref|XP_017154647.1| PREDICTED: N-acetylgalactosaminyltransferase 4-like [Drosoph
>gi|1036912081|ref|XP_017154130.1| PREDICTED: ATP-dependent DNA helicase PIF1 [Drosophila mirar
>gi|1036912099|ref|XP_017154131.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036911403|ref|XP_017154094.1| PREDICTED: uncharacterized protein LOC108163360 isoform X1
>gi|1036911421|ref|XP_017154095.1| PREDICTED: uncharacterized protein LOC108163360 isoform X2
>gi|1036911499|ref|XP_017154099.1| PREDICTED: structural maintenance of chromosomes protein 6-l
>gi|1036911461|ref|XP_017154097.1| PREDICTED: structural maintenance of chromosomes protein 6-l
>gi|1036911479|ref|XP_017154098.1| PREDICTED: structural maintenance of chromosomes protein 6-l
>gi|1036911521|ref|XP_017154100.1| PREDICTED: trypsin inhibitor-like [Drosophila miranda]
>gi|1036911441|ref|XP_017154096.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036884033|ref|XP_017152583.1| PREDICTED: BTB/POZ domain-containing protein 9 [Drosophila r
>gi|1036884147|ref|XP_017152589.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036884128|ref|XP_017152588.1| PREDICTED: probable small nuclear ribonucleoprotein E [Droso
>gi|1036884014|ref|XP_017152581.1| PREDICTED: uncharacterized protein LOC108162376 isoform X2
>gi|1036883978|ref|XP_017152579.1| PREDICTED: uncharacterized protein LOC108162376 isoform X1
>gi|1036883997|ref|XP_017152580.1| PREDICTED: uncharacterized protein LOC108162376 isoform X1
>gi|1036884109|ref|XP_017152587.1| PREDICTED: uncharacterized protein LOC108162381 [Drosophila
>gi|1036884090|ref|XP_017152586.1| PREDICTED: proline-rich extensin-like protein EPR1 [Drosoph
>gi|1036883941|ref|XP_017152577.1| PREDICTED: kinesin-like protein GA13060 [Drosophila miranda]
>gi|1036883960|ref|XP_017152578.1| PREDICTED: kinesin-like protein GA13060 [Drosophila miranda]
>gi|1036884071|ref|XP_017152585.1| PREDICTED: uncharacterized protein LOC108162379 [Drosophila
>gi|1036884052|ref|XP_017152584.1| PREDICTED: serine/threonine-protein phosphatase PP2A [Drosop
>gi|1036929140|ref|XP_017155102.1| PREDICTED: uncharacterized protein LOC108164034 [Drosophila
>gi|1036901913|ref|XP_017153581.1| PREDICTED: guanine nucleotide-binding protein subunit beta-l
>gi|1036901895|ref|XP_017153580.1| PREDICTED: protein phosphatase 1L [Drosophila miranda]
>gi|1036901929|ref|XP_017153582.1| PREDICTED: histidine triad nucleotide-binding protein 3 [Dro
>gi|1036932312|ref|XP_017155269.1| PREDICTED: serine proteases 1/2 isoform X2 [Drosophila mirar
>gi|1036932290|ref|XP_017155268.1| PREDICTED: serine proteases 1/2 isoform X1 [Drosophila mirar
>gi|1036854993|ref|XP_017150981.1| PREDICTED: uncharacterized protein LOC108161272 [Drosophila
>gi|1036918002|ref|XP_017154480.1| PREDICTED: F-box/LRR-repeat protein 15 isoform X2 [Drosophil
>gi|1036917985|ref|XP_017154479.1| PREDICTED: F-box/LRR-repeat protein 15 isoform X1 [Drosophil
>gi|1036917967|ref|XP_017154478.1| PREDICTED: geranylgeranyl transferase type-1 subunit beta [[
>gi|1036920476|ref|XP_017154617.1| PREDICTED: probable 39S ribosomal protein L24, mitochondrial
>gi|1036920456|ref|XP_017154616.1| PREDICTED: uncharacterized protein LOC108163694 isoform X2
>gi|1036920438|ref|XP_017154615.1| PREDICTED: uncharacterized protein LOC108163694 isoform X1
>gi|1036920420|ref|XP_017154614.1| PREDICTED: P protein [Drosophila miranda]
>gi|1036912265|ref|XP_017154141.1| PREDICTED: P protein isoform X1 [Drosophila miranda]
>gi|1036912282|ref|XP_017154142.1| PREDICTED: P protein isoform X2 [Drosophila miranda]
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>gi|1036912316|ref|XP_017154144.1| PREDICTED: uncharacterized protein LOC108163395 [Drosophila
>gi|1036912300|ref|XP_017154143.1| PREDICTED: P protein isoform X3 [Drosophila miranda]
>gi|1036857978|ref|XP_017151154.1| PREDICTED: uncharacterized protein LOC108161413 [Drosophila
>gi|1036857997|ref|XP_017151155.1| PREDICTED: uncharacterized protein LOC108161413 [Drosophila
>gi|1036858016|ref|XP_017151156.1| PREDICTED: uncharacterized protein LOC108161413 [Drosophila
>gi|1036903065|ref|XP_017153645.1| PREDICTED: sialin [Drosophila miranda]
>gi|1036903085|ref|XP_017153646.1| PREDICTED: sialin [Drosophila miranda]
>gi|1036903103|ref|XP_017153647.1| PREDICTED: uncharacterized protein LOC108163078 [Drosophila
>gi|1036874068|ref|XP_017152038.1| PREDICTED: serine/threonine-protein kinase RIO3 [Drosophila
>gi|1036874145|ref|XP_017152041.1| PREDICTED: chorion transcription factor Cf2 isoform X1 [Dros
>gi|1036874164|ref|XP_017152042.1| PREDICTED: chorion transcription factor Cf2 isoform X2 [Dros
>gi|1036874182|ref|XP_017152043.1| PREDICTED: chorion transcription factor Cf2 isoform X3 [Dros
>gi|1036874200|ref|XP_017152044.1| PREDICTED: chorion transcription factor Cf2 isoform X3 [Dros
>gi|1036874219|ref|XP_017152046.1| PREDICTED: chorion transcription factor Cf2 isoform X3 [Dros
>gi|1036855012|ref|XP_017150982.1| PREDICTED: probable G-protein coupled receptor 158 [Drosophi
>gi|1036855033|ref|XP_017150983.1| PREDICTED: uncharacterized protein LOC108161275 [Drosophila
>gi|1036874087|ref|XP_017152039.1| PREDICTED: vacuolar fusion protein MON1 homolog A [Drosophil
```

In [44]: *## Redirección de la salida a un archivo*
         grep ">" protein.fa > resultados.txt
         ll

```
total 25M
-rwxrwxr-x 1 4.5M 10-01 21:22 genoma.fna
-rw-rw-r-- 1    0 10-01 21:06 mundo.txt
-rw-rw-r-- 1  18M 10-01 21:32 protein.fa
-rw-rw-r-- 1 2.4M 10-01 21:40 resultados.txt
```

In [41]: cat resultados.txt

```
>gi|1036701783|ref|XP_017155988.1| PREDICTED: LOW QUALITY PROTEIN: ATP-dependent RNA helicase
>gi|1036701631|ref|XP_017155883.1| PREDICTED: LOW QUALITY PROTEIN: probable cytochrome P450 31
>gi|1036701733|ref|XP_017155947.1| PREDICTED: uncharacterized protein LOC108164616 [Drosophila
>gi|1036701800|ref|XP_017155998.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036701338|ref|XP_017155122.1| PREDICTED: serine/threonine-protein kinase par-1 [Drosophila
>gi|1036701601|ref|XP_017155861.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701488|ref|XP_017155806.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701509|ref|XP_017155814.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701526|ref|XP_017155824.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701545|ref|XP_017155832.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701562|ref|XP_017155842.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701582|ref|XP_017155851.1| PREDICTED: G-protein coupled receptor moody [Drosophila mira
>gi|1036701768|ref|XP_017155968.1| PREDICTED: E3 ubiquitin-protein ligase RNF128 [Drosophila mi
>gi|1036701422|ref|XP_017155494.1| PREDICTED: alpha-actinin, sarcomeric isoform X4 [Drosophila
>gi|1036701616|ref|XP_017155871.1| PREDICTED: protein ultraspiracle [Drosophila miranda]
>gi|1036701355|ref|XP_017155231.1| PREDICTED: alpha-actinin, sarcomeric isoform X1 [Drosophila
>gi|1036701374|ref|XP_017155306.1| PREDICTED: alpha-actinin, sarcomeric isoform X1 [Drosophila
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>gi|1036701390|ref|XP_017155378.1| PREDICTED: alpha-actinin, sarcomeric isoform X2 [Drosophila
>gi|1036701406|ref|XP_017155453.1| PREDICTED: alpha-actinin, sarcomeric isoform X3 [Drosophila
>gi|1036701287|ref|XP_017154817.1| PREDICTED: uncharacterized protein LOC108163645 isoform X4
>gi|1036701272|ref|XP_017154742.1| PREDICTED: uncharacterized protein LOC108163645 isoform X3
>gi|1036701257|ref|XP_017154665.1| PREDICTED: uncharacterized protein LOC108163645 isoform X2
>gi|1036701243|ref|XP_017154584.1| PREDICTED: uncharacterized protein LOC108163645 isoform X1
>gi|1036701701|ref|XP_017155930.1| PREDICTED: putative GPI-anchor transamidase [Drosophila mira
>gi|1036701717|ref|XP_017155937.1| PREDICTED: putative GPI-anchor transamidase [Drosophila mira
>gi|1036698287|ref|XP_017139785.1| PREDICTED: AF4/FMR2 family member 4-like [Drosophila miranda
>gi|1036701323|ref|XP_017155006.1| PREDICTED: hormone receptor 4-like isoform X2 [Drosophila mi
>gi|1036701305|ref|XP_017154930.1| PREDICTED: hormone receptor 4-like isoform X1 [Drosophila mi
>gi|1036701752|ref|XP_017155958.1| PREDICTED: protein KTI12 homolog [Drosophila miranda]
>gi|1036701647|ref|XP_017155896.1| PREDICTED: protein SMG9 [Drosophila miranda]
>gi|1036701680|ref|XP_017155916.1| PREDICTED: carnosine N-methyltransferase isoform X2 [Drosoph
>gi|1036701663|ref|XP_017155907.1| PREDICTED: carnosine N-methyltransferase isoform X1 [Drosoph
>gi|1036701439|ref|XP_017155626.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila mirar
>gi|1036701454|ref|XP_017155715.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila mirar
>gi|1036701471|ref|XP_017155795.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila mirar
>gi|1036711564|ref|XP_017139547.1| PREDICTED: dehydrodolichyl diphosphate synthase complex subu
>gi|1036711597|ref|XP_017139567.1| PREDICTED: 40S ribosomal protein S14 [Drosophila miranda]
>gi|1036711581|ref|XP_017139558.1| PREDICTED: 40S ribosomal protein S14-like [Drosophila mirand
>gi|1036711500|ref|XP_017139511.1| PREDICTED: E3 ubiquitin-protein ligase UBR3 [Drosophila mira
>gi|1036711515|ref|XP_017139518.1| PREDICTED: E3 ubiquitin-protein ligase UBR3 [Drosophila mira
>gi|1036711534|ref|XP_017139529.1| PREDICTED: uncharacterized protein LOC108153940 isoform X1
>gi|1036711550|ref|XP_017139537.1| PREDICTED: uncharacterized protein LOC108153940 isoform X2
>gi|1036751956|ref|XP_017156406.1| PREDICTED: DM7 family protein GG17591 [Drosophila miranda]
>gi|1036729081|ref|XP_017149347.1| PREDICTED: neuropathy target esterase sws isoform X4 [Drosop
>gi|1036729050|ref|XP_017149329.1| PREDICTED: neuropathy target esterase sws isoform X2 [Drosop
>gi|1036729021|ref|XP_017149318.1| PREDICTED: neuropathy target esterase sws isoform X1 [Drosop
>gi|1036729063|ref|XP_017149338.1| PREDICTED: neuropathy target esterase sws isoform X3 [Drosop
>gi|1036737636|ref|XP_017154022.1| PREDICTED: protein singed [Drosophila miranda]
>gi|1036702955|ref|XP_017156697.1| PREDICTED: protein arginine N-methyltransferase 1-like [Dros
>gi|1036707540|ref|XP_017137034.1| PREDICTED: E3 ubiquitin ligase complex SCF subunit sconC-lik
>gi|1036702891|ref|XP_017156653.1| PREDICTED: uncharacterized protein LOC108165104 [Drosophila
>gi|1036756219|ref|XP_017156657.1| PREDICTED: tyramine beta-hydroxylase [Drosophila miranda]
>gi|1036756236|ref|XP_017156658.1| PREDICTED: tyramine beta-hydroxylase [Drosophila miranda]
>gi|1036756255|ref|XP_017156659.1| PREDICTED: tyramine beta-hydroxylase [Drosophila miranda]
>gi|1036756728|ref|XP_017156687.1| PREDICTED: uncharacterized protein LOC108165131 isoform X3
>gi|1036756709|ref|XP_017156686.1| PREDICTED: uncharacterized protein LOC108165131 isoform X2
>gi|1036756690|ref|XP_017156685.1| PREDICTED: uncharacterized protein LOC108165131 isoform X1
>gi|1036709552|ref|XP_017138295.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709468|ref|XP_017138254.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709484|ref|XP_017138263.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709501|ref|XP_017138272.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709518|ref|XP_017138278.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709535|ref|XP_017138286.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709568|ref|XP_017138303.1| PREDICTED: phosphorylase b kinase gamma catalytic chain, ske
>gi|1036709612|ref|XP_017138327.1| PREDICTED: uncharacterized protein LOC108153062 [Drosophila
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>gi|1036709344|ref|XP_017138193.1| PREDICTED: alpha-(1,6)-fucosyltransferase [Drosophila miran
>gi|1036709365|ref|XP_017138200.1| PREDICTED: alpha-(1,6)-fucosyltransferase [Drosophila miran
>gi|1036709381|ref|XP_017138209.1| PREDICTED: alpha-(1,6)-fucosyltransferase [Drosophila miran
>gi|1036709430|ref|XP_017138234.1| PREDICTED: uncharacterized protein LOC108153001 [Drosophila
>gi|1036709449|ref|XP_017138244.1| PREDICTED: uncharacterized protein LOC108153001 [Drosophila
>gi|1036709597|ref|XP_017138315.1| PREDICTED: thioredoxin domain-containing protein 5 [Drosophi
>gi|1036709398|ref|XP_017138222.1| PREDICTED: BTB/POZ domain-containing protein 17 [Drosophila
>gi|1036752027|ref|XP_017156411.1| PREDICTED: putative transporter svop-1 [Drosophila miranda]
>gi|1036752042|ref|XP_017156412.1| PREDICTED: uncharacterized protein LOC108164934 [Drosophila
>gi|1036698181|ref|XP_017138639.1| PREDICTED: uncharacterized protein LOC108150747 [Drosophila
>gi|1036752060|ref|XP_017156413.1| PREDICTED: uncharacterized protein LOC108164935 [Drosophila
>gi|1036698196|ref|XP_017150822.1| PREDICTED: uncharacterized protein LOC108158651 [Drosophila
>gi|1036756638|ref|XP_017156682.1| PREDICTED: uncharacterized protein LOC108165128 [Drosophila
>gi|1036756653|ref|XP_017156683.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036706619|ref|XP_017136424.1| PREDICTED: uncharacterized protein LOC108151975 [Drosophila
>gi|1036755818|ref|XP_017156635.1| PREDICTED: uncharacterized protein LOC108165094 isoform X1
>gi|1036755837|ref|XP_017156636.1| PREDICTED: uncharacterized protein LOC108165094 isoform X2
>gi|1036755645|ref|XP_017156624.1| PREDICTED: insecticyanin-A [Drosophila miranda]
>gi|1036729511|ref|XP_017149608.1| PREDICTED: protein wings apart-like [Drosophila miranda]
>gi|1036698304|ref|XP_017140963.1| PREDICTED: uncharacterized protein LOC108154713 [Drosophila
>gi|1036755698|ref|XP_017156628.1| PREDICTED: uncharacterized protein LOC108165087 [Drosophila
>gi|1036755716|ref|XP_017156629.1| PREDICTED: uncharacterized protein LOC108165088 [Drosophila
>gi|1036730248|ref|XP_017150076.1| PREDICTED: LOW QUALITY PROTEIN: histone H2B-like [Drosophila
>gi|1036730189|ref|XP_017150041.1| PREDICTED: histone H1-like isoform X1 [Drosophila miranda]
>gi|1036730208|ref|XP_017150051.1| PREDICTED: histone H1-like isoform X2 [Drosophila miranda]
>gi|1036730229|ref|XP_017150062.1| PREDICTED: protein lethal(2)denticleless-like [Drosophila mi
>gi|1036729548|ref|XP_017149634.1| PREDICTED: centrosomal protein of 162 kDa isoform X1 [Drosop
>gi|1036729567|ref|XP_017149643.1| PREDICTED: centrosomal protein of 162 kDa isoform X2 [Drosop
>gi|1036729580|ref|XP_017149655.1| PREDICTED: carboxypeptidase B isoform X1 [Drosophila miranda
>gi|1036729600|ref|XP_017149664.1| PREDICTED: carboxypeptidase B isoform X2 [Drosophila miranda
>gi|1036698208|ref|XP_017156463.1| PREDICTED: zinc carboxypeptidase A 1 [Drosophila miranda]
>gi|1036729529|ref|XP_017149621.1| PREDICTED: histone acetyltransferase KAT6A [Drosophila miran
>gi|1036755785|ref|XP_017156633.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036703982|ref|XP_017134622.1| PREDICTED: proteasome subunit beta type-7 [Drosophila miran
>gi|1036703732|ref|XP_017157117.1| PREDICTED: dopamine N-acetyltransferase [Drosophila miranda]
>gi|1036747149|ref|XP_017156152.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036704091|ref|XP_017134680.1| PREDICTED: bax inhibitor 1-like [Drosophila miranda]
>gi|1036698319|ref|XP_017142208.1| PREDICTED: uncharacterized protein LOC108155475 [Drosophila
>gi|1036741943|ref|XP_017155857.1| PREDICTED: glycosylphosphatidylinositol anchor attachment 1
>gi|1036713747|ref|XP_017140860.1| PREDICTED: uncharacterized protein LOC108154906 isoform X2
>gi|1036713732|ref|XP_017140851.1| PREDICTED: uncharacterized protein LOC108154906 isoform X1
>gi|1036713793|ref|XP_017140898.1| PREDICTED: G2/M phase-specific E3 ubiquitin-protein ligase
>gi|1036713764|ref|XP_017140871.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 30 homolog
>gi|1036713827|ref|XP_017140923.1| PREDICTED: LOW QUALITY PROTEIN: T-complex protein 11-like pr
>gi|1036713779|ref|XP_017140883.1| PREDICTED: DDB1- and CUL4-associated factor 10 homolog [Dros
>gi|1036713810|ref|XP_017140913.1| PREDICTED: patatin-like phospholipase domain-containing prot
>gi|1036741536|ref|XP_017155836.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036741517|ref|XP_017155835.1| PREDICTED: lipid storage droplets surface-binding protein 2
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>gi|1036741577|ref|XP_017155838.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036741558|ref|XP_017155837.1| PREDICTED: lipid storage droplets surface-binding protein 2
>gi|1036706037|ref|XP_017136019.1| PREDICTED: tubulin-specific chaperone A-like [Drosophila mi
>gi|1036739823|ref|XP_017155212.1| PREDICTED: microsomal glutathione S-transferase 1 [Drosophil
>gi|1036739857|ref|XP_017155232.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X
>gi|1036739839|ref|XP_017155224.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X
>gi|1036739875|ref|XP_017155242.1| PREDICTED: cytochrome c oxidase copper chaperone [Drosophila
>gi|1036739893|ref|XP_017155248.1| PREDICTED: cytochrome c oxidase copper chaperone [Drosophila
>gi|1036739805|ref|XP_017155202.1| PREDICTED: translation machinery-associated protein 16 homol
>gi|1036739787|ref|XP_017155190.1| PREDICTED: LOW QUALITY PROTEIN: lipoamide acyltransferase co
>gi|1036738684|ref|XP_017154541.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738702|ref|XP_017154547.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738720|ref|XP_017154556.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738740|ref|XP_017154564.1| PREDICTED: MAP kinase-activating death domain protein isofor
>gi|1036738758|ref|XP_017154576.1| PREDICTED: tubulin-specific chaperone A [Drosophila miranda]
>gi|1036752119|ref|XP_017156416.1| PREDICTED: probable cytochrome P450 4s3 [Drosophila miranda]
>gi|1036740350|ref|XP_017155473.1| PREDICTED: uncharacterized protein LOC108164335 isoform X1
>gi|1036740366|ref|XP_017155474.1| PREDICTED: uncharacterized protein LOC108164335 isoform X2
>gi|1036740384|ref|XP_017155475.1| PREDICTED: probable 60S ribosomal protein L37-A [Drosophila
>gi|1036743313|ref|XP_017155936.1| PREDICTED: uncharacterized protein LOC108164608 [Drosophila
>gi|1036743295|ref|XP_017155935.1| PREDICTED: uncharacterized protein LOC108164607 [Drosophila
>gi|1036705018|ref|XP_017135315.1| PREDICTED: uncharacterized protein LOC108151289 [Drosophila
>gi|1036753611|ref|XP_017156508.1| PREDICTED: uncharacterized protein LOC108164994 isoform X1
>gi|1036753631|ref|XP_017156510.1| PREDICTED: uncharacterized protein LOC108164994 isoform X2
>gi|1036748586|ref|XP_017156220.1| PREDICTED: uncharacterized protein LOC108164812 [Drosophila
>gi|1036698334|ref|XP_017143452.1| PREDICTED: uncharacterized protein LOC108156218 [Drosophila
>gi|1036751287|ref|XP_017156370.1| PREDICTED: uncharacterized protein LOC108164904 [Drosophila
>gi|1036751308|ref|XP_017156371.1| PREDICTED: uncharacterized protein LOC108164904 [Drosophila
>gi|1036751329|ref|XP_017156372.1| PREDICTED: uncharacterized protein LOC108164905 [Drosophila
>gi|1036705899|ref|XP_017135911.1| PREDICTED: uncharacterized protein LOC108151676 [Drosophila
>gi|1036739354|ref|XP_017154947.1| PREDICTED: uncharacterized protein LOC108163921 [Drosophila
>gi|1036738971|ref|XP_017154717.1| PREDICTED: uncharacterized protein LOC108163755 [Drosophila
>gi|1036738989|ref|XP_017154725.1| PREDICTED: uncharacterized protein LOC108163755 [Drosophila
>gi|1036739007|ref|XP_017154733.1| PREDICTED: uncharacterized protein LOC108163755 [Drosophila
>gi|1036715019|ref|XP_017141618.1| PREDICTED: lymphokine-activated killer T-cell-originated pro
>gi|1036715000|ref|XP_017141607.1| PREDICTED: partitioning defective protein 6 [Drosophila mira
>gi|1036715036|ref|XP_017141631.1| PREDICTED: ubiquitin-conjugating enzyme E2 S isoform X1 [Dro
>gi|1036715053|ref|XP_017141639.1| PREDICTED: ubiquitin-conjugating enzyme E2 S isoform X2 [Dro
>gi|1036715091|ref|XP_017141660.1| PREDICTED: uncharacterized protein LOC108155384 isoform X2
>gi|1036715072|ref|XP_017141652.1| PREDICTED: uncharacterized protein LOC108155384 isoform X1
>gi|1036715109|ref|XP_017141670.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036756166|ref|XP_017156654.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036709204|ref|XP_017138102.1| PREDICTED: spectrin beta chain isoform X2 [Drosophila mirand
>gi|1036709154|ref|XP_017138077.1| PREDICTED: spectrin beta chain isoform X1 [Drosophila mirand
>gi|1036709171|ref|XP_017138085.1| PREDICTED: spectrin beta chain isoform X1 [Drosophila mirand
>gi|1036709188|ref|XP_017138093.1| PREDICTED: spectrin beta chain isoform X1 [Drosophila mirand
>gi|1036709219|ref|XP_017138111.1| PREDICTED: spectrin beta chain isoform X3 [Drosophila mirand
>gi|1036709232|ref|XP_017138123.1| PREDICTED: integrator complex subunit 2 [Drosophila miranda]
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>gi|1036709251|ref|XP_017138135.1| PREDICTED: rho GTPase-activating protein 190 isoform X1 [Dro
>gi|1036709267|ref|XP_017138143.1| PREDICTED: rho GTPase-activating protein 190 isoform X1 [Dro
>gi|1036709281|ref|XP_017138151.1| PREDICTED: rho GTPase-activating protein 190 isoform X2 [Dro
>gi|1036709296|ref|XP_017138161.1| PREDICTED: rho GTPase-activating protein 190 isoform X3 [Dro
>gi|1036709312|ref|XP_017138170.1| PREDICTED: rho GTPase-activating protein 190 isoform X4 [Dro
>gi|1036709327|ref|XP_017138181.1| PREDICTED: rho GTPase-activating protein 190 isoform X5 [Dro
>gi|1036742108|ref|XP_017155865.1| PREDICTED: probable ATP-dependent RNA helicase DDX10 [Drosop
>gi|1036736749|ref|XP_017153521.1| PREDICTED: calpain-D isoform X1 [Drosophila miranda]
>gi|1036736770|ref|XP_017153530.1| PREDICTED: calpain-D isoform X2 [Drosophila miranda]
>gi|1036736789|ref|XP_017153539.1| PREDICTED: calpain-D isoform X3 [Drosophila miranda]
>gi|1036736807|ref|XP_017153548.1| PREDICTED: calpain-D isoform X4 [Drosophila miranda]
>gi|1036738006|ref|XP_017154223.1| PREDICTED: septin-1 isoform X2 [Drosophila miranda]
>gi|1036737988|ref|XP_017154215.1| PREDICTED: septin-1 isoform X1 [Drosophila miranda]
>gi|1036718395|ref|XP_017143507.1| PREDICTED: translation factor waclaw, mitochondrial [Drosop
>gi|1036718226|ref|XP_017143423.1| PREDICTED: uncharacterized protein LOC108156467 [Drosophila
>gi|1036718246|ref|XP_017143430.1| PREDICTED: uncharacterized protein LOC108156467 [Drosophila
>gi|1036718264|ref|XP_017143439.1| PREDICTED: uncharacterized protein LOC108156467 [Drosophila
>gi|1036718303|ref|XP_017143462.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X2
>gi|1036718377|ref|XP_017143496.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X4
>gi|1036718283|ref|XP_017143454.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X1
>gi|1036718322|ref|XP_017143472.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X1
>gi|1036718343|ref|XP_017143479.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X1
>gi|1036718360|ref|XP_017143487.1| PREDICTED: proline dehydrogenase 1, mitochondrial isoform X3
>gi|1036718410|ref|XP_017143518.1| PREDICTED: probable ATP-dependent RNA helicase DDX56 [Drosop
>gi|1036751607|ref|XP_017156387.1| PREDICTED: uncharacterized protein LOC108164912 [Drosophila
>gi|1036742446|ref|XP_017155886.1| PREDICTED: mpv17-like protein [Drosophila miranda]
>gi|1036742427|ref|XP_017155885.1| PREDICTED: protein misato [Drosophila miranda]
>gi|1036698349|ref|XP_017144690.1| PREDICTED: myosin heavy chain IB-like [Drosophila miranda]
>gi|1036706022|ref|XP_017136003.1| PREDICTED: proline-rich receptor-like protein kinase PERK9
>gi|1036753255|ref|XP_017156488.1| PREDICTED: neurotrimin isoform X4 [Drosophila miranda]
>gi|1036753219|ref|XP_017156486.1| PREDICTED: neurotrimin isoform X2 [Drosophila miranda]
>gi|1036753238|ref|XP_017156487.1| PREDICTED: neuronal growth regulator 1 isoform X3 [Drosophi
>gi|1036753204|ref|XP_017156485.1| PREDICTED: neuronal growth regulator 1 isoform X1 [Drosophi
>gi|1036706318|ref|XP_017136222.1| PREDICTED: uncharacterized protein LOC108151845 [Drosophila
>gi|1036735067|ref|XP_017152556.1| PREDICTED: uncharacterized protein LOC108162358 [Drosophila
>gi|1036728354|ref|XP_017148921.1| PREDICTED: neutral alpha-glucosidase AB [Drosophila miranda]
>gi|1036728373|ref|XP_017148929.1| PREDICTED: neutral alpha-glucosidase AB [Drosophila miranda]
>gi|1036728412|ref|XP_017148951.1| PREDICTED: tektin-3-like isoform X2 [Drosophila miranda]
>gi|1036728391|ref|XP_017148943.1| PREDICTED: tektin-3-like isoform X1 [Drosophila miranda]
>gi|1036728433|ref|XP_017148962.1| PREDICTED: NFU1 iron-sulfur cluster scaffold homolog, mitoch
>gi|1036742771|ref|XP_017155904.1| PREDICTED: uncharacterized protein LOC108164590 [Drosophila
>gi|1036725812|ref|XP_017147391.1| PREDICTED: solute carrier family 35 member E1 homolog [Dros
>gi|1036725850|ref|XP_017147422.1| PREDICTED: uncharacterized protein LOC108159019 [Drosophila
>gi|1036725831|ref|XP_017147407.1| PREDICTED: DDB1- and CUL4-associated factor 7 [Drosophila m
>gi|1036725012|ref|XP_017146926.1| PREDICTED: cell wall protein AWA1 isoform X4 [Drosophila mi
>gi|1036725030|ref|XP_017146937.1| PREDICTED: cell wall protein AWA1 isoform X4 [Drosophila mi
>gi|1036725051|ref|XP_017146946.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 2 isoform
>gi|1036724951|ref|XP_017146896.1| PREDICTED: myb-like protein A isoform X1 [Drosophila miranda
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>gi|1036724993|ref|XP_017146916.1| PREDICTED: mucin-12 isoform X3 [Drosophila miranda]
>gi|1036724972|ref|XP_017146906.1| PREDICTED: myb-like protein A isoform X2 [Drosophila miranda
>gi|1036723743|ref|XP_017146122.1| PREDICTED: uncharacterized protein C18orf19 homolog A [Droso
>gi|1036723724|ref|XP_017146111.1| PREDICTED: tRNA methyltransferase 10 homolog A [Drosophila r
>gi|1036723666|ref|XP_017146082.1| PREDICTED: uncharacterized protein LOC108158350 [Drosophila
>gi|1036723685|ref|XP_017146092.1| PREDICTED: uncharacterized protein LOC108158350 [Drosophila
>gi|1036723704|ref|XP_017146101.1| PREDICTED: uncharacterized protein LOC108158350 [Drosophila
>gi|1036722128|ref|XP_017145358.1| PREDICTED: uncharacterized protein LOC108157696 [Drosophila
>gi|1036721934|ref|XP_017145274.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721953|ref|XP_017145283.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722053|ref|XP_017145325.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721970|ref|XP_017145291.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722071|ref|XP_017145333.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721919|ref|XP_017145266.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
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>gi|1036721864|ref|XP_017145241.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722090|ref|XP_017145340.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721883|ref|XP_017145249.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036722111|ref|XP_017145346.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721847|ref|XP_017145232.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721902|ref|XP_017145258.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036721989|ref|XP_017145300.1| PREDICTED: inositol hexakisphosphate and diphosphoinositol-p
>gi|1036727249|ref|XP_017148291.1| PREDICTED: uncharacterized protein LOC108159474 isoform X1
>gi|1036727232|ref|XP_017148278.1| PREDICTED: ribosomal protein S6 kinase 2 beta [Drosophila mi
>gi|1036727279|ref|XP_017148311.1| PREDICTED: protein phosphatase 1H [Drosophila miranda]
>gi|1036727262|ref|XP_017148299.1| PREDICTED: formin BNI1 isoform X2 [Drosophila miranda]
>gi|1036704459|ref|XP_017134908.1| PREDICTED: clusterin-associated protein 1 [Drosophila mirand
>gi|1036736279|ref|XP_017153243.1| PREDICTED: adenylate cyclase type 9 isoform X1 [Drosophila r
>gi|1036736297|ref|XP_017153251.1| PREDICTED: adenylate cyclase type 9 isoform X2 [Drosophila r
>gi|1036736315|ref|XP_017153260.1| PREDICTED: adenylate cyclase type 9 isoform X3 [Drosophila r
>gi|1036715889|ref|XP_017142115.1| PREDICTED: thyroid receptor-interacting protein 11 [Drosophi
>gi|1036715959|ref|XP_017142159.1| PREDICTED: F-box-like/WD repeat-containing protein TBL1XR1
>gi|1036715941|ref|XP_017142147.1| PREDICTED: arginine/serine-rich coiled-coil protein 2 isofor
>gi|1036715923|ref|XP_017142139.1| PREDICTED: arginine/serine-rich coiled-coil protein 2 isofor
>gi|1036715907|ref|XP_017142126.1| PREDICTED: uncharacterized protein LOC108155670 [Drosophila
>gi|1036742178|ref|XP_017155869.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036742197|ref|XP_017155870.1| PREDICTED: UDP-N-acetylglucosamine transferase subunit ALG14
>gi|1036724525|ref|XP_017146626.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036724579|ref|XP_017146667.1| PREDICTED: uncharacterized protein LOC108158667 [Drosophila
>gi|1036724560|ref|XP_017146653.1| PREDICTED: glycolipid transfer protein [Drosophila miranda]
>gi|1036698364|ref|XP_017147071.1| PREDICTED: cell surface glycoprotein 1-like [Drosophila mira
>gi|1036724541|ref|XP_017146641.1| PREDICTED: uncharacterized protein LOC108158654 [Drosophila
>gi|1036749432|ref|XP_017156265.1| PREDICTED: putative inositol monophosphatase 3 [Drosophila r
>gi|1036733728|ref|XP_017151782.1| PREDICTED: histone deacetylase 4 isoform X1 [Drosophila mira
>gi|1036733765|ref|XP_017151798.1| PREDICTED: histone deacetylase 5 isoform X3 [Drosophila mira
>gi|1036733747|ref|XP_017151790.1| PREDICTED: histone deacetylase 4 isoform X2 [Drosophila mira
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>gi|1036733782|ref|XP_017151808.1| PREDICTED: histone deacetylase 4 isoform X4 [Drosophila mira
>gi|1036713039|ref|XP_017140388.1| PREDICTED: probable peroxisomal acyl-coenzyme A oxidase 1 [I
>gi|1036713072|ref|XP_017140410.1| PREDICTED: leucine-rich repeat-containing protein 57 [Drosop
>gi|1036713055|ref|XP_017140402.1| PREDICTED: proteasome activator complex subunit 3 [Drosophil
>gi|1036713006|ref|XP_017140366.1| PREDICTED: integrin alpha-PS1 isoform X2 [Drosophila miranda
>gi|1036712989|ref|XP_017140358.1| PREDICTED: integrin alpha-PS1 isoform X1 [Drosophila miranda
>gi|1036713088|ref|XP_017140418.1| PREDICTED: uncharacterized protein LOC108154609 [Drosophila
>gi|1036713023|ref|XP_017140377.1| PREDICTED: vesicle-fusing ATPase 1 [Drosophila miranda]
>gi|1036753070|ref|XP_017156476.1| PREDICTED: protein tilB [Drosophila miranda]
>gi|1036735331|ref|XP_017152710.1| PREDICTED: histone-lysine N-methyltransferase setd3 [Drosop
>gi|1036735282|ref|XP_017152687.1| PREDICTED: uncharacterized protein LOC108162452 isoform X1
>gi|1036735301|ref|XP_017152695.1| PREDICTED: uncharacterized protein LOC108162452 isoform X2
>gi|1036702908|ref|XP_017156665.1| PREDICTED: uncharacterized protein LOC108165114 [Drosophila
>gi|1036724066|ref|XP_017146325.1| PREDICTED: LOW QUALITY PROTEIN: serine-rich adhesin for pla
>gi|1036754168|ref|XP_017156542.1| PREDICTED: uncharacterized protein LOC108165015 [Drosophila
>gi|1036714661|ref|XP_017141388.1| PREDICTED: uncharacterized protein LOC108155212 [Drosophila
>gi|1036714605|ref|XP_017141352.1| PREDICTED: uncharacterized protein LOC108155188 [Drosophila
>gi|1036714679|ref|XP_017141400.1| PREDICTED: 28 kDa heat- and acid-stable phosphoprotein [Dros
>gi|1036714642|ref|XP_017141375.1| PREDICTED: ribosomal RNA small subunit methyltransferase NEI
>gi|1036714698|ref|XP_017141412.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036714510|ref|XP_017141305.1| PREDICTED: probable cytochrome P450 4d14 [Drosophila miranda
>gi|1036714473|ref|XP_017141284.1| PREDICTED: cytochrome P450 4d2 [Drosophila miranda]
>gi|1036714492|ref|XP_017141293.1| PREDICTED: cytochrome P450 4d2 [Drosophila miranda]
>gi|1036714567|ref|XP_017141332.1| PREDICTED: cytochrome P450 4ae1 isoform X3 [Drosophila miran
>gi|1036714586|ref|XP_017141340.1| PREDICTED: cytochrome P450 4ae1 isoform X4 [Drosophila miran
>gi|1036714529|ref|XP_017141317.1| PREDICTED: cytochrome P450 4ae1 isoform X1 [Drosophila miran
>gi|1036714548|ref|XP_017141325.1| PREDICTED: cytochrome P450 4ae1 isoform X2 [Drosophila miran
>gi|1036714624|ref|XP_017141363.1| PREDICTED: protein prune homolog [Drosophila miranda]
>gi|1036714454|ref|XP_017141272.1| PREDICTED: 60S ribosomal export protein NMD3 [Drosophila mir
>gi|1036714718|ref|XP_017141421.1| PREDICTED: uncharacterized protein LOC108155239 [Drosophila
>gi|1036705377|ref|XP_017135563.1| PREDICTED: keratin-associated protein 19-2 [Drosophila miran
>gi|1036702307|ref|XP_017156278.1| PREDICTED: uncharacterized protein LOC108164857 [Drosophila
>gi|1036736396|ref|XP_017153315.1| PREDICTED: monocarboxylate transporter 12 [Drosophila mirand
>gi|1036736415|ref|XP_017153326.1| PREDICTED: fatty acyl-CoA reductase 1 [Drosophila miranda]
>gi|1036727562|ref|XP_017148473.1| PREDICTED: protein msta isoform X1 [Drosophila miranda]
>gi|1036727581|ref|XP_017148482.1| PREDICTED: protein msta isoform X2 [Drosophila miranda]
>gi|1036727524|ref|XP_017148448.1| PREDICTED: vinculin [Drosophila miranda]
>gi|1036727541|ref|XP_017148459.1| PREDICTED: vinculin [Drosophila miranda]
>gi|1036727598|ref|XP_017148494.1| PREDICTED: histidine-rich glycoprotein isoform X1 [Drosophil
>gi|1036727616|ref|XP_017148503.1| PREDICTED: histidine-rich glycoprotein isoform X1 [Drosophil
>gi|1036727634|ref|XP_017148513.1| PREDICTED: histidine-rich glycoprotein isoform X2 [Drosophil
>gi|1036715126|ref|XP_017141684.1| PREDICTED: protein pecanex [Drosophila miranda]
>gi|1036715144|ref|XP_017141695.1| PREDICTED: probable ATP-dependent RNA helicase kurz [Drosop
>gi|1036744253|ref|XP_017155981.1| PREDICTED: DNA-binding protein K10 [Drosophila miranda]
>gi|1036705611|ref|XP_017135701.1| PREDICTED: odorant receptor 2a [Drosophila miranda]
>gi|1036743910|ref|XP_017155964.1| PREDICTED: protein crooked neck [Drosophila miranda]
>gi|1036698381|ref|XP_017148371.1| PREDICTED: uncharacterized protein LOC108159290 [Drosophila
>gi|1036729414|ref|XP_017149549.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
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>gi|1036729322|ref|XP_017149494.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036729341|ref|XP_017149503.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036729357|ref|XP_017149511.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036729395|ref|XP_017149535.1| PREDICTED: protein prenyltransferase alpha subunit repeat-cc
>gi|1036729376|ref|XP_017149523.1| PREDICTED: U3 small nucleolar RNA-associated protein 15 homc
>gi|1036732027|ref|XP_017150947.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X2 [Di
>gi|1036731916|ref|XP_017150904.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Di
>gi|1036731935|ref|XP_017150911.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Di
>gi|1036731954|ref|XP_017150918.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Di
>gi|1036731971|ref|XP_017150926.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Di
>gi|1036731989|ref|XP_017150933.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Di
>gi|1036732008|ref|XP_017150940.1| PREDICTED: ubiquitin-conjugating enzyme E2 Q2 isoform X1 [Di
>gi|1036733364|ref|XP_017151590.1| PREDICTED: hypoxia up-regulated protein 1 [Drosophila mirand
>gi|1036733383|ref|XP_017151599.1| PREDICTED: hypoxia up-regulated protein 1 [Drosophila mirand
>gi|1036733402|ref|XP_017151612.1| PREDICTED: uncharacterized protein LOC108161779 [Drosophila
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>gi|1036698396|ref|XP_017149616.1| PREDICTED: uncharacterized protein LOC108159997 [Drosophila
>gi|1036757665|ref|XP_017156736.1| PREDICTED: uncharacterized protein LOC108165176 [Drosophila
>gi|1036750976|ref|XP_017156351.1| PREDICTED: probable insulin-like peptide 6 [Drosophila mira
>gi|1036757496|ref|XP_017156730.1| PREDICTED: uncharacterized protein LOC108165167 [Drosophila
>gi|1036724596|ref|XP_017146682.1| PREDICTED: apoptosis-inducing factor 3 isoform X1 [Drosophil
>gi|1036724612|ref|XP_017146693.1| PREDICTED: apoptosis-inducing factor 3 isoform X2 [Drosophil
>gi|1036724692|ref|XP_017146728.1| PREDICTED: uncharacterized protein LOC108158699 [Drosophila
>gi|1036724632|ref|XP_017146701.1| PREDICTED: apoptosis-inducing factor 3 isoform X3 [Drosophil
>gi|1036724671|ref|XP_017146716.1| PREDICTED: vacuolar protein sorting-associated protein 26 [I
>gi|1036724711|ref|XP_017146740.1| PREDICTED: ribosomal protein 63, mitochondrial [Drosophila r
>gi|1036739662|ref|XP_017155117.1| PREDICTED: required for meiotic nuclear division protein 1 l
>gi|1036739648|ref|XP_017155106.1| PREDICTED: anaphase-promoting complex subunit 7 [Drosophila
>gi|1036708557|ref|XP_017137723.1| PREDICTED: uncharacterized protein LOC108152708 isoform X1
>gi|1036708577|ref|XP_017137733.1| PREDICTED: uncharacterized protein LOC108152708 isoform X1
>gi|1036708594|ref|XP_017137741.1| PREDICTED: uncharacterized protein LOC108152708 isoform X2
>gi|1036708685|ref|XP_017137795.1| PREDICTED: calcyclin-binding protein [Drosophila miranda]
>gi|1036708668|ref|XP_017137782.1| PREDICTED: protein swallow [Drosophila miranda]
>gi|1036708611|ref|XP_017137753.1| PREDICTED: cell division cycle 7-related protein kinase [Drc
>gi|1036708630|ref|XP_017137762.1| PREDICTED: cell division cycle 7-related protein kinase [Drc
>gi|1036708651|ref|XP_017137769.1| PREDICTED: cell division cycle 7-related protein kinase [Drc
>gi|1036708766|ref|XP_017137847.1| PREDICTED: zinc finger protein 593 homolog [Drosophila miran
>gi|1036708702|ref|XP_017137808.1| PREDICTED: high affinity copper uptake protein 1 isoform X1
>gi|1036708719|ref|XP_017137816.1| PREDICTED: high affinity copper uptake protein 1 isoform X1
>gi|1036708733|ref|XP_017137826.1| PREDICTED: high affinity copper uptake protein 1 isoform X1
>gi|1036708750|ref|XP_017137834.1| PREDICTED: high affinity copper uptake protein 1 isoform X2
>gi|1036708814|ref|XP_017137869.1| PREDICTED: luc7-like protein 3 [Drosophila miranda]
>gi|1036735172|ref|XP_017152624.1| PREDICTED: DNA replication licensing factor Mcm6 [Drosophila
>gi|1036735189|ref|XP_017152635.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036713484|ref|XP_017140666.1| PREDICTED: fumarate hydratase, mitochondrial isoform X1 [Drc
>gi|1036713500|ref|XP_017140674.1| PREDICTED: fumarate hydratase, mitochondrial isoform X2 [Drc
>gi|1036713472|ref|XP_017140655.1| PREDICTED: fumarate hydratase, mitochondrial-like [Drosophil
>gi|1036713457|ref|XP_017140643.1| PREDICTED: uncharacterized protein LOC108154768 [Drosophila
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>gi|1036713537|ref|XP_017140699.1| PREDICTED: uncharacterized protein LOC108154807 [Drosophila
>gi|1036713516|ref|XP_017140687.1| PREDICTED: diphthine methyltransferase [Drosophila miranda]
>gi|1036713440|ref|XP_017140632.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila r
>gi|1036713424|ref|XP_017140622.1| PREDICTED: AF4/FMR2 family member 4 isoform X1 [Drosophila r
>gi|1036713569|ref|XP_017140717.1| PREDICTED: transcription factor Maf [Drosophila miranda]
>gi|1036704626|ref|XP_017135013.1| PREDICTED: uncharacterized protein LOC108151114 [Drosophila
>gi|1036704640|ref|XP_017135023.1| PREDICTED: uncharacterized protein LOC108151114 [Drosophila
>gi|1036698225|ref|XP_017134906.1| PREDICTED: breakpoint cluster region protein [Drosophila mir
>gi|1036704659|ref|XP_017135035.1| PREDICTED: probable RNA-binding protein 19 [Drosophila mirar
>gi|1036704714|ref|XP_017135067.1| PREDICTED: uncharacterized protein LOC108151149 [Drosophila
>gi|1036698240|ref|XP_017136144.1| PREDICTED: uncharacterized protein LOC108151567 [Drosophila
>gi|1036704677|ref|XP_017135047.1| PREDICTED: uncharacterized protein LOC108151131 [Drosophila
>gi|1036704697|ref|XP_017135055.1| PREDICTED: beta-1,3-galactosyltransferase brn [Drosophila m
>gi|1036704549|ref|XP_017134967.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704564|ref|XP_017134976.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704579|ref|XP_017134983.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704598|ref|XP_017134990.1| PREDICTED: uncharacterized protein LOC108151088 isoform X1
>gi|1036704613|ref|XP_017134999.1| PREDICTED: histone-lysine N-methyltransferase EHMT2 isoform
>gi|1036702005|ref|XP_017156117.1| PREDICTED: uncharacterized oxidoreductase SSP0419 [Drosophi
>gi|1036702021|ref|XP_017156125.1| PREDICTED: uncharacterized oxidoreductase SSP0419 [Drosophi
>gi|1036701898|ref|XP_017156051.1| PREDICTED: molybdenum cofactor synthesis protein cinnamon [I
>gi|1036702165|ref|XP_017156206.1| PREDICTED: cytochrome c oxidase assembly factor 6 homolog [I
>gi|1036702042|ref|XP_017156137.1| PREDICTED: uncharacterized protein LOC108164748 [Drosophila
>gi|1036701865|ref|XP_017156030.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofor
>gi|1036701881|ref|XP_017156038.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofor
>gi|1036701833|ref|XP_017156014.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofor
>gi|1036701848|ref|XP_017156023.1| PREDICTED: histone-lysine N-methyltransferase Suv4-20 isofor
>gi|1036702148|ref|XP_017156196.1| PREDICTED: RING-box protein 1A [Drosophila miranda]
>gi|1036701989|ref|XP_017156104.1| PREDICTED: uncharacterized protein LOC108164722 [Drosophila
>gi|1036701915|ref|XP_017156063.1| PREDICTED: protein SPT2 homolog [Drosophila miranda]
>gi|1036701951|ref|XP_017156084.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X2 [Drosop
>gi|1036701932|ref|XP_017156075.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X1 [Drosop
>gi|1036701970|ref|XP_017156092.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X3 [Drosop
>gi|1036702059|ref|XP_017156150.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702076|ref|XP_017156160.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702093|ref|XP_017156167.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702114|ref|XP_017156176.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036702131|ref|XP_017156185.1| PREDICTED: uncharacterized protein LOC108164755 [Drosophila
>gi|1036698410|ref|XP_017150823.1| PREDICTED: uncharacterized protein LOC108160753 [Drosophila
>gi|1036707387|ref|XP_017136928.1| PREDICTED: putative odorant receptor 92a [Drosophila miranda
>gi|1036705458|ref|XP_017135617.1| PREDICTED: uncharacterized protein LOC108151486 [Drosophila
>gi|1036703751|ref|XP_017157128.1| PREDICTED: uncharacterized protein LOC108165562 [Drosophila
>gi|1036713104|ref|XP_017140430.1| PREDICTED: protein sidekick isoform X1 [Drosophila miranda]
>gi|1036713121|ref|XP_017140436.1| PREDICTED: protein sidekick isoform X2 [Drosophila miranda]
>gi|1036713144|ref|XP_017140443.1| PREDICTED: protein sidekick isoform X3 [Drosophila miranda]
>gi|1036713161|ref|XP_017140449.1| PREDICTED: protein sidekick isoform X4 [Drosophila miranda]
>gi|1036713177|ref|XP_017140458.1| PREDICTED: protein sidekick isoform X5 [Drosophila miranda]
>gi|1036713194|ref|XP_017140466.1| PREDICTED: protein sidekick isoform X6 [Drosophila miranda]
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>gi|1036713211|ref|XP_017140475.1| PREDICTED: protein sidekick isoform X7 [Drosophila miranda]
>gi|1036704270|ref|XP_017134784.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036698425|ref|XP_017151896.1| PREDICTED: uncharacterized protein LOC108161692 [Drosophila
>gi|1036755177|ref|XP_017156601.1| PREDICTED: mitochondrial 2-oxodicarboxylate carrier [Drosoph
>gi|1036751123|ref|XP_017156360.1| PREDICTED: beta-amyloid-like protein isoform X1 [Drosophila
>gi|1036751141|ref|XP_017156361.1| PREDICTED: beta-amyloid-like protein isoform X1 [Drosophila
>gi|1036751215|ref|XP_017156365.1| PREDICTED: beta-amyloid-like protein isoform X5 [Drosophila
>gi|1036751197|ref|XP_017156364.1| PREDICTED: beta-amyloid-like protein isoform X4 [Drosophila
>gi|1036751179|ref|XP_017156363.1| PREDICTED: beta-amyloid-like protein isoform X3 [Drosophila
>gi|1036751234|ref|XP_017156366.1| PREDICTED: beta-amyloid-like protein isoform X6 [Drosophila
>gi|1036751160|ref|XP_017156362.1| PREDICTED: beta-amyloid-like protein isoform X2 [Drosophila
>gi|1036751251|ref|XP_017156367.1| PREDICTED: beta-amyloid-like protein isoform X7 [Drosophila
>gi|1036751269|ref|XP_017156369.1| PREDICTED: beta-amyloid-like protein isoform X8 [Drosophila
>gi|1036698440|ref|XP_017153096.1| PREDICTED: uncharacterized protein LOC108162461 [Drosophila
>gi|1036745739|ref|XP_017156066.1| PREDICTED: homeobox protein vnd isoform X2 [Drosophila mirar
>gi|1036745721|ref|XP_017156065.1| PREDICTED: homeobox protein vnd isoform X1 [Drosophila mirar
>gi|1036729959|ref|XP_017149863.1| PREDICTED: cytospin-A isoform X2 [Drosophila miranda]
>gi|1036729943|ref|XP_017149855.1| PREDICTED: cytospin-A isoform X1 [Drosophila miranda]
>gi|1036746629|ref|XP_017156120.1| PREDICTED: uncharacterized protein LOC108164738 [Drosophila
>gi|1036746612|ref|XP_017156119.1| PREDICTED: vacuolar protein sorting-associated protein 37C
>gi|1036753497|ref|XP_017156502.1| PREDICTED: ribokinase [Drosophila miranda]
>gi|1036708830|ref|XP_017137879.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036708846|ref|XP_017137887.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036708862|ref|XP_017137893.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036708879|ref|XP_017137902.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036708896|ref|XP_017137911.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036708913|ref|XP_017137920.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036708949|ref|XP_017137945.1| PREDICTED: histone acetyltransferase Tip60 [Drosophila mirar
>gi|1036708964|ref|XP_017137958.1| PREDICTED: uncharacterized protein LOC108152849 [Drosophila
>gi|1036708930|ref|XP_017137932.1| PREDICTED: tRNA (cytosine(34)-C(5))-methyltransferase [Drosc
>gi|1036708981|ref|XP_017137971.1| PREDICTED: uncharacterized protein LOC108152857 [Drosophila
>gi|1036749808|ref|XP_017156286.1| PREDICTED: beta-1,3-galactosyltransferase brn [Drosophila mi
>gi|1036740489|ref|XP_017155531.1| PREDICTED: DNA repair protein complementing XP-A cells homol
>gi|1036740455|ref|XP_017155506.1| PREDICTED: uncharacterized protein LOC108164395 [Drosophila
>gi|1036740472|ref|XP_017155516.1| PREDICTED: uncharacterized protein LOC108164395 [Drosophila
>gi|1036735086|ref|XP_017152569.1| PREDICTED: T-complex protein 1 subunit zeta-like [Drosophila
>gi|1036735103|ref|XP_017152582.1| PREDICTED: phosphatidylinositol 4,5-bisphosphate 5-phosphata
>gi|1036750227|ref|XP_017156309.1| PREDICTED: ubiquitin-conjugating enzyme E2 G2 [Drosophila mi
>gi|1036721608|ref|XP_017145191.1| PREDICTED: WD repeat-containing protein 18 [Drosophila mirar
>gi|1036721646|ref|XP_017145215.1| PREDICTED: UPF0184 protein CG14818 [Drosophila miranda]
>gi|1036721625|ref|XP_017145204.1| PREDICTED: APOPT family protein CG14806, mitochondrial [Dros
>gi|1036721591|ref|XP_017145180.1| PREDICTED: histone-lysine N-methyltransferase trr [Drosophil
>gi|1036708117|ref|XP_017137433.1| PREDICTED: 39S ribosomal protein L16, mitochondrial isoform
>gi|1036708134|ref|XP_017137440.1| PREDICTED: 39S ribosomal protein L16, mitochondrial isoform
>gi|1036708032|ref|XP_017137380.1| PREDICTED: armadillo segment polarity protein isoform X1 [Dr
>gi|1036708049|ref|XP_017137389.1| PREDICTED: armadillo segment polarity protein isoform X1 [Dr
>gi|1036708066|ref|XP_017137398.1| PREDICTED: armadillo segment polarity protein isoform X1 [Dr
>gi|1036708083|ref|XP_017137407.1| PREDICTED: armadillo segment polarity protein isoform X2 [Dr
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>gi|1036708151|ref|XP_017137451.1| PREDICTED: uncharacterized protein LOC108152546 isoform X1
>gi|1036708167|ref|XP_017137461.1| PREDICTED: uncharacterized protein LOC108152546 isoform X2
>gi|1036708182|ref|XP_017137474.1| PREDICTED: uncharacterized protein LOC108152557 isoform X1
>gi|1036708199|ref|XP_017137484.1| PREDICTED: uncharacterized protein LOC108152557 isoform X2
>gi|1036707981|ref|XP_017137343.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036707947|ref|XP_017137326.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036707965|ref|XP_017137335.1| PREDICTED: ER degradation-enhancing alpha-mannosidase-like p
>gi|1036707930|ref|XP_017137314.1| PREDICTED: WD repeat-containing protein 7 [Drosophila miranc
>gi|1036707997|ref|XP_017137355.1| PREDICTED: protein lin-9 homolog [Drosophila miranda]
>gi|1036708100|ref|XP_017137419.1| PREDICTED: translation initiation factor eIF-2B subunit eps:
>gi|1036708013|ref|XP_017137366.1| PREDICTED: type II inositol 1,4,5-trisphosphate 5-phosphatas
>gi|1036717028|ref|XP_017142775.1| PREDICTED: synaptobrevin homolog YKT6 [Drosophila miranda]
>gi|1036716881|ref|XP_017142683.1| PREDICTED: protein hold'em isoform X1 [Drosophila miranda]
>gi|1036716898|ref|XP_017142691.1| PREDICTED: protein hold'em isoform X2 [Drosophila miranda]
>gi|1036716864|ref|XP_017142672.1| PREDICTED: sorting nexin-25 [Drosophila miranda]
>gi|1036716994|ref|XP_017142754.1| PREDICTED: very-long-chain 3-oxoacyl-CoA reductase-B [Drosop
>gi|1036717013|ref|XP_017142763.1| PREDICTED: very-long-chain 3-oxoacyl-CoA reductase-B [Drosop
>gi|1036716937|ref|XP_017142720.1| PREDICTED: fibroblast growth factor receptor 3 [Drosophila r
>gi|1036716956|ref|XP_017142729.1| PREDICTED: fibroblast growth factor receptor 3 [Drosophila r
>gi|1036716975|ref|XP_017142739.1| PREDICTED: fibroblast growth factor receptor 3 [Drosophila r
>gi|1036716918|ref|XP_017142704.1| PREDICTED: bystin [Drosophila miranda]
>gi|1036749133|ref|XP_017156249.1| PREDICTED: 40S ribosomal protein S6 [Drosophila miranda]
>gi|1036742788|ref|XP_017155905.1| PREDICTED: general vesicular transport factor p115 [Drosoph:
>gi|1036724795|ref|XP_017146802.1| PREDICTED: uncharacterized protein LOC108158725 [Drosophila
>gi|1036724812|ref|XP_017146818.1| PREDICTED: N(G),N(G)-dimethylarginine dimethylaminohydrolase
>gi|1036728008|ref|XP_017148715.1| PREDICTED: cyclin-L1 [Drosophila miranda]
>gi|1036728027|ref|XP_017148722.1| PREDICTED: cyclin-L1 [Drosophila miranda]
>gi|1036728095|ref|XP_017148769.1| PREDICTED: DNA damage-regulated autophagy modulator protein
>gi|1036728040|ref|XP_017148734.1| PREDICTED: RNA 3'-terminal phosphate cyclase [Drosophila mi:
>gi|1036728059|ref|XP_017148748.1| PREDICTED: tRNA (guanine-N(7)-)-methyltransferase [Drosophil
>gi|1036728076|ref|XP_017148757.1| PREDICTED: tRNA (guanine-N(7)-)-methyltransferase [Drosophil
>gi|1036728114|ref|XP_017148782.1| PREDICTED: uncharacterized protein LOC108159749 [Drosophila
>gi|1036742463|ref|XP_017155887.1| PREDICTED: fasciculation and elongation protein zeta-2 [Dros
>gi|1036735539|ref|XP_017152824.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X3
>gi|1036735520|ref|XP_017152816.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X2
>gi|1036735502|ref|XP_017152808.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X1
>gi|1036735558|ref|XP_017152833.1| PREDICTED: tyrosine-protein phosphatase corkscrew isoform X4
>gi|1036745758|ref|XP_017156067.1| PREDICTED: uncharacterized protein LOC108164703 [Drosophila
>gi|1036749989|ref|XP_017156297.1| PREDICTED: AT-rich binding protein [Drosophila miranda]
>gi|1036717424|ref|XP_017142996.1| PREDICTED: uncharacterized protein LOC108156174 [Drosophila
>gi|1036717443|ref|XP_017143007.1| PREDICTED: protein suppressor of forked [Drosophila miranda]
>gi|1036717461|ref|XP_017143015.1| PREDICTED: protein suppressor of forked [Drosophila miranda]
>gi|1036698454|ref|XP_017154303.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036717511|ref|XP_017143038.1| PREDICTED: uncharacterized protein LOC108156208 [Drosophila
>gi|1036717530|ref|XP_017143050.1| PREDICTED: uncharacterized protein LOC108156220 [Drosophila
>gi|1036717480|ref|XP_017143027.1| PREDICTED: smad nuclear-interacting protein 1 [Drosophila mi
>gi|1036751937|ref|XP_017156405.1| PREDICTED: uncharacterized protein LOC108164928 [Drosophila
>gi|1036756860|ref|XP_017156694.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X1
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>gi|1036756876|ref|XP_017156695.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X2
>gi|1036757154|ref|XP_017156708.1| PREDICTED: sodium/potassium/calcium exchanger 4 [Drosophila
>gi|1036707741|ref|XP_017137192.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X1
>gi|1036707777|ref|XP_017137210.1| PREDICTED: sodium/potassium/calcium exchanger 4 isoform X3
>gi|1036707758|ref|XP_017137201.1| PREDICTED: sodium/potassium/calcium exchanger 3 isoform X2
>gi|1036707843|ref|XP_017137261.1| PREDICTED: cytochrome c oxidase assembly factor 7 homolog [
>gi|1036698468|ref|XP_017155452.1| PREDICTED: NAD-dependent protein deacylase Sirt4 [Drosophila
>gi|1036707671|ref|XP_017137148.1| PREDICTED: RNA-binding protein 25 isoform X1 [Drosophila mi
>gi|1036707688|ref|XP_017137157.1| PREDICTED: RNA-binding protein 25 isoform X2 [Drosophila mi
>gi|1036707870|ref|XP_017137276.1| PREDICTED: ras-related protein Rab-18 [Drosophila miranda]
>gi|1036707889|ref|XP_017137289.1| PREDICTED: 60S ribosomal protein L35 [Drosophila miranda]
>gi|1036707794|ref|XP_017137222.1| PREDICTED: protein trapped in endoderm-1 [Drosophila miranda
>gi|1036698483|ref|XP_017155879.1| PREDICTED: LOW QUALITY PROTEIN: gustatory receptor 5a for t
>gi|1036698498|ref|XP_017155992.1| PREDICTED: sulfated surface glycoprotein 185 [Drosophila mi
>gi|1036707707|ref|XP_017137170.1| PREDICTED: GPI inositol-deacylase [Drosophila miranda]
>gi|1036707826|ref|XP_017137249.1| PREDICTED: protein twisted gastrulation [Drosophila miranda]
>gi|1036707724|ref|XP_017137182.1| PREDICTED: paraplegin [Drosophila miranda]
>gi|1036707809|ref|XP_017137235.1| PREDICTED: circadian clock-controlled protein [Drosophila mi
>gi|1036707635|ref|XP_017137127.1| PREDICTED: period circadian protein isoform X2 [Drosophila m
>gi|1036707620|ref|XP_017137119.1| PREDICTED: period circadian protein isoform X1 [Drosophila m
>gi|1036707654|ref|XP_017137135.1| PREDICTED: period circadian protein isoform X3 [Drosophila m
>gi|1036698513|ref|XP_017156109.1| PREDICTED: uncharacterized protein LOC108164705 [Drosophila
>gi|1036756602|ref|XP_017156680.1| PREDICTED: tryptophan 2,3-dioxygenase isoform X1 [Drosophila
>gi|1036756619|ref|XP_017156681.1| PREDICTED: tryptophan 2,3-dioxygenase isoform X2 [Drosophila
>gi|1036751436|ref|XP_017156377.1| PREDICTED: poly(U)-specific endoribonuclease homolog isoform
>gi|1036751455|ref|XP_017156378.1| PREDICTED: poly(U)-specific endoribonuclease homolog isoform
>gi|1036735782|ref|XP_017152955.1| PREDICTED: unconventional myosin-XV isoform X1 [Drosophila m
>gi|1036735799|ref|XP_017152964.1| PREDICTED: uncharacterized protein LOC108162639 isoform X2
>gi|1036698528|ref|XP_017156222.1| PREDICTED: tenascin-X [Drosophila miranda]
>gi|1036750155|ref|XP_017156304.1| PREDICTED: uncharacterized protein LOC108164872 [Drosophila
>gi|1036750174|ref|XP_017156306.1| PREDICTED: uncharacterized protein LOC108164872 [Drosophila
>gi|1036750190|ref|XP_017156307.1| PREDICTED: uncharacterized protein LOC108164873 [Drosophila
>gi|1036698545|ref|XP_017156336.1| PREDICTED: uncharacterized protein LOC108164869 [Drosophila
>gi|1036754905|ref|XP_017156586.1| PREDICTED: protein msta isoform X1 [Drosophila miranda]
>gi|1036754924|ref|XP_017156587.1| PREDICTED: protein msta isoform X2 [Drosophila miranda]
>gi|1036698562|ref|XP_017156464.1| PREDICTED: polycystic kidney disease 2-like 2 protein [Droso
>gi|1036698577|ref|XP_017156582.1| PREDICTED: uncharacterized protein LOC108165018 [Drosophila
>gi|1036704892|ref|XP_017135217.1| PREDICTED: uncharacterized protein CG1552 isoform X2 [Drosop
>gi|1036704876|ref|XP_017135209.1| PREDICTED: uncharacterized protein CG1552 isoform X1 [Drosop
>gi|1036698592|ref|XP_017156696.1| PREDICTED: surface protein-like [Drosophila miranda]
>gi|1036705118|ref|XP_017135393.1| PREDICTED: uncharacterized protein LOC108151345 [Drosophila
>gi|1036705865|ref|XP_017135889.1| PREDICTED: uncharacterized protein LOC108151664 [Drosophila
>gi|1036703770|ref|XP_017157149.1| PREDICTED: uncharacterized protein LOC108165584 [Drosophila
>gi|1036741499|ref|XP_017155834.1| PREDICTED: protein sevenless isoform X2 [Drosophila miranda]
>gi|1036741462|ref|XP_017155831.1| PREDICTED: protein sevenless isoform X1 [Drosophila miranda]
>gi|1036741481|ref|XP_017155833.1| PREDICTED: protein sevenless isoform X1 [Drosophila miranda]
>gi|1036735870|ref|XP_017153010.1| PREDICTED: uncharacterized protein LOC108162674 isoform X1
>gi|1036735890|ref|XP_017153018.1| PREDICTED: uncharacterized protein LOC108162674 isoform X2
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>gi|1036735851|ref|XP_017152998.1| PREDICTED: growth hormone-inducible transmembrane protein [
>gi|1036735815|ref|XP_017152977.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036735832|ref|XP_017152986.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036755417|ref|XP_017156615.1| PREDICTED: apyrase isoform X2 [Drosophila miranda]
>gi|1036755398|ref|XP_017156614.1| PREDICTED: apyrase isoform X1 [Drosophila miranda]
>gi|1036710670|ref|XP_017139039.1| PREDICTED: RNA polymerase II subunit A C-terminal domain ph
>gi|1036710653|ref|XP_017139026.1| PREDICTED: glucose-6-phosphate 1-dehydrogenase [Drosophila n
>gi|1036710620|ref|XP_017139002.1| PREDICTED: cytokine receptor isoform X2 [Drosophila miranda]
>gi|1036710602|ref|XP_017138993.1| PREDICTED: cytokine receptor isoform X1 [Drosophila miranda]
>gi|1036710585|ref|XP_017138981.1| PREDICTED: cytokine receptor [Drosophila miranda]
>gi|1036710636|ref|XP_017139013.1| PREDICTED: ubiquilin-1 [Drosophila miranda]
>gi|1036742409|ref|XP_017155884.1| PREDICTED: moesin/ezrin/radixin homolog 2 [Drosophila miranc
>gi|1036752212|ref|XP_017156422.1| PREDICTED: uncharacterized protein LOC108164944 isoform X2
>gi|1036752191|ref|XP_017156421.1| PREDICTED: uncharacterized protein LOC108164944 isoform X1
>gi|1036744272|ref|XP_017155982.1| PREDICTED: cdc42 homolog [Drosophila miranda]
>gi|1036743892|ref|XP_017155963.1| PREDICTED: probable tRNA N6-adenosine threonylcarbamoyltrans
>gi|1036743875|ref|XP_017155962.1| PREDICTED: Golgi resident protein GCP60 [Drosophila miranda]
>gi|1036736935|ref|XP_017153635.1| PREDICTED: probable RNA-binding protein CG14230 [Drosophila
>gi|1036736951|ref|XP_017153648.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036736968|ref|XP_017153655.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036736986|ref|XP_017153664.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036737004|ref|XP_017153672.1| PREDICTED: uncharacterized protein LOC108163076 [Drosophila
>gi|1036754508|ref|XP_017156561.1| PREDICTED: high affinity cationic amino acid transporter 1
>gi|1036748701|ref|XP_017156226.1| PREDICTED: chorion peroxidase [Drosophila miranda]
>gi|1036750387|ref|XP_017156316.1| PREDICTED: dipeptidyl aminopeptidase-like protein 6 isoform
>gi|1036750368|ref|XP_017156315.1| PREDICTED: dipeptidyl aminopeptidase-like protein 6 isoform
>gi|1036750407|ref|XP_017156317.1| PREDICTED: uncharacterized protein LOC108164881 [Drosophila
>gi|1036734133|ref|XP_017152045.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X2
>gi|1036734076|ref|XP_017152018.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X1
>gi|1036734092|ref|XP_017152029.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X1
>gi|1036734114|ref|XP_017152037.1| PREDICTED: uncharacterized protein DDB_G0287625 isoform X1
>gi|1036749674|ref|XP_017156279.1| PREDICTED: uncharacterized protein LOC108164860 isoform X1
>gi|1036749693|ref|XP_017156280.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036749712|ref|XP_017156281.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036749731|ref|XP_017156282.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036749749|ref|XP_017156283.1| PREDICTED: uncharacterized protein LOC108164860 isoform X2
>gi|1036698606|ref|XP_017156772.1| PREDICTED: uncharacterized protein LOC108165203 [Drosophila
>gi|1036757208|ref|XP_017156711.1| PREDICTED: uncharacterized protein LOC108165156 [Drosophila
>gi|1036733239|ref|XP_017151520.1| PREDICTED: proton-coupled amino acid transporter 4 isoform >
>gi|1036733292|ref|XP_017151545.1| PREDICTED: proton-coupled amino acid transporter 1 [Drosophi
>gi|1036733255|ref|XP_017151527.1| PREDICTED: proton-coupled amino acid transporter 4 isoform >
>gi|1036733273|ref|XP_017151535.1| PREDICTED: proton-coupled amino acid transporter 4 isoform >
>gi|1036733347|ref|XP_017151579.1| PREDICTED: actin-related protein 2/3 complex subunit 3 [Dros
>gi|1036733329|ref|XP_017151569.1| PREDICTED: LOW QUALITY PROTEIN: rab-like protein 3 [Drosophi
>gi|1036733311|ref|XP_017151554.1| PREDICTED: decapping nuclease DXO homolog [Drosophila miranc
>gi|1036744610|ref|XP_017156000.1| PREDICTED: uncharacterized protein LOC108164657 isoform X2
>gi|1036744591|ref|XP_017155999.1| PREDICTED: uncharacterized protein LOC108164657 isoform X1
>gi|1036698621|ref|XP_017156875.1| PREDICTED: uncharacterized protein LOC108165318 [Drosophila
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>gi|1036711462|ref|XP_017139492.1| PREDICTED: TM2 domain-containing protein almondex [Drosophil
>gi|1036711445|ref|XP_017139482.1| PREDICTED: la-related protein 7 [Drosophila miranda]
>gi|1036711340|ref|XP_017139437.1| PREDICTED: uncharacterized protein LOC108153846 isoform X1
>gi|1036711357|ref|XP_017139442.1| PREDICTED: uncharacterized protein LOC108153846 isoform X2
>gi|1036711479|ref|XP_017139501.1| PREDICTED: uncharacterized protein LOC108153913 [Drosophila
>gi|1036711374|ref|XP_017139452.1| PREDICTED: protein GPR107 isoform X1 [Drosophila miranda]
>gi|1036711393|ref|XP_017139460.1| PREDICTED: protein GPR107 isoform X2 [Drosophila miranda]
>gi|1036711408|ref|XP_017139468.1| PREDICTED: protein GPR107 isoform X2 [Drosophila miranda]
>gi|1036711428|ref|XP_017139474.1| PREDICTED: protein GPR107 isoform X2 [Drosophila miranda]
>gi|1036698635|ref|XP_017156987.1| PREDICTED: gustatory receptor 8a [Drosophila miranda]
>gi|1036698651|ref|XP_017157107.1| PREDICTED: uncharacterized protein LOC108165520 [Drosophila
>gi|1036752267|ref|XP_017156425.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036752282|ref|XP_017156426.1| PREDICTED: uncharacterized protein LOC108164948 [Drosophila
>gi|1036752300|ref|XP_017156428.1| PREDICTED: uncharacterized protein LOC108164948 [Drosophila
>gi|1036752320|ref|XP_017156429.1| PREDICTED: uncharacterized protein LOC108164948 [Drosophila
>gi|1036754740|ref|XP_017156576.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036754756|ref|XP_017156577.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036754771|ref|XP_017156578.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036754787|ref|XP_017156579.1| PREDICTED: uncharacterized protein LOC108165047 [Drosophila
>gi|1036743966|ref|XP_017155967.1| PREDICTED: kinesin-associated protein 3 [Drosophila miranda]
>gi|1036730760|ref|XP_017150382.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036730779|ref|XP_017150390.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036730797|ref|XP_017150400.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036730815|ref|XP_017150408.1| PREDICTED: receptor-mediated endocytosis protein 6 homolog
>gi|1036698667|ref|XP_017134572.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036740943|ref|XP_017155800.1| PREDICTED: uncharacterized protein C1683.06c [Drosophila mi
>gi|1036740959|ref|XP_017155801.1| PREDICTED: inosine-uridine preferring nucleoside hydrolase
>gi|1036714435|ref|XP_017141260.1| PREDICTED: probable glutamine-dependent NAD(+) synthetase [
>gi|1036714235|ref|XP_017141152.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X1 [Dros
>gi|1036714273|ref|XP_017141171.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X3 [Dros
>gi|1036714345|ref|XP_017141205.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X7 [Dros
>gi|1036714290|ref|XP_017141181.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X4 [Dros
>gi|1036714309|ref|XP_017141190.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X5 [Dros
>gi|1036714327|ref|XP_017141199.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X6 [Dros
>gi|1036714254|ref|XP_017141160.1| PREDICTED: cytosolic carboxypeptidase NnaD isoform X2 [Dros
>gi|1036714363|ref|XP_017141214.1| PREDICTED: uncharacterized protein LOC108155072 isoform X8
>gi|1036714380|ref|XP_017141222.1| PREDICTED: uncharacterized protein LOC108155072 isoform X9
>gi|1036714416|ref|XP_017141248.1| PREDICTED: RING finger protein 157 [Drosophila miranda]
>gi|1036714395|ref|XP_017141236.1| PREDICTED: DNA topoisomerase 2-binding protein 1-A [Drosophi
>gi|1036739911|ref|XP_017155258.1| PREDICTED: AP-3 complex subunit delta [Drosophila miranda]
>gi|1036731592|ref|XP_017150790.1| PREDICTED: gamma-tubulin complex component 3 homolog [Droso
>gi|1036731609|ref|XP_017150801.1| PREDICTED: probable methylthioribulose-1-phosphate dehydrata
>gi|1036731626|ref|XP_017150813.1| PREDICTED: peroxisomal multifunctional enzyme type 2 [Droso
>gi|1036711983|ref|XP_017139800.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036711932|ref|XP_017139760.1| PREDICTED: programmed cell death protein 4 [Drosophila mira
>gi|1036712017|ref|XP_017139818.1| PREDICTED: gametocyte-specific factor 1 homolog isoform X2
>gi|1036712031|ref|XP_017139828.1| PREDICTED: gametocyte-specific factor 1 isoform X3 [Drosophi
>gi|1036712000|ref|XP_017139812.1| PREDICTED: gametocyte-specific factor 1 homolog isoform X1
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>gi|1036711966|ref|XP_017139787.1| PREDICTED: probable RNA 3'-terminal phosphate cyclase-like p
>gi|1036711949|ref|XP_017139770.1| PREDICTED: vitellogenin-3 [Drosophila miranda]
>gi|1036711881|ref|XP_017139731.1| PREDICTED: protein retinal degeneration B isoform X3 [Droso
>gi|1036711897|ref|XP_017139739.1| PREDICTED: protein retinal degeneration B isoform X4 [Droso
>gi|1036711914|ref|XP_017139748.1| PREDICTED: protein retinal degeneration B isoform X5 [Droso
>gi|1036711829|ref|XP_017139707.1| PREDICTED: protein retinal degeneration B isoform X1 [Droso
>gi|1036711847|ref|XP_017139715.1| PREDICTED: protein retinal degeneration B isoform X1 [Droso
>gi|1036711864|ref|XP_017139723.1| PREDICTED: protein retinal degeneration B isoform X2 [Droso
>gi|1036746543|ref|XP_017156114.1| PREDICTED: cathepsin B [Drosophila miranda]
>gi|1036702340|ref|XP_017156305.1| PREDICTED: TM2 domain-containing protein CG11103 [Drosophila
>gi|1036721009|ref|XP_017144895.1| PREDICTED: probable tubulin polyglutamylase TTLL1 isoform X2
>gi|1036721046|ref|XP_017144916.1| PREDICTED: myb-like protein I [Drosophila miranda]
>gi|1036720990|ref|XP_017144886.1| PREDICTED: probable tubulin polyglutamylase TTLL1 isoform X1
>gi|1036720953|ref|XP_017144868.1| PREDICTED: partitioning defective 3 homolog isoform X1 [Dros
>gi|1036720971|ref|XP_017144875.1| PREDICTED: partitioning defective 3 homolog isoform X2 [Dros
>gi|1036721028|ref|XP_017144902.1| PREDICTED: probable tubulin polyglutamylase TTLL1 isoform X3
>gi|1036706297|ref|XP_017136210.1| PREDICTED: keratin-associated protein 6-2 [Drosophila mirand
>gi|1036757481|ref|XP_017156729.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1036707913|ref|XP_017137303.1| PREDICTED: uncharacterized protein LOC108152451 [Drosophila
>gi|1036698686|ref|XP_017134677.1| PREDICTED: uncharacterized protein LOC108150846, partial [Dr
>gi|1036698703|ref|XP_017134789.1| PREDICTED: phospholipase A1 2-like [Drosophila miranda]
>gi|1036753104|ref|XP_017156478.1| PREDICTED: probable phospholipase A1 magnifin [Drosophila mi
>gi|1036753087|ref|XP_017156477.1| PREDICTED: vitellogenin-3 [Drosophila miranda]
>gi|1036707490|ref|XP_017136991.1| PREDICTED: heterogeneous nuclear ribonucleoprotein A1 [Droso
>gi|1036706066|ref|XP_017136048.1| PREDICTED: glycine-rich cell wall structural protein 1.0 [Dr
>gi|1036705208|ref|XP_017135451.1| PREDICTED: glycine-rich cell wall structural protein 2 [Dros
>gi|1036707604|ref|XP_017137094.1| PREDICTED: ctenidin-1 [Drosophila miranda]
>gi|1036698719|ref|XP_017134907.1| PREDICTED: glycine-rich cell wall structural protein 1.0 [Dr
>gi|1036737299|ref|XP_017153840.1| PREDICTED: DExH-box ATP-dependent RNA helicase DExH6 [Drosop
>gi|1036737317|ref|XP_017153852.1| PREDICTED: uncharacterized protein C9orf85 homolog [Drosophi
>gi|1036737249|ref|XP_017153811.1| PREDICTED: uncharacterized protein LOC108163182 isoform X1
>gi|1036737267|ref|XP_017153820.1| PREDICTED: myb-like protein I isoform X2 [Drosophila miranda
>gi|1036737281|ref|XP_017153828.1| PREDICTED: espin isoform X3 [Drosophila miranda]
>gi|1036704126|ref|XP_017134701.1| PREDICTED: uncharacterized protein LOC108150899 [Drosophila
>gi|1036704439|ref|XP_017134894.1| PREDICTED: uncharacterized protein LOC108151036 [Drosophila
>gi|1036733525|ref|XP_017151683.1| PREDICTED: plastin-1 isoform X1 [Drosophila miranda]
>gi|1036733543|ref|XP_017151691.1| PREDICTED: plastin-1 isoform X2 [Drosophila miranda]
>gi|1036746470|ref|XP_017156110.1| PREDICTED: uncharacterized protein C6orf106 homolog [Drosoph
>gi|1036754219|ref|XP_017156545.1| PREDICTED: homeobox protein B-H2 [Drosophila miranda]
>gi|1036755435|ref|XP_017156616.1| PREDICTED: homeobox protein B-H1 isoform X1 [Drosophila mira
>gi|1036755454|ref|XP_017156617.1| PREDICTED: homeobox protein B-H1 isoform X2 [Drosophila mira
>gi|1036704924|ref|XP_017135242.1| PREDICTED: E3 SUMO-protein ligase PIAS1-like [Drosophila mir
>gi|1036735263|ref|XP_017152676.1| PREDICTED: probable ATP-dependent RNA helicase CG8611 [Droso
>gi|1036707111|ref|XP_017136761.1| PREDICTED: uncharacterized protein LOC108152152 [Drosophila
>gi|1036741379|ref|XP_017155826.1| PREDICTED: cytosolic endo-beta-N-acetylglucosaminidase [Dros
>gi|1036741396|ref|XP_017155827.1| PREDICTED: cytosolic endo-beta-N-acetylglucosaminidase [Dros
>gi|1036741445|ref|XP_017155830.1| PREDICTED: uncharacterized protein LOC108164539 isoform X2
>gi|1036741426|ref|XP_017155829.1| PREDICTED: uncharacterized protein LOC108164539 isoform X1
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>gi|1036741413|ref|XP_017155828.1| PREDICTED: uncharacterized protein LOC108164538 [Drosophila
>gi|1036757284|ref|XP_017156716.1| PREDICTED: uncharacterized protein LOC108165160 [Drosophila
>gi|1036698734|ref|XP_017135031.1| PREDICTED: acidic leucine-rich nuclear phosphoprotein 32-rel
>gi|1036747464|ref|XP_017156170.1| PREDICTED: ubiquitin-like modifier-activating enzyme 5 [Dros
>gi|1036747798|ref|XP_017156190.1| PREDICTED: signal peptidase complex subunit 2 [Drosophila mi
>gi|1036747779|ref|XP_017156189.1| PREDICTED: probable tubulin beta chain CG32396 [Drosophila m
>gi|1036710810|ref|XP_017139125.1| PREDICTED: death-associated protein kinase related-like [Dro
>gi|1036710827|ref|XP_017139133.1| PREDICTED: death-associated protein kinase related-like [Dro
>gi|1036698751|ref|XP_017135152.1| PREDICTED: death-associated protein kinase related, partial
>gi|1036698770|ref|XP_017135281.1| PREDICTED: cytochrome P450 4g15-like, partial [Drosophila mi
>gi|1036704155|ref|XP_017134724.1| PREDICTED: uncharacterized protein LOC108150906 isoform X2
>gi|1036704141|ref|XP_017134715.1| PREDICTED: uncharacterized protein LOC108150906 isoform X1
>gi|1036704173|ref|XP_017134732.1| PREDICTED: uncharacterized protein LOC108150906 isoform X3
>gi|1036698781|ref|XP_017135401.1| PREDICTED: maternal protein exuperantia-2-like [Drosophila m
>gi|1036751808|ref|XP_017156397.1| PREDICTED: cytochrome P450 4g15 isoform X1 [Drosophila miran
>gi|1036751825|ref|XP_017156398.1| PREDICTED: cytochrome P450 4g15 isoform X2 [Drosophila miran
>gi|1036750023|ref|XP_017156299.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750042|ref|XP_017156300.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750062|ref|XP_017156301.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750006|ref|XP_017156298.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750083|ref|XP_017156302.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036750102|ref|XP_017156303.1| PREDICTED: TWiK family of potassium channels protein 7 isofo
>gi|1036720713|ref|XP_017144781.1| PREDICTED: neurofilament heavy polypeptide [Drosophila miran
>gi|1036720694|ref|XP_017144769.1| PREDICTED: protein flightless-1 [Drosophila miranda]
>gi|1036747985|ref|XP_017156197.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748002|ref|XP_017156198.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748021|ref|XP_017156199.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748038|ref|XP_017156200.1| PREDICTED: protein tweety isoform X1 [Drosophila miranda]
>gi|1036748053|ref|XP_017156201.1| PREDICTED: protein tweety isoform X2 [Drosophila miranda]
>gi|1036748070|ref|XP_017156202.1| PREDICTED: protein tweety isoform X3 [Drosophila miranda]
>gi|1036742160|ref|XP_017155868.1| PREDICTED: pyridoxal-dependent decarboxylase domain-containi
>gi|1036757908|ref|XP_017156749.1| PREDICTED: OTU domain-containing protein 1-like [Drosophila
>gi|1036757927|ref|XP_017156750.1| PREDICTED: OTU domain-containing protein 1-like [Drosophila
>gi|1036726043|ref|XP_017147530.1| PREDICTED: acetylcholine receptor subunit alpha-like [Droso
>gi|1036726076|ref|XP_017147552.1| PREDICTED: uncharacterized protein LOC108159091 isoform X2
>gi|1036726063|ref|XP_017147543.1| PREDICTED: uncharacterized protein LOC108159091 isoform X1
>gi|1036726093|ref|XP_017147565.1| PREDICTED: pyridoxine/pyridoxamine 5'-phosphate oxidase [Dr
>gi|1036726024|ref|XP_017147517.1| PREDICTED: uncharacterized protein LOC108159070 isoform X2
>gi|1036726005|ref|XP_017147510.1| PREDICTED: uncharacterized protein LOC108159070 isoform X1
>gi|1036726112|ref|XP_017147578.1| PREDICTED: uncharacterized protein LOC108159109 [Drosophila
>gi|1036747817|ref|XP_017156191.1| PREDICTED: protein RFT1 homolog [Drosophila miranda]
>gi|1036726492|ref|XP_017147819.1| PREDICTED: uncharacterized protein LOC108159225 [Drosophila
>gi|1036726505|ref|XP_017147832.1| PREDICTED: UDP-N-acetylglucosamine transporter [Drosophila m
>gi|1036726524|ref|XP_017147846.1| PREDICTED: translation initiation factor eIF-2B subunit beta
>gi|1036726541|ref|XP_017147858.1| PREDICTED: phosphoglycolate phosphatase 2 [Drosophila miran
>gi|1036726439|ref|XP_017147786.1| PREDICTED: uncharacterized protein LOC108159214 isoform X1
>gi|1036726458|ref|XP_017147797.1| PREDICTED: uncharacterized protein LOC108159214 isoform X2
>gi|1036726476|ref|XP_017147807.1| PREDICTED: uncharacterized protein LOC108159214 isoform X3
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>gi|1036745594|ref|XP_017156057.1| PREDICTED: WW domain-binding protein 11 [Drosophila miranda]
>gi|1036732702|ref|XP_017151275.1| PREDICTED: activating signal cointegrator 1 complex subunit
>gi|1036732723|ref|XP_017151287.1| PREDICTED: mitosis initiation protein fs(1)Ya [Drosophila m
>gi|1036732742|ref|XP_017151298.1| PREDICTED: RING finger protein vilya isoform X1 [Drosophila
>gi|1036732761|ref|XP_017151306.1| PREDICTED: RING finger protein vilya isoform X2 [Drosophila
>gi|1036718943|ref|XP_017143840.1| PREDICTED: zinc finger protein 391 [Drosophila miranda]
>gi|1036718964|ref|XP_017143851.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036698798|ref|XP_017135524.1| PREDICTED: zinc finger protein 569 [Drosophila miranda]
>gi|1036718923|ref|XP_017143828.1| PREDICTED: protein cramped [Drosophila miranda]
>gi|1036718983|ref|XP_017143862.1| PREDICTED: syntaxin-4 isoform X1 [Drosophila miranda]
>gi|1036719003|ref|XP_017143871.1| PREDICTED: syntaxin-4 isoform X2 [Drosophila miranda]
>gi|1036705326|ref|XP_017135528.1| PREDICTED: uncharacterized protein LOC108151430 isoform X2
>gi|1036705311|ref|XP_017135519.1| PREDICTED: uncharacterized protein LOC108151430 isoform X1
>gi|1036752884|ref|XP_017156465.1| PREDICTED: uncharacterized protein LOC108164964 [Drosophila
>gi|1036752901|ref|XP_017156466.1| PREDICTED: uncharacterized protein LOC108164964 [Drosophila
>gi|1036698813|ref|XP_017135648.1| PREDICTED: POC1 centriolar protein homolog A-like [Drosophil
>gi|1036752919|ref|XP_017156467.1| PREDICTED: ARL-6-interacting protein 1 homolog [Drosophila r
>gi|1036703713|ref|XP_017157108.1| PREDICTED: uncharacterized protein LOC108165541 [Drosophila
>gi|1036707146|ref|XP_017136781.1| PREDICTED: uncharacterized protein LOC108152167 [Drosophila
>gi|1036698828|ref|XP_017135766.1| PREDICTED: uncharacterized protein LOC108151568 [Drosophila
>gi|1036757172|ref|XP_017156709.1| PREDICTED: TAF5-like RNA polymerase II p300/CBP-associated f
>gi|1036705033|ref|XP_017135326.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1-like [Drosophil
>gi|1036735617|ref|XP_017152866.1| PREDICTED: transmembrane protein 120 homolog isoform X1 [Dro
>gi|1036735635|ref|XP_017152876.1| PREDICTED: transmembrane protein 120 homolog isoform X2 [Dro
>gi|1036735598|ref|XP_017152854.1| PREDICTED: protein white isoform X2 [Drosophila miranda]
>gi|1036735579|ref|XP_017152845.1| PREDICTED: protein white isoform X1 [Drosophila miranda]
>gi|1036735650|ref|XP_017152888.1| PREDICTED: protein angel-like [Drosophila miranda]
>gi|1036705224|ref|XP_017135465.1| PREDICTED: RNA polymerase-associated protein Rtf1 [Drosophil
>gi|1036704350|ref|XP_017134836.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036705156|ref|XP_017135417.1| PREDICTED: protein germ cell-less-like [Drosophila miranda]
>gi|1036705135|ref|XP_017135405.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036698842|ref|XP_017135892.1| PREDICTED: anaphase-promoting complex subunit cdh1 [Drosophi
>gi|1036698859|ref|XP_017136018.1| PREDICTED: uncharacterized protein LOC108151713 [Drosophila
>gi|1036698874|ref|XP_017136145.1| PREDICTED: putative cyclin-dependent serine/threonine-protei
>gi|1036706528|ref|XP_017136355.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036756826|ref|XP_017156692.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 [Drosophila mir
>gi|1036755996|ref|XP_017156644.1| PREDICTED: glycine, alanine and asparagine-rich protein [Dro
>gi|1036708233|ref|XP_017137508.1| PREDICTED: uncharacterized protein LOC108152577 [Drosophila
>gi|1036698889|ref|XP_017136270.1| PREDICTED: eukaryotic translation initiation factor 4 gamma
>gi|1036706634|ref|XP_017136437.1| PREDICTED: uncharacterized protein LOC108151980 [Drosophila
>gi|1036698904|ref|XP_017136392.1| PREDICTED: PH domain-containing protein DDB_G0287875 [Drosop
>gi|1036698917|ref|XP_017136516.1| PREDICTED: putative uncharacterized protein DDB_G0274435 [Di
>gi|1036698931|ref|XP_017136644.1| PREDICTED: protein PRY2 [Drosophila miranda]
>gi|1036698948|ref|XP_017136777.1| PREDICTED: uncharacterized protein LOC108152142 [Drosophila
>gi|1036698961|ref|XP_017136902.1| PREDICTED: uncharacterized protein LOC108152210 [Drosophila
>gi|1036754447|ref|XP_017156558.1| PREDICTED: estradiol 17-beta-dehydrogenase 8 [Drosophila mir
>gi|1036739106|ref|XP_017154782.1| PREDICTED: irregular chiasm C-roughest protein [Drosophila r
>gi|1036703531|ref|XP_017157001.1| PREDICTED: uncharacterized protein LOC108165462 [Drosophila
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>gi|1036751645|ref|XP_017156389.1| PREDICTED: uncharacterized protein LOC108164914 [Drosophila
>gi|1036698975|ref|XP_017137021.1| PREDICTED: jacalin-related lectin 34 [Drosophila miranda]
>gi|1036755261|ref|XP_017156606.1| PREDICTED: hepatoma-derived growth factor-related protein 2-
>gi|1036755277|ref|XP_017156607.1| PREDICTED: uncharacterized protein LOC108165069 isoform X2
>gi|1036715997|ref|XP_017142187.1| PREDICTED: irregular chiasm C-roughest protein isoform X1 [
>gi|1036716017|ref|XP_017142194.1| PREDICTED: irregular chiasm C-roughest protein isoform X2 [
>gi|1036716037|ref|XP_017142203.1| PREDICTED: irregular chiasm C-roughest protein isoform X2 [
>gi|1036715979|ref|XP_017142172.1| PREDICTED: neurogenic locus Notch protein [Drosophila miranc
>gi|1036752611|ref|XP_017156446.1| PREDICTED: UPF0729 protein GD16342 [Drosophila miranda]
>gi|1036742016|ref|XP_017155862.1| PREDICTED: follicle cell protein 3C-1 [Drosophila miranda]
>gi|1036742037|ref|XP_017155863.1| PREDICTED: uncharacterized protein LOC108164557 [Drosophila
>gi|1036742056|ref|XP_017155864.1| PREDICTED: thioredoxin domain-containing protein 17 [Drosopl
>gi|1036719280|ref|XP_017144018.1| PREDICTED: uncharacterized protein LOC108156826 [Drosophila
>gi|1036719338|ref|XP_017144056.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036719319|ref|XP_017144043.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036719360|ref|XP_017144068.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036719299|ref|XP_017144031.1| PREDICTED: alpha-tubulin N-acetyltransferase 1-like [Drosopl
>gi|1036719037|ref|XP_017143911.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719054|ref|XP_017143919.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719074|ref|XP_017143927.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719095|ref|XP_017143936.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719120|ref|XP_017143944.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719139|ref|XP_017143952.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719159|ref|XP_017143961.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719179|ref|XP_017143969.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719197|ref|XP_017143976.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase, iso
>gi|1036719262|ref|XP_017144005.1| PREDICTED: katanin p60 ATPase-containing subunit A-like 2 [
>gi|1036719219|ref|XP_017143984.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036719244|ref|XP_017143993.1| PREDICTED: cAMP-specific 3',5'-cyclic phosphodiesterase iso
>gi|1036739750|ref|XP_017155172.1| PREDICTED: septin-5 isoform X1 [Drosophila miranda]
>gi|1036739769|ref|XP_017155178.1| PREDICTED: septin-5 isoform X2 [Drosophila miranda]
>gi|1036698998|ref|XP_017137147.1| PREDICTED: transmembrane protease serine 9-like [Drosophila
>gi|1036746332|ref|XP_017156100.1| PREDICTED: trypsin beta [Drosophila miranda]
>gi|1036746368|ref|XP_017156102.1| PREDICTED: serine protease SP24D [Drosophila miranda]
>gi|1036746351|ref|XP_017156101.1| PREDICTED: serine protease SP24D [Drosophila miranda]
>gi|1036746225|ref|XP_017156094.1| PREDICTED: carboxypeptidase D isoform X1 [Drosophila mirand;
>gi|1036746244|ref|XP_017156095.1| PREDICTED: carboxypeptidase D isoform X1 [Drosophila mirand;
>gi|1036746263|ref|XP_017156096.1| PREDICTED: carboxypeptidase D isoform X2 [Drosophila mirand;
>gi|1036746282|ref|XP_017156097.1| PREDICTED: carboxypeptidase D isoform X3 [Drosophila mirand;
>gi|1036746297|ref|XP_017156098.1| PREDICTED: carboxypeptidase D isoform X4 [Drosophila mirand;
>gi|1036746314|ref|XP_017156099.1| PREDICTED: carboxypeptidase D isoform X5 [Drosophila mirand;
>gi|1036746209|ref|XP_017156093.1| PREDICTED: uncharacterized protein LOC108164719 [Drosophila
>gi|1036728992|ref|XP_017149297.1| PREDICTED: transforming growth factor beta-1-induced transci
>gi|1036729007|ref|XP_017149305.1| PREDICTED: transforming growth factor beta-1-induced transci
>gi|1036728909|ref|XP_017149239.1| PREDICTED: uncharacterized protein LOC108160007 isoform X2
>gi|1036728889|ref|XP_017149230.1| PREDICTED: uncharacterized protein LOC108160007 isoform X1
>gi|1036728873|ref|XP_017149217.1| PREDICTED: endothelin-converting enzyme 1-like isoform X2 [
>gi|1036728855|ref|XP_017149208.1| PREDICTED: endothelin-converting enzyme 1-like isoform X1 [
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>gi|1036728940|ref|XP_017149263.1| PREDICTED: uncharacterized protein LOC108160031 [Drosophila
>gi|1036728959|ref|XP_017149274.1| PREDICTED: uncharacterized protein LOC108160038 [Drosophila
>gi|1036728976|ref|XP_017149286.1| PREDICTED: uncharacterized protein LOC108160049 [Drosophila
>gi|1036728925|ref|XP_017149251.1| PREDICTED: keratin, type I cytoskeletal 10-like [Drosophila
>gi|1036704855|ref|XP_017135194.1| PREDICTED: casein kinase II subunit beta'-like [Drosophila r
>gi|1036753532|ref|XP_017156504.1| PREDICTED: TWiK family of potassium channels protein 7 [Dros
>gi|1036753553|ref|XP_017156505.1| PREDICTED: lanC-like protein 3 homolog [Drosophila miranda]
>gi|1036752754|ref|XP_017156455.1| PREDICTED: transmembrane emp24 domain-containing protein 2
>gi|1036743155|ref|XP_017155927.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036743136|ref|XP_017155926.1| PREDICTED: uncharacterized protein LOC108164598 [Drosophila
>gi|1036730851|ref|XP_017150432.1| PREDICTED: uncharacterized protein LOC108160747 [Drosophila
>gi|1036730833|ref|XP_017150421.1| PREDICTED: structural maintenance of chromosomes protein 3
>gi|1036727974|ref|XP_017148695.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein pho
>gi|1036736569|ref|XP_017153421.1| PREDICTED: actin-related protein 2 isoform X1 [Drosophila m:
>gi|1036736587|ref|XP_017153430.1| PREDICTED: actin-related protein 2 isoform X1 [Drosophila m:
>gi|1036736607|ref|XP_017153438.1| PREDICTED: actin-related protein 2 isoform X2 [Drosophila m:
>gi|1036756469|ref|XP_017156672.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila r
>gi|1036756487|ref|XP_017156673.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila r
>gi|1036756505|ref|XP_017156674.1| PREDICTED: proteasome subunit beta type-3-like [Drosophila r
>gi|1036738091|ref|XP_017154268.1| PREDICTED: sodium channel protein para isoform X1 [Drosophil
>gi|1036738109|ref|XP_017154276.1| PREDICTED: sodium channel protein para isoform X2 [Drosophil
>gi|1036738359|ref|XP_017154388.1| PREDICTED: sodium channel protein para isoform X15 [Drosoph:
>gi|1036738197|ref|XP_017154320.1| PREDICTED: sodium channel protein para isoform X7 [Drosophil
>gi|1036738461|ref|XP_017154434.1| PREDICTED: sodium channel protein para isoform X21 [Drosoph:
>gi|1036738127|ref|XP_017154285.1| PREDICTED: sodium channel protein para isoform X3 [Drosophil
>gi|1036738145|ref|XP_017154294.1| PREDICTED: sodium channel protein para isoform X4 [Drosophil
>gi|1036738429|ref|XP_017154417.1| PREDICTED: sodium channel protein para isoform X19 [Drosoph:
>gi|1036738262|ref|XP_017154357.1| PREDICTED: sodium channel protein para isoform X11 [Drosoph:
>gi|1036738279|ref|XP_017154364.1| PREDICTED: sodium channel protein para isoform X12 [Drosoph:
>gi|1036738325|ref|XP_017154373.1| PREDICTED: sodium channel protein para isoform X13 [Drosoph:
>gi|1036738507|ref|XP_017154459.1| PREDICTED: sodium channel protein para isoform X24 [Drosoph:
>gi|1036738375|ref|XP_017154396.1| PREDICTED: sodium channel protein para isoform X16 [Drosoph:
>gi|1036738559|ref|XP_017154484.1| PREDICTED: sodium channel protein para isoform X27 [Drosoph:
>gi|1036738343|ref|XP_017154380.1| PREDICTED: sodium channel protein para isoform X14 [Drosoph:
>gi|1036738163|ref|XP_017154304.1| PREDICTED: sodium channel protein para isoform X5 [Drosophil
>gi|1036738230|ref|XP_017154337.1| PREDICTED: sodium channel protein para isoform X9 [Drosophil
>gi|1036738212|ref|XP_017154329.1| PREDICTED: sodium channel protein para isoform X8 [Drosophil
>gi|1036738475|ref|XP_017154442.1| PREDICTED: sodium channel protein para isoform X22 [Drosoph:
>gi|1036738525|ref|XP_017154468.1| PREDICTED: sodium channel protein para isoform X25 [Drosoph:
>gi|1036738247|ref|XP_017154348.1| PREDICTED: sodium channel protein para isoform X10 [Drosoph:
>gi|1036738491|ref|XP_017154451.1| PREDICTED: sodium channel protein para isoform X23 [Drosoph:
>gi|1036738411|ref|XP_017154411.1| PREDICTED: sodium channel protein para isoform X18 [Drosoph:
>gi|1036738393|ref|XP_017154404.1| PREDICTED: sodium channel protein para isoform X17 [Drosoph:
>gi|1036738592|ref|XP_017154499.1| PREDICTED: sodium channel protein para isoform X29 [Drosoph:
>gi|1036738576|ref|XP_017154491.1| PREDICTED: sodium channel protein para isoform X28 [Drosoph:
>gi|1036738610|ref|XP_017154508.1| PREDICTED: sodium channel protein para isoform X30 [Drosoph:
>gi|1036738179|ref|XP_017154312.1| PREDICTED: sodium channel protein para isoform X6 [Drosophil
>gi|1036738646|ref|XP_017154525.1| PREDICTED: sodium channel protein para isoform X32 [Drosoph:
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>gi|1036738628|ref|XP_017154515.1| PREDICTED: sodium channel protein para isoform X31 [Drosophi
>gi|1036738543|ref|XP_017154476.1| PREDICTED: sodium channel protein para isoform X26 [Drosophi
>gi|1036738447|ref|XP_017154426.1| PREDICTED: sodium channel protein para isoform X20 [Drosophi
>gi|1036738666|ref|XP_017154530.1| PREDICTED: sodium channel protein para isoform X33 [Drosophi
>gi|1036706974|ref|XP_017136670.1| PREDICTED: eukaryotic translation initiation factor 4H isofo
>gi|1036706934|ref|XP_017136649.1| PREDICTED: eukaryotic translation initiation factor 4H isofo
>gi|1036706955|ref|XP_017136659.1| PREDICTED: eukaryotic translation initiation factor 4H isofo
>gi|1036707024|ref|XP_017136699.1| PREDICTED: eukaryotic translation initiation factor 2 subuni
>gi|1036706875|ref|XP_017136615.1| PREDICTED: protein DDI1 homolog 2 [Drosophila miranda]
>gi|1036706991|ref|XP_017136682.1| PREDICTED: cell cycle control protein 50A [Drosophila mirand
>gi|1036706825|ref|XP_017136586.1| PREDICTED: zinc finger protein hangover isoform X2 [Drosophi
>gi|1036706808|ref|XP_017136578.1| PREDICTED: zinc finger protein hangover isoform X1 [Drosophi
>gi|1036706859|ref|XP_017136604.1| PREDICTED: zinc finger protein hangover isoform X4 [Drosophi
>gi|1036706842|ref|XP_017136596.1| PREDICTED: zinc finger protein hangover isoform X3 [Drosophi
>gi|1036706894|ref|XP_017136629.1| PREDICTED: regucalcin isoform X1 [Drosophila miranda]
>gi|1036706911|ref|XP_017136637.1| PREDICTED: regucalcin isoform X2 [Drosophila miranda]
>gi|1036707075|ref|XP_017136735.1| PREDICTED: chromatin accessibility complex protein 1 [Droso
>gi|1036707041|ref|XP_017136712.1| PREDICTED: protein LSM12 homolog A isoform X1 [Drosophila m
>gi|1036707058|ref|XP_017136721.1| PREDICTED: protein LSM12 homolog A isoform X2 [Drosophila m
>gi|1036710860|ref|XP_017139156.1| PREDICTED: pleiotropic regulator 1 [Drosophila miranda]
>gi|1036710844|ref|XP_017139146.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 7 [Drosophi
>gi|1036710877|ref|XP_017139168.1| PREDICTED: uncharacterized protein LOC108153594 [Drosophila
>gi|1036710928|ref|XP_017139206.1| PREDICTED: rhythmically expressed gene 2 protein [Drosophila
>gi|1036710894|ref|XP_017139181.1| PREDICTED: 26S protease regulatory subunit 10B [Drosophila
>gi|1036710911|ref|XP_017139193.1| PREDICTED: tRNA (guanine-N(7)-)-methyltransferase non-cataly
>gi|1036710232|ref|XP_017138730.1| PREDICTED: kinesin-like protein Klp10A [Drosophila miranda]
>gi|1036710253|ref|XP_017138737.1| PREDICTED: kinesin-like protein Klp10A [Drosophila miranda]
>gi|1036710287|ref|XP_017138765.1| PREDICTED: uncharacterized protein LOC108153337 isoform X1
>gi|1036710309|ref|XP_017138776.1| PREDICTED: homeotic protein ocelliless isoform X2 [Drosophi
>gi|1036710412|ref|XP_017138847.1| PREDICTED: venom toxin OcyC11 [Drosophila miranda]
>gi|1036710393|ref|XP_017138836.1| PREDICTED: uncharacterized protein LOC108153373 [Drosophila
>gi|1036710376|ref|XP_017138824.1| PREDICTED: uncharacterized protein LOC108153357 isoform X2
>gi|1036710359|ref|XP_017138816.1| PREDICTED: uncharacterized protein LOC108153357 isoform X1
>gi|1036710342|ref|XP_017138805.1| PREDICTED: uncharacterized protein LOC108153349 [Drosophila
>gi|1036710326|ref|XP_017138791.1| PREDICTED: GTP-binding nuclear protein Ran [Drosophila mira
>gi|1036710270|ref|XP_017138751.1| PREDICTED: probable G-protein coupled receptor Mth-like 1 [
>gi|1036741020|ref|XP_017155802.1| PREDICTED: DNA topoisomerase 3-beta [Drosophila miranda]
>gi|1036719809|ref|XP_017144302.1| PREDICTED: ubiquitin [Drosophila miranda]
>gi|1036719789|ref|XP_017144290.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036719769|ref|XP_017144277.1| PREDICTED: zinc finger SWIM domain-containing protein 8 [Dro
>gi|1036699011|ref|XP_017137275.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1036723238|ref|XP_017145878.1| PREDICTED: CAD protein [Drosophila miranda]
>gi|1036723248|ref|XP_017145886.1| PREDICTED: uncharacterized protein LOC108158176 [Drosophila
>gi|1036699026|ref|XP_017137403.1| PREDICTED: LOW QUALITY PROTEIN: protein BUD31 homolog [Droso
>gi|1036723373|ref|XP_017145935.1| PREDICTED: uncharacterized protein LOC108158217 [Drosophila
>gi|1036723354|ref|XP_017145922.1| PREDICTED: protein sisterless A [Drosophila miranda]
>gi|1036699039|ref|XP_017137528.1| PREDICTED: uncharacterized protein LOC108152566 [Drosophila
>gi|1036723390|ref|XP_017145948.1| PREDICTED: uncharacterized protein LOC108158224 [Drosophila
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>gi|1036723337|ref|XP_017145911.1| PREDICTED: protein naked cuticle [Drosophila miranda]
>gi|1036707179|ref|XP_017136808.1| PREDICTED: uncharacterized protein LOC108152181 isoform X1
>gi|1036707200|ref|XP_017136817.1| PREDICTED: uncharacterized protein LOC108152181 isoform X2
>gi|1036704367|ref|XP_017134846.1| PREDICTED: uncharacterized protein LOC108151003 [Drosophila
>gi|1036704030|ref|XP_017134646.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036725562|ref|XP_017147235.1| PREDICTED: cytoplasmic dynein 1 light intermediate chain 1
>gi|1036725581|ref|XP_017147245.1| PREDICTED: cytoplasmic dynein 1 light intermediate chain 1
>gi|1036725683|ref|XP_017147298.1| PREDICTED: armadillo repeat-containing protein 2 [Drosophila
>gi|1036725661|ref|XP_017147286.1| PREDICTED: uncharacterized protein LOC108158962 [Drosophila
>gi|1036725600|ref|XP_017147257.1| PREDICTED: uncharacterized protein LOC108158945 isoform X1
>gi|1036725619|ref|XP_017147266.1| PREDICTED: uncharacterized protein LOC108158945 isoform X2
>gi|1036725638|ref|XP_017147273.1| PREDICTED: uncharacterized protein LOC108158945 isoform X3
>gi|1036725545|ref|XP_017147223.1| PREDICTED: uncharacterized protein LOC108158926 [Drosophila
>gi|1036735206|ref|XP_017152647.1| PREDICTED: bleomycin hydrolase [Drosophila miranda]
>gi|1036735225|ref|XP_017152659.1| PREDICTED: uncharacterized protein F58A4.6 [Drosophila mira
>gi|1036735244|ref|XP_017152667.1| PREDICTED: uncharacterized protein F58A4.6 [Drosophila mira
>gi|1036705294|ref|XP_017135507.1| PREDICTED: uncharacterized protein LOC108151424 [Drosophila
>gi|1036754095|ref|XP_017156537.1| PREDICTED: T-box transcription factor TBX1 [Drosophila mira
>gi|1036704783|ref|XP_017135141.1| PREDICTED: uncharacterized protein LOC108151194 [Drosophila
>gi|1036754656|ref|XP_017156570.1| PREDICTED: uncharacterized protein LOC108165043 [Drosophila
>gi|1036703385|ref|XP_017156917.1| PREDICTED: very low-density lipoprotein receptor-like [Droso
>gi|1036703400|ref|XP_017156925.1| PREDICTED: very low-density lipoprotein receptor-like [Droso
>gi|1036744849|ref|XP_017156011.1| PREDICTED: protein DGCR14 homolog [Drosophila miranda]
>gi|1036744868|ref|XP_017156012.1| PREDICTED: sepiapterin reductase [Drosophila miranda]
>gi|1036744784|ref|XP_017156009.1| PREDICTED: sepiapterin reductase [Drosophila miranda]
>gi|1036744748|ref|XP_017156007.1| PREDICTED: uncharacterized protein LOC108164665 isoform X1
>gi|1036744766|ref|XP_017156008.1| PREDICTED: uncharacterized protein LOC108164665 isoform X2
>gi|1036744731|ref|XP_017156006.1| PREDICTED: uncharacterized protein LOC108164663 isoform X2
>gi|1036744713|ref|XP_017156005.1| PREDICTED: uncharacterized protein LOC108164663 isoform X1
>gi|1036704836|ref|XP_017135182.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036704247|ref|XP_017134774.1| PREDICTED: chorion protein S36 [Drosophila miranda]
>gi|1036757301|ref|XP_017156717.1| PREDICTED: chorion protein S38 [Drosophila miranda]
>gi|1036726169|ref|XP_017147613.1| PREDICTED: protein ovarian tumor locus isoform X2 [Drosophil
>gi|1036726150|ref|XP_017147604.1| PREDICTED: protein ovarian tumor locus isoform X1 [Drosophil
>gi|1036726189|ref|XP_017147621.1| PREDICTED: protein ovarian tumor locus isoform X1 [Drosophil
>gi|1036726280|ref|XP_017147668.1| PREDICTED: solute carrier family 41 member 1 isoform X5 [Dro
>gi|1036726257|ref|XP_017147658.1| PREDICTED: solute carrier family 41 member 1 isoform X4 [Dro
>gi|1036726241|ref|XP_017147648.1| PREDICTED: solute carrier family 41 member 2 isoform X3 [Dro
>gi|1036726204|ref|XP_017147632.1| PREDICTED: solute carrier family 41 member 1 isoform X1 [Dro
>gi|1036726222|ref|XP_017147640.1| PREDICTED: solute carrier family 41 member 1 isoform X2 [Dro
>gi|1036734410|ref|XP_017152186.1| PREDICTED: kinetochore protein NDC80 homolog [Drosophila mi
>gi|1036734394|ref|XP_017152174.1| PREDICTED: late secretory pathway protein AVL9 homolog [Dros
>gi|1036750613|ref|XP_017156329.1| PREDICTED: troponin T, skeletal muscle isoform X2 [Drosophil
>gi|1036750594|ref|XP_017156328.1| PREDICTED: troponin T, skeletal muscle isoform X1 [Drosophil
>gi|1036750665|ref|XP_017156332.1| PREDICTED: troponin T, skeletal muscle isoform X5 [Drosophil
>gi|1036750629|ref|XP_017156330.1| PREDICTED: troponin T, skeletal muscle isoform X3 [Drosophil
>gi|1036750832|ref|XP_017156343.1| PREDICTED: troponin T, skeletal muscle isoform X14 [Drosoph
>gi|1036750648|ref|XP_017156331.1| PREDICTED: troponin T, skeletal muscle isoform X4 [Drosophil
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>gi|1036750812|ref|XP_017156342.1| PREDICTED: troponin T, skeletal muscle isoform X13 [Drosophi
>gi|1036750760|ref|XP_017156339.1| PREDICTED: troponin T, skeletal muscle isoform X10 [Drosophi
>gi|1036750741|ref|XP_017156338.1| PREDICTED: troponin T, skeletal muscle isoform X9 [Drosophil
>gi|1036750795|ref|XP_017156341.1| PREDICTED: troponin T, skeletal muscle isoform X12 [Drosophi
>gi|1036750777|ref|XP_017156340.1| PREDICTED: troponin T, skeletal muscle isoform X11 [Drosophi
>gi|1036750684|ref|XP_017156333.1| PREDICTED: troponin T, skeletal muscle isoform X6 [Drosophil
>gi|1036750703|ref|XP_017156334.1| PREDICTED: troponin T, skeletal muscle isoform X7 [Drosophil
>gi|1036750722|ref|XP_017156335.1| PREDICTED: troponin T, skeletal muscle isoform X8 [Drosophil
>gi|1036727090|ref|XP_017148180.1| PREDICTED: protein-tyrosine sulfotransferase isoform X1 [Dro
>gi|1036727109|ref|XP_017148188.1| PREDICTED: protein-tyrosine sulfotransferase isoform X1 [Dro
>gi|1036727126|ref|XP_017148197.1| PREDICTED: protein-tyrosine sulfotransferase isoform X2 [Dro
>gi|1036703433|ref|XP_017156947.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036703450|ref|XP_017156955.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036703463|ref|XP_017156961.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036707302|ref|XP_017136877.1| PREDICTED: uncharacterized protein LOC108152215 [Drosophila
>gi|1036706709|ref|XP_017136517.1| PREDICTED: uncharacterized protein LOC108152033 [Drosophila
>gi|1036705173|ref|XP_017135429.1| PREDICTED: uncharacterized protein LOC108151374 [Drosophila
>gi|1036705190|ref|XP_017135436.1| PREDICTED: uncharacterized protein LOC108151374 [Drosophila
>gi|1036706404|ref|XP_017136278.1| PREDICTED: larval cuticle protein LCP-14 [Drosophila miranda
>gi|1036705689|ref|XP_017135760.1| PREDICTED: uncharacterized protein LOC108151588 [Drosophila
>gi|1036707234|ref|XP_017136842.1| PREDICTED: uncharacterized protein LOC108152195 [Drosophila
>gi|1036737335|ref|XP_017153863.1| PREDICTED: uncharacterized protein LOC108163215 [Drosophila
>gi|1036737350|ref|XP_017153874.1| PREDICTED: mucin-2-like [Drosophila miranda]
>gi|1036757320|ref|XP_017156718.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036757339|ref|XP_017156719.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036757354|ref|XP_017156720.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036752846|ref|XP_017156461.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036752865|ref|XP_017156462.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036699054|ref|XP_017137653.1| PREDICTED: proteasome subunit alpha type-2-like, partial [Di
>gi|1036725774|ref|XP_017147365.1| PREDICTED: protein LST8 homolog [Drosophila miranda]
>gi|1036725755|ref|XP_017147351.1| PREDICTED: ADP-ribosylation factor-binding protein GGA2 [Dro
>gi|1036725793|ref|XP_017147377.1| PREDICTED: 40S ribosomal protein S5a [Drosophila miranda]
>gi|1036699069|ref|XP_017137778.1| PREDICTED: uncharacterized protein LOC108152715 [Drosophila
>gi|1036742141|ref|XP_017155867.1| PREDICTED: uncharacterized protein LOC108164561 isoform X2
>gi|1036742123|ref|XP_017155866.1| PREDICTED: uncharacterized protein LOC108164561 isoform X1
>gi|1036752972|ref|XP_017156470.1| PREDICTED: nuclear transcription factor Y subunit gamma-like
>gi|1036752955|ref|XP_017156469.1| PREDICTED: uncharacterized protein LOC108164968 [Drosophila
>gi|1036731537|ref|XP_017150762.1| PREDICTED: uncharacterized protein LOC108160971 isoform X1
>gi|1036731556|ref|XP_017150770.1| PREDICTED: sterol 3-beta-glucosyltransferase isoform X2 [Dro
>gi|1036731573|ref|XP_017150778.1| PREDICTED: pleckstrin homology-like domain family B member 2
>gi|1036745632|ref|XP_017156059.1| PREDICTED: coatomer subunit delta [Drosophila miranda]
>gi|1036712871|ref|XP_017140293.1| PREDICTED: LOW QUALITY PROTEIN: acidic repeat-containing pro
>gi|1036712972|ref|XP_017140349.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036712888|ref|XP_017140304.1| PREDICTED: peroxisomal targeting signal 1 receptor [Drosophi
>gi|1036712853|ref|XP_017140283.1| PREDICTED: protein MMS22-like [Drosophila miranda]
>gi|1036712955|ref|XP_017140341.1| PREDICTED: serine/threonine-protein phosphatase Pgam5, mito
>gi|1036712938|ref|XP_017140333.1| PREDICTED: serine/threonine-protein phosphatase Pgam5, mito
>gi|1036712905|ref|XP_017140315.1| PREDICTED: calphotin isoform X1 [Drosophila miranda]
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>gi|1036712922|ref|XP_017140323.1| PREDICTED: GATA type zinc finger protein asd-4 isoform X2 [
>gi|1036748739|ref|XP_017156228.1| PREDICTED: cytochrome P450 4d1 isoform X2 [Drosophila miran
>gi|1036748720|ref|XP_017156227.1| PREDICTED: cytochrome P450 4d1 isoform X1 [Drosophila miran
>gi|1036748758|ref|XP_017156229.1| PREDICTED: uncharacterized protein LOC108164819 [Drosophila
>gi|1036748815|ref|XP_017156232.1| PREDICTED: E3 ubiquitin-protein ligase RNF125-like [Drosoph
>gi|1036748777|ref|XP_017156230.1| PREDICTED: uncharacterized protein LOC108164820 isoform X1
>gi|1036748796|ref|XP_017156231.1| PREDICTED: uncharacterized protein LOC108164820 isoform X2
>gi|1036723794|ref|XP_017146152.1| PREDICTED: open rectifier potassium channel protein 1 isofo
>gi|1036723776|ref|XP_017146144.1| PREDICTED: open rectifier potassium channel protein 1 isofo
>gi|1036723813|ref|XP_017146163.1| PREDICTED: open rectifier potassium channel protein 1 isofo
>gi|1036723758|ref|XP_017146133.1| PREDICTED: putative ATP-dependent RNA helicase DHX57 [Droso
>gi|1036723830|ref|XP_017146173.1| PREDICTED: uncharacterized protein LOC108158413 [Drosophila
>gi|1036705723|ref|XP_017135784.1| PREDICTED: uncharacterized protein LOC108151605 [Drosophila
>gi|1036706264|ref|XP_017136184.1| PREDICTED: E3 ubiquitin-protein ligase RNF125-like [Drosoph
>gi|1036707319|ref|XP_017136890.1| PREDICTED: cytochrome P450 4d1-like [Drosophila miranda]
>gi|1036745068|ref|XP_017156025.1| PREDICTED: uncharacterized protein LOC108164679 isoform X1
>gi|1036745087|ref|XP_017156026.1| PREDICTED: uncharacterized protein LOC108164679 isoform X1
>gi|1036745106|ref|XP_017156027.1| PREDICTED: uncharacterized protein LOC108164679 isoform X2
>gi|1036745125|ref|XP_017156028.1| PREDICTED: uncharacterized protein LOC108164679 isoform X3
>gi|1036745143|ref|XP_017156029.1| PREDICTED: uncharacterized protein LOC108164679 isoform X4
>gi|1036745809|ref|XP_017156068.1| PREDICTED: methylmalonic aciduria and homocystinuria type D
>gi|1036745827|ref|XP_017156069.1| PREDICTED: methylmalonic aciduria and homocystinuria type D
>gi|1036745846|ref|XP_017156070.1| PREDICTED: COMM domain-containing protein 3 [Drosophila mir
>gi|1036753185|ref|XP_017156483.1| PREDICTED: neuferricin homolog [Drosophila miranda]
>gi|1036726315|ref|XP_017147713.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726334|ref|XP_017147722.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726350|ref|XP_017147730.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726367|ref|XP_017147738.1| PREDICTED: glutamate--cysteine ligase [Drosophila miranda]
>gi|1036726386|ref|XP_017147752.1| PREDICTED: nudC domain-containing protein 1 [Drosophila mir
>gi|1036726404|ref|XP_017147764.1| PREDICTED: transmembrane emp24 domain-containing protein 3
>gi|1036726422|ref|XP_017147774.1| PREDICTED: transmembrane emp24 domain-containing protein 3
>gi|1036699084|ref|XP_017137898.1| PREDICTED: transcription factor SPT20 homolog [Drosophila m
>gi|1036703584|ref|XP_017157032.1| PREDICTED: probable cytochrome P450 28c1 [Drosophila mirand
>gi|1036699098|ref|XP_017138026.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036709050|ref|XP_017138009.1| PREDICTED: breast carcinoma-amplified sequence 3 homolog is
>gi|1036708999|ref|XP_017137984.1| PREDICTED: uncharacterized protein LOC108152864 isoform X1
>gi|1036709016|ref|XP_017137992.1| PREDICTED: uncharacterized protein LOC108152864 isoform X1
>gi|1036709033|ref|XP_017138000.1| PREDICTED: uncharacterized protein LOC108152864 isoform X1
>gi|1036709065|ref|XP_017138021.1| PREDICTED: uncharacterized protein LOC108152890 [Drosophila
>gi|1036709079|ref|XP_017138031.1| PREDICTED: uncharacterized protein LOC108152890 [Drosophila
>gi|1036709100|ref|XP_017138042.1| PREDICTED: heat shock 70 kDa protein cognate 3 [Drosophila
>gi|1036709119|ref|XP_017138051.1| PREDICTED: heat shock 70 kDa protein cognate 3 [Drosophila
>gi|1036709136|ref|XP_017138061.1| PREDICTED: heat shock 70 kDa protein cognate 3 [Drosophila
>gi|1036718613|ref|XP_017143633.1| PREDICTED: casein kinase II subunit beta isoform X1 [Drosop
>gi|1036718631|ref|XP_017143639.1| PREDICTED: casein kinase II subunit beta isoform X1 [Drosop
>gi|1036718647|ref|XP_017143648.1| PREDICTED: casein kinase II subunit beta isoform X2 [Drosop
>gi|1036718576|ref|XP_017143609.1| PREDICTED: testis-expressed sequence 10 protein homolog [Dr
>gi|1036718594|ref|XP_017143621.1| PREDICTED: WD repeat-containing protein 46 [Drosophila mira
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>gi|1036755664|ref|XP_017156625.1| PREDICTED: uncharacterized protein LOC108165085 [Drosophila
>gi|1036710446|ref|XP_017138874.1| PREDICTED: transcriptional repressor CTCF [Drosophila mirand
>gi|1036710429|ref|XP_017138860.1| PREDICTED: dynein regulatory complex protein 1 homolog [Dros
>gi|1036710523|ref|XP_017138934.1| PREDICTED: zinc finger protein 311 isoform X1 [Drosophila mi
>gi|1036710536|ref|XP_017138943.1| PREDICTED: zinc finger protein 311 isoform X2 [Drosophila mi
>gi|1036710480|ref|XP_017138900.1| PREDICTED: mothers against decapentaplegic homolog 3 [Drosop
>gi|1036710553|ref|XP_017138955.1| PREDICTED: uncharacterized protein LOC108153446 [Drosophila
>gi|1036710463|ref|XP_017138887.1| PREDICTED: phenylalanine--tRNA ligase alpha subunit [Drosopl
>gi|1036710496|ref|XP_017138912.1| PREDICTED: zinc finger protein 311 [Drosophila miranda]
>gi|1036710570|ref|XP_017138968.1| PREDICTED: zinc finger protein 32 [Drosophila miranda]
>gi|1036710510|ref|XP_017138923.1| PREDICTED: zinc finger protein 813 [Drosophila miranda]
>gi|1036754365|ref|XP_017156553.1| PREDICTED: guanine nucleotide-binding protein subunit beta-5
>gi|1036751364|ref|XP_017156373.1| PREDICTED: zinc finger protein 32 [Drosophila miranda]
>gi|1036745562|ref|XP_017156055.1| PREDICTED: hepatic triacylglycerol lipase-like [Drosophila r
>gi|1036745544|ref|XP_017156054.1| PREDICTED: TNF receptor-associated factor 6 [Drosophila mira
>gi|1036748332|ref|XP_017156215.1| PREDICTED: acidic phospholipase A2 PA4 [Drosophila miranda]
>gi|1036748314|ref|XP_017156214.1| PREDICTED: uncharacterized protein LOC108164803 [Drosophila
>gi|1036699113|ref|XP_017138156.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 [Drosophila mi
>gi|1036699128|ref|XP_017138281.1| PREDICTED: LOW QUALITY PROTEIN: odorant receptor 7a [Drosopl
>gi|1036751974|ref|XP_017156407.1| PREDICTED: uncharacterized protein LOC108164930 [Drosophila
>gi|1036735669|ref|XP_017152900.1| PREDICTED: mitogen-activated protein kinase kinase kinase [I
>gi|1036727799|ref|XP_017148607.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036727834|ref|XP_017148623.1| PREDICTED: uncharacterized protein LOC108159648 isoform X3
>gi|1036727899|ref|XP_017148657.1| PREDICTED: uncharacterized protein LOC108159648 isoform X5
>gi|1036727848|ref|XP_017148632.1| PREDICTED: uncharacterized protein LOC108159648 isoform X4
>gi|1036727864|ref|XP_017148640.1| PREDICTED: uncharacterized protein LOC108159648 isoform X4
>gi|1036727880|ref|XP_017148648.1| PREDICTED: uncharacterized protein LOC108159648 isoform X4
>gi|1036727917|ref|XP_017148666.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X6 [Drosophi
>gi|1036727815|ref|XP_017148616.1| PREDICTED: uncharacterized protein LOC108159648 isoform X2
>gi|1036727936|ref|XP_017148676.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X7 [Drosophi
>gi|1036727955|ref|XP_017148684.1| PREDICTED: MAGUK p55 subfamily member 7 isoform X7 [Drosophi
>gi|1036756326|ref|XP_017156663.1| PREDICTED: rRNA 2'-O-methyltransferase fibrillarin [Drosophi
>gi|1036699142|ref|XP_017138396.1| PREDICTED: glycine-rich protein DOT1 [Drosophila miranda]
>gi|1036736056|ref|XP_017153110.1| PREDICTED: DNA-binding protein Ewg isoform X1 [Drosophila mi
>gi|1036736097|ref|XP_017153121.1| PREDICTED: DNA-binding protein Ewg isoform X3 [Drosophila mi
>gi|1036736079|ref|XP_017153113.1| PREDICTED: DNA-binding protein Ewg isoform X2 [Drosophila mi
>gi|1036754185|ref|XP_017156543.1| PREDICTED: ras-related protein Rap-2b [Drosophila miranda]
>gi|1036754202|ref|XP_017156544.1| PREDICTED: uncharacterized protein LOC108165017 [Drosophila
>gi|1036706676|ref|XP_017136483.1| PREDICTED: uncharacterized protein LOC108152014 [Drosophila
>gi|1036757081|ref|XP_017156704.1| PREDICTED: uncharacterized protein LOC108165148 [Drosophila
>gi|1036699157|ref|XP_017138516.1| PREDICTED: odorant receptor 1a [Drosophila miranda]
>gi|1036755921|ref|XP_017156640.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036755940|ref|XP_017156641.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036755959|ref|XP_017156642.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036755978|ref|XP_017156643.1| PREDICTED: homeobox protein 12 [Drosophila miranda]
>gi|1036753933|ref|XP_017156528.1| PREDICTED: nuclear pore complex protein DDB_G0274915 [Drosop
>gi|1036699171|ref|XP_017138641.1| PREDICTED: uncharacterized protein LOC108153252 [Drosophila
>gi|1036699186|ref|XP_017138770.1| PREDICTED: uncharacterized protein LOC108153321 [Drosophila
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>gi|1036757116|ref|XP_017156706.1| PREDICTED: protein yellow [Drosophila miranda]
>gi|1036702374|ref|XP_017156337.1| PREDICTED: achaete-scute complex protein T5 [Drosophila mira
>gi|1036702357|ref|XP_017156322.1| PREDICTED: achaete-scute complex protein T4 [Drosophila mira
>gi|1036702271|ref|XP_017156257.1| PREDICTED: achaete-scute complex protein T3 [Drosophila mira
>gi|1036757246|ref|XP_017156714.1| PREDICTED: pepsin-2B [Drosophila miranda]
>gi|1036754816|ref|XP_017156580.1| PREDICTED: achaete-scute complex protein T8 [Drosophila mira
>gi|1036727688|ref|XP_017148548.1| PREDICTED: cytochrome P450 4g1 [Drosophila miranda]
>gi|1036727670|ref|XP_017148537.1| PREDICTED: exportin-6-A [Drosophila miranda]
>gi|1036743242|ref|XP_017155933.1| PREDICTED: apoptosis regulatory protein Siva isoform X2 [Dro
>gi|1036743224|ref|XP_017155932.1| PREDICTED: apoptosis regulatory protein Siva isoform X1 [Dro
>gi|1036743277|ref|XP_017155934.1| PREDICTED: uncharacterized protein LOC108164606 [Drosophila
>gi|1036738775|ref|XP_017154589.1| PREDICTED: ATP-binding cassette sub-family B member 10, mito
>gi|1036738787|ref|XP_017154598.1| PREDICTED: probable methylmalonate-semialdehyde dehydrogenas
>gi|1036704384|ref|XP_017134859.1| PREDICTED: uncharacterized protein LOC108151009 [Drosophila
>gi|1036726558|ref|XP_017147870.1| PREDICTED: carboxypeptidase D isoform X1 [Drosophila miranda
>gi|1036726577|ref|XP_017147878.1| PREDICTED: carboxypeptidase D isoform X2 [Drosophila miranda
>gi|1036722294|ref|XP_017145462.1| PREDICTED: arginase-1 [Drosophila miranda]
>gi|1036722311|ref|XP_017145474.1| PREDICTED: protein elav [Drosophila miranda]
>gi|1036722277|ref|XP_017145450.1| PREDICTED: endoplasmic reticulum-Golgi intermediate compartm
>gi|1036740402|ref|XP_017155476.1| PREDICTED: chromodomain-helicase-DNA-binding protein 1 isofc
>gi|1036740420|ref|XP_017155480.1| PREDICTED: chromodomain-helicase-DNA-binding protein 1 isofc
>gi|1036740438|ref|XP_017155495.1| PREDICTED: NADH dehydrogenase [ubiquinone] flavoprotein 2, r
>gi|1036754256|ref|XP_017156547.1| PREDICTED: uncharacterized protein LOC108165021 [Drosophila
>gi|1036749413|ref|XP_017156264.1| PREDICTED: uncharacterized protein LOC108164846 [Drosophila
>gi|1036742717|ref|XP_017155901.1| PREDICTED: transmembrane protein 41 homolog [Drosophila mira
>gi|1036740314|ref|XP_017155458.1| PREDICTED: sn1-specific diacylglycerol lipase alpha isoform
>gi|1036740298|ref|XP_017155448.1| PREDICTED: sn1-specific diacylglycerol lipase alpha isoform
>gi|1036740332|ref|XP_017155469.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036732353|ref|XP_017151106.1| PREDICTED: regulator of nonsense transcripts 1 homolog [Dros
>gi|1036744658|ref|XP_017156002.1| PREDICTED: poly(A) RNA polymerase gld-2 homolog B [Drosophil
>gi|1036699201|ref|XP_017138899.1| PREDICTED: NADH dehydrogenase (ubiquinone) complex I, assemb
>gi|1036732299|ref|XP_017151071.1| PREDICTED: glycine-rich selenoprotein [Drosophila miranda]
>gi|1036732220|ref|XP_017151039.1| PREDICTED: anoctamin-6 isoform X1 [Drosophila miranda]
>gi|1036732243|ref|XP_017151048.1| PREDICTED: anoctamin-6 isoform X1 [Drosophila miranda]
>gi|1036732281|ref|XP_017151065.1| PREDICTED: anoctamin-6 isoform X3 [Drosophila miranda]
>gi|1036732264|ref|XP_017151057.1| PREDICTED: anoctamin-6 isoform X2 [Drosophila miranda]
>gi|1036752810|ref|XP_017156458.1| PREDICTED: 4-nitrophenylphosphatase [Drosophila miranda]
>gi|1036752791|ref|XP_017156457.1| PREDICTED: pyridoxal phosphate phosphatase [Drosophila mirar
>gi|1036752482|ref|XP_017156439.1| PREDICTED: loss of heterozygosity 12 chromosomal region 1 pr
>gi|1036751699|ref|XP_017156390.1| PREDICTED: AH receptor-interacting protein [Drosophila mirar
>gi|1036747387|ref|XP_017156165.1| PREDICTED: SH3 domain-binding protein 5-like isoform X2 [Dro
>gi|1036747368|ref|XP_017156164.1| PREDICTED: keratin, type II cytoskeletal 1 isoform X1 [Droso
>gi|1036732370|ref|XP_017151115.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732388|ref|XP_017151121.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732407|ref|XP_017151128.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732447|ref|XP_017151142.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732483|ref|XP_017151158.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732426|ref|XP_017151134.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
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>gi|1036732464|ref|XP_017151149.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036732500|ref|XP_017151166.1| PREDICTED: Ca(2+)/calmodulin-responsive adenylate cyclase is
>gi|1036705277|ref|XP_017135496.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036703079|ref|XP_017156747.1| PREDICTED: venom allergen 5-like [Drosophila miranda]
>gi|1036703214|ref|XP_017156818.1| PREDICTED: venom allergen 5-like [Drosophila miranda]
>gi|1036751106|ref|XP_017156359.1| PREDICTED: solute carrier family 46 member 3 [Drosophila mi
>gi|1036735975|ref|XP_017153058.1| PREDICTED: netrin-B [Drosophila miranda]
>gi|1036703251|ref|XP_017156840.1| PREDICTED: gamma-butyrobetaine dioxygenase [Drosophila miran
>gi|1036736208|ref|XP_017153196.1| PREDICTED: netrin-A isoform X1 [Drosophila miranda]
>gi|1036736226|ref|XP_017153204.1| PREDICTED: netrin-A isoform X2 [Drosophila miranda]
>gi|1036741231|ref|XP_017155816.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X2
>gi|1036741282|ref|XP_017155819.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X5
>gi|1036741213|ref|XP_017155815.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X1
>gi|1036741248|ref|XP_017155817.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X3
>gi|1036741265|ref|XP_017155818.1| PREDICTED: b(0,+)-type amino acid transporter 1 isoform X4
>gi|1036741343|ref|XP_017155823.1| PREDICTED: nucleoside diphosphate kinase 6 [Drosophila miran
>gi|1036741326|ref|XP_017155822.1| PREDICTED: probable 28S ribosomal protein S25, mitochondrial
>gi|1036741299|ref|XP_017155820.1| PREDICTED: probable RNA-binding protein 18 isoform X1 [Droso
>gi|1036741313|ref|XP_017155821.1| PREDICTED: probable RNA-binding protein 18 isoform X2 [Droso
>gi|1036706228|ref|XP_017136158.1| PREDICTED: uncharacterized protein LOC108151811 [Drosophila
>gi|1036699216|ref|XP_017139141.1| PREDICTED: uncharacterized protein LOC108153545 [Drosophila
>gi|1036758052|ref|XP_017156753.1| PREDICTED: atrophin-1 [Drosophila miranda]
>gi|1036746085|ref|XP_017156085.1| PREDICTED: titin homolog isoform X2 [Drosophila miranda]
>gi|1036746028|ref|XP_017156081.1| PREDICTED: titin homolog isoform X1 [Drosophila miranda]
>gi|1036746047|ref|XP_017156082.1| PREDICTED: titin homolog isoform X1 [Drosophila miranda]
>gi|1036746066|ref|XP_017156083.1| PREDICTED: titin homolog isoform X1 [Drosophila miranda]
>gi|1036746103|ref|XP_017156086.1| PREDICTED: protein TsetseEP isoform X1 [Drosophila miranda]
>gi|1036746139|ref|XP_017156088.1| PREDICTED: transcriptional regulatory protein AlgP isoform
>gi|1036746120|ref|XP_017156087.1| PREDICTED: protein TsetseEP isoform X2 [Drosophila miranda]
>gi|1036745996|ref|XP_017156079.1| PREDICTED: uncharacterized protein LOC108164713 [Drosophila
>gi|1036746011|ref|XP_017156080.1| PREDICTED: uncharacterized protein LOC108164713 [Drosophila
>gi|1036703967|ref|XP_017134609.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036699231|ref|XP_017139260.1| PREDICTED: uncharacterized protein LOC108153628 [Drosophila
>gi|1036732853|ref|XP_017151354.1| PREDICTED: transcription factor TBF1-like [Drosophila miranc
>gi|1036732871|ref|XP_017151361.1| PREDICTED: transcription factor TBF1-like [Drosophila miranc
>gi|1036712798|ref|XP_017140241.1| PREDICTED: sodium leak channel non-selective protein isoform
>gi|1036712764|ref|XP_017140226.1| PREDICTED: sodium leak channel non-selective protein isoform
>gi|1036712781|ref|XP_017140234.1| PREDICTED: sodium leak channel non-selective protein isoform
>gi|1036712832|ref|XP_017140273.1| PREDICTED: plasminogen receptor (KT) [Drosophila miranda]
>gi|1036712815|ref|XP_017140258.1| PREDICTED: tyrosine aminotransferase [Drosophila miranda]
>gi|1036712747|ref|XP_017140216.1| PREDICTED: golgin subfamily B member 1 [Drosophila miranda]
>gi|1036751473|ref|XP_017156379.1| PREDICTED: uncharacterized protein LOC108164910 isoform X1
>gi|1036751492|ref|XP_017156380.1| PREDICTED: uncharacterized protein LOC108164910 isoform X2
>gi|1036751511|ref|XP_017156382.1| PREDICTED: uncharacterized protein LOC108164910 isoform X2
>gi|1036714882|ref|XP_017141522.1| PREDICTED: mRNA-capping enzyme [Drosophila miranda]
>gi|1036714937|ref|XP_017141563.1| PREDICTED: tRNA-dihydrouridine(20) synthase [NAD(P)+]-like
>gi|1036714950|ref|XP_017141577.1| PREDICTED: vacuolar protein sorting-associated protein 4B [[
>gi|1036714965|ref|XP_017141587.1| PREDICTED: vacuolar protein sorting-associated protein 4B [[
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>gi|1036714981|ref|XP_017141595.1| PREDICTED: vacuolar protein sorting-associated protein 4B [
>gi|1036714920|ref|XP_017141550.1| PREDICTED: uncharacterized protein LOC108155323 [Drosophila
>gi|1036714903|ref|XP_017141535.1| PREDICTED: serine/threonine-protein kinase ATR [Drosophila
>gi|1036734889|ref|XP_017152458.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734907|ref|XP_017152466.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734761|ref|XP_017152404.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734923|ref|XP_017152473.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734871|ref|XP_017152450.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734852|ref|XP_017152441.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734816|ref|XP_017152426.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734833|ref|XP_017152434.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734960|ref|XP_017152490.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734780|ref|XP_017152408.1| PREDICTED: uncharacterized protein LOC108162264 isoform X2
>gi|1036734944|ref|XP_017152482.1| PREDICTED: potassium voltage-gated channel protein Shaker is
>gi|1036734797|ref|XP_017152419.1| PREDICTED: dual specificity protein kinase splA isoform X3
>gi|1036734979|ref|XP_017152503.1| PREDICTED: uncharacterized protein LOC108162326 isoform X1
>gi|1036734998|ref|XP_017152510.1| PREDICTED: uncharacterized protein LOC108162326 isoform X2
>gi|1036699246|ref|XP_017139376.1| PREDICTED: axonemal 84 kDa protein [Drosophila miranda]
>gi|1036718432|ref|XP_017143530.1| PREDICTED: dynein heavy chain 6, axonemal [Drosophila miranc
>gi|1036718451|ref|XP_017143542.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036718474|ref|XP_017143551.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036718518|ref|XP_017143576.1| PREDICTED: transcriptional adapter 3 [Drosophila miranda]
>gi|1036718556|ref|XP_017143600.1| PREDICTED: 3-hydroxyacyl-CoA dehydrogenase type-2 [Drosophil
>gi|1036718495|ref|XP_017143563.1| PREDICTED: zinc finger protein 132-like [Drosophila miranda]
>gi|1036718536|ref|XP_017143590.1| PREDICTED: rho GTPase-activating protein 15 [Drosophila mira
>gi|1036735017|ref|XP_017152522.1| PREDICTED: multivesicular body subunit 12A isoform X1 [Dros
>gi|1036735033|ref|XP_017152531.1| PREDICTED: multivesicular body subunit 12A isoform X2 [Dros
>gi|1036735051|ref|XP_017152545.1| PREDICTED: transcription initiation factor TFIID subunit 8
>gi|1036736366|ref|XP_017153295.1| PREDICTED: amyloid protein-binding protein 2 isoform X1 [Dr
>gi|1036736383|ref|XP_017153303.1| PREDICTED: amyloid protein-binding protein 2 isoform X2 [Dr
>gi|1036748202|ref|XP_017156210.1| PREDICTED: uncharacterized protein LOC108164797 isoform X1
>gi|1036748221|ref|XP_017156211.1| PREDICTED: uncharacterized protein LOC108164797 isoform X2
>gi|1036748240|ref|XP_017156212.1| PREDICTED: type-1 angiotensin II receptor-associated protei
>gi|1036740280|ref|XP_017155437.1| PREDICTED: uncharacterized protein LOC108164297 [Drosophila
>gi|1036740262|ref|XP_017155426.1| PREDICTED: putative inorganic phosphate cotransporter [Dros
>gi|1036724452|ref|XP_017146586.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1036724469|ref|XP_017146594.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1036724486|ref|XP_017146603.1| PREDICTED: neprilysin-2 [Drosophila miranda]
>gi|1036724505|ref|XP_017146615.1| PREDICTED: multiple inositol polyphosphate phosphatase 1 [D
>gi|1036724436|ref|XP_017146576.1| PREDICTED: regulatory-associated protein of mTOR [Drosophil
>gi|1036756986|ref|XP_017156699.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036757005|ref|XP_017156700.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036757024|ref|XP_017156701.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036754997|ref|XP_017156591.1| PREDICTED: protein THEM6 [Drosophila miranda]
>gi|1036754867|ref|XP_017156583.1| PREDICTED: tetraspanin-9 [Drosophila miranda]
>gi|1036754886|ref|XP_017156584.1| PREDICTED: uncharacterized protein LOC108165054 [Drosophila
>gi|1036704420|ref|XP_017134880.1| PREDICTED: uncharacterized protein LOC108151028 [Drosophila
>gi|1036751718|ref|XP_017156391.1| PREDICTED: hyphally regulated cell wall protein 3 [Drosophil
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>gi|1036743445|ref|XP_017155944.1| PREDICTED: brefeldin A-inhibited guanine nucleotide-exchang
>gi|1036752773|ref|XP_017156456.1| PREDICTED: malignant T-cell-amplified sequence 1 homolog [Dr
>gi|1036752230|ref|XP_017156423.1| PREDICTED: cell cycle negative regulator roughex [Drosophila
>gi|1036751899|ref|XP_017156403.1| PREDICTED: protein mab-21 [Drosophila miranda]
>gi|1036754710|ref|XP_017156574.1| PREDICTED: protein mab-21-like [Drosophila miranda]
>gi|1036754725|ref|XP_017156575.1| PREDICTED: protein mab-21-like [Drosophila miranda]
>gi|1036750957|ref|XP_017156349.1| PREDICTED: pancreatic triacylglycerol lipase [Drosophila mi
>gi|1036741159|ref|XP_017155811.1| PREDICTED: glutamate receptor-interacting protein 1 isoform
>gi|1036741141|ref|XP_017155810.1| PREDICTED: glutamate receptor-interacting protein 1 isoform
>gi|1036740711|ref|XP_017155683.1| PREDICTED: uncharacterized protein LOC108164460 [Drosophila
>gi|1036740729|ref|XP_017155697.1| PREDICTED: uncharacterized protein LOC108164466 [Drosophila
>gi|1036706194|ref|XP_017136132.1| PREDICTED: trypsin eta-like [Drosophila miranda]
>gi|1036704001|ref|XP_017134631.1| PREDICTED: trypsin-1-like [Drosophila miranda]
>gi|1036745287|ref|XP_017156037.1| PREDICTED: trypsin I-P1 isoform X1 [Drosophila miranda]
>gi|1036745304|ref|XP_017156039.1| PREDICTED: trypsin I-P1 isoform X2 [Drosophila miranda]
>gi|1036745321|ref|XP_017156040.1| PREDICTED: trypsin I-P1 isoform X3 [Drosophila miranda]
>gi|1036745338|ref|XP_017156041.1| PREDICTED: uncharacterized protein LOC108164685 isoform X4
>gi|1036745354|ref|XP_017156042.1| PREDICTED: urokinase-type plasminogen activator isoform X5
>gi|1036745370|ref|XP_017156043.1| PREDICTED: trypsin eta isoform X1 [Drosophila miranda]
>gi|1036745387|ref|XP_017156044.1| PREDICTED: trypsin eta isoform X2 [Drosophila miranda]
>gi|1036745404|ref|XP_017156045.1| PREDICTED: trypsin eta isoform X3 [Drosophila miranda]
>gi|1036745421|ref|XP_017156046.1| PREDICTED: uncharacterized protein LOC108164687 [Drosophila
>gi|1036745233|ref|XP_017156034.1| PREDICTED: protein bric-a-brac 1 isoform X1 [Drosophila mira
>gi|1036745269|ref|XP_017156036.1| PREDICTED: protein bric-a-brac 1 isoform X3 [Drosophila mira
>gi|1036745252|ref|XP_017156035.1| PREDICTED: protein bric-a-brac 1 isoform X2 [Drosophila mira
>gi|1036712511|ref|XP_017140078.1| PREDICTED: serine hydroxymethyltransferase, cytosolic [Dros
>gi|1036712594|ref|XP_017140128.1| PREDICTED: transmembrane protein 256 homolog [Drosophila mi
>gi|1036712528|ref|XP_017140088.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036712545|ref|XP_017140096.1| PREDICTED: modifier of mdg4 isoform X2 [Drosophila miranda]
>gi|1036712561|ref|XP_017140108.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosoph
>gi|1036712577|ref|XP_017140118.1| PREDICTED: actin-5C [Drosophila miranda]
>gi|1036742251|ref|XP_017155874.1| PREDICTED: serine/threonine-protein phosphatase beta isoform
>gi|1036742233|ref|XP_017155873.1| PREDICTED: serine/threonine-protein phosphatase beta isoform
>gi|1036755592|ref|XP_017156621.1| PREDICTED: uncharacterized protein LOC108165082 isoform X1
>gi|1036755609|ref|XP_017156622.1| PREDICTED: uncharacterized protein LOC108165082 isoform X2
>gi|1036703787|ref|XP_017157160.1| PREDICTED: sphingomyelin synthase-related 1-like [Drosophila
>gi|1036699261|ref|XP_017139456.1| PREDICTED: odorant receptor 9a [Drosophila miranda]
>gi|1036743789|ref|XP_017155959.1| PREDICTED: uncharacterized protein LOC108164624 [Drosophila
>gi|1036743624|ref|XP_017155952.1| PREDICTED: uncharacterized protein LOC108164621 isoform X4
>gi|1036743549|ref|XP_017155948.1| PREDICTED: uncharacterized protein LOC108164621 isoform X1
>gi|1036743568|ref|XP_017155949.1| PREDICTED: uncharacterized protein LOC108164621 isoform X1
>gi|1036743587|ref|XP_017155950.1| PREDICTED: uncharacterized protein LOC108164621 isoform X2
>gi|1036743606|ref|XP_017155951.1| PREDICTED: uncharacterized protein LOC108164621 isoform X3
>gi|1036743716|ref|XP_017155957.1| PREDICTED: uncharacterized protein LOC108164621 isoform X8
>gi|1036743643|ref|XP_017155953.1| PREDICTED: uncharacterized protein LOC108164621 isoform X5
>gi|1036743661|ref|XP_017155954.1| PREDICTED: uncharacterized protein LOC108164621 isoform X6
>gi|1036743680|ref|XP_017155955.1| PREDICTED: uncharacterized protein LOC108164621 isoform X6
>gi|1036743699|ref|XP_017155956.1| PREDICTED: uncharacterized protein LOC108164621 isoform X7
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>gi|1036748257|ref|XP_017156213.1| PREDICTED: up-regulated during skeletal muscle growth protei
>gi|1036719849|ref|XP_017144338.1| PREDICTED: neurogenic protein mastermind [Drosophila miranda
>gi|1036719827|ref|XP_017144316.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036733434|ref|XP_017151633.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1036733452|ref|XP_017151641.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1036733470|ref|XP_017151650.1| PREDICTED: serine/threonine-protein phosphatase 4 regulatory
>gi|1036733489|ref|XP_017151662.1| PREDICTED: leucine-rich repeat-containing protein 58 [Drosop
>gi|1036722475|ref|XP_017145554.1| PREDICTED: protein sprint isoform X3 [Drosophila miranda]
>gi|1036722494|ref|XP_017145555.1| PREDICTED: protein sprint isoform X3 [Drosophila miranda]
>gi|1036722512|ref|XP_017145556.1| PREDICTED: protein sprint isoform X4 [Drosophila miranda]
>gi|1036722437|ref|XP_017145540.1| PREDICTED: protein sprint isoform X1 [Drosophila miranda]
>gi|1036722456|ref|XP_017145548.1| PREDICTED: protein sprint isoform X2 [Drosophila miranda]
>gi|1036722529|ref|XP_017145557.1| PREDICTED: protein sprint isoform X5 [Drosophila miranda]
>gi|1036722550|ref|XP_017145558.1| PREDICTED: protein sprint isoform X5 [Drosophila miranda]
>gi|1036722569|ref|XP_017145559.1| PREDICTED: protein sprint isoform X6 [Drosophila miranda]
>gi|1036722646|ref|XP_017145587.1| PREDICTED: uncharacterized protein LOC108157950 [Drosophila
>gi|1036722667|ref|XP_017145595.1| PREDICTED: uncharacterized protein LOC108157950 [Drosophila
>gi|1036722592|ref|XP_017145560.1| PREDICTED: microspherule protein 1-like [Drosophila miranda]
>gi|1036722609|ref|XP_017145567.1| PREDICTED: microspherule protein 1-like [Drosophila miranda]
>gi|1036722682|ref|XP_017145606.1| PREDICTED: uncharacterized protein LOC108157968 [Drosophila
>gi|1036722628|ref|XP_017145576.1| PREDICTED: uncharacterized protein F09G8.5-like [Drosophila
>gi|1036706161|ref|XP_017136113.1| PREDICTED: GTP-binding protein ypt1-like [Drosophila miranda
>gi|1036731735|ref|XP_017150828.1| PREDICTED: uncharacterized protein LOC108161083 [Drosophila
>gi|1036738937|ref|XP_017154693.1| PREDICTED: FAST kinase domain-containing protein 5, mitochon
>gi|1036738953|ref|XP_017154706.1| PREDICTED: LOW QUALITY PROTEIN: mitochondrial ornithine tra
>gi|1036751088|ref|XP_017156357.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036741801|ref|XP_017155852.1| PREDICTED: uncharacterized protein LOC108164549 [Drosophila
>gi|1036741693|ref|XP_017155845.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741710|ref|XP_017155846.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741731|ref|XP_017155847.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741747|ref|XP_017155848.1| PREDICTED: uncharacterized protein LOC108164547 isoform X1
>gi|1036741768|ref|XP_017155849.1| PREDICTED: uncharacterized protein LOC108164547 isoform X2
>gi|1036741784|ref|XP_017155850.1| PREDICTED: uncharacterized protein LOC108164548 [Drosophila
>gi|1036705425|ref|XP_017135596.1| PREDICTED: rRNA biogenesis protein rrp36-like [Drosophila m
>gi|1036725961|ref|XP_017147489.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase a
>gi|1036725984|ref|XP_017147497.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase a
>gi|1036725923|ref|XP_017147468.1| PREDICTED: BTB/POZ domain-containing protein 9 isoform X1 [
>gi|1036725942|ref|XP_017147476.1| PREDICTED: BTB/POZ domain-containing protein 9 isoform X2 [
>gi|1036717963|ref|XP_017143290.1| PREDICTED: uncharacterized protein LOC108156382 [Drosophila
>gi|1036717885|ref|XP_017143246.1| PREDICTED: ADP,ATP carrier protein [Drosophila miranda]
>gi|1036717904|ref|XP_017143257.1| PREDICTED: ADP,ATP carrier protein [Drosophila miranda]
>gi|1036717926|ref|XP_017143266.1| PREDICTED: ADP,ATP carrier protein [Drosophila miranda]
>gi|1036717866|ref|XP_017143234.1| PREDICTED: protein penguin [Drosophila miranda]
>gi|1036717944|ref|XP_017143278.1| PREDICTED: putative peptidyl-prolyl cis-trans isomerase dod
>gi|1036728836|ref|XP_017149195.1| PREDICTED: sphingosine kinase 2 [Drosophila miranda]
>gi|1036728748|ref|XP_017149147.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728767|ref|XP_017149154.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728784|ref|XP_017149163.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
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>gi|1036728801|ref|XP_017149173.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036728819|ref|XP_017149182.1| PREDICTED: ecotropic viral integration site 5 ortholog isof
>gi|1036746435|ref|XP_017156107.1| PREDICTED: glutamine synthetase 2 cytoplasmic isoform X3 [Di
>gi|1036746403|ref|XP_017156105.1| PREDICTED: glutamine synthetase 2 cytoplasmic isoform X1 [Di
>gi|1036746420|ref|XP_017156106.1| PREDICTED: glutamine synthetase 2 cytoplasmic isoform X2 [Di
>gi|1036746453|ref|XP_017156108.1| PREDICTED: odorant receptor 10a [Drosophila miranda]
>gi|1036746385|ref|XP_017156103.1| PREDICTED: gustatory receptor 10a [Drosophila miranda]
>gi|1036707473|ref|XP_017136981.1| PREDICTED: putative gustatory receptor 10b [Drosophila miran
>gi|1036715328|ref|XP_017141794.1| PREDICTED: uncharacterized protein LOC108155479 [Drosophila
>gi|1036715568|ref|XP_017141910.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715549|ref|XP_017141902.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715511|ref|XP_017141885.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715475|ref|XP_017141866.1| PREDICTED: disks large 1 tumor suppressor protein isoform X8
>gi|1036715606|ref|XP_017141927.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715458|ref|XP_017141857.1| PREDICTED: disks large 1 tumor suppressor protein isoform X7
>gi|1036715437|ref|XP_017141848.1| PREDICTED: disks large 1 tumor suppressor protein isoform X6
>gi|1036715419|ref|XP_017141841.1| PREDICTED: disks large 1 tumor suppressor protein isoform X5
>gi|1036715402|ref|XP_017141833.1| PREDICTED: disks large 1 tumor suppressor protein isoform X4
>gi|1036715383|ref|XP_017141823.1| PREDICTED: disks large 1 tumor suppressor protein isoform X3
>gi|1036715366|ref|XP_017141814.1| PREDICTED: disks large 1 tumor suppressor protein isoform X2
>gi|1036715347|ref|XP_017141805.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715587|ref|XP_017141918.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715530|ref|XP_017141893.1| PREDICTED: disks large 1 tumor suppressor protein isoform X1
>gi|1036715494|ref|XP_017141876.1| PREDICTED: disks large 1 tumor suppressor protein isoform X9
>gi|1036715624|ref|XP_017141939.1| PREDICTED: uncharacterized protein LOC108155555 [Drosophila
>gi|1036752008|ref|XP_017156410.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036732597|ref|XP_017151220.1| PREDICTED: tyrosine-protein kinase hopscotch [Drosophila mi
>gi|1036751627|ref|XP_017156388.1| PREDICTED: uncharacterized protein LOC108164913 [Drosophila
>gi|1036743173|ref|XP_017155928.1| PREDICTED: segment polarity protein dishevelled [Drosophila
>gi|1036724085|ref|XP_017146341.1| PREDICTED: uncharacterized protein LOC108158504 [Drosophila
>gi|1036724143|ref|XP_017146384.1| PREDICTED: uncharacterized protein LOC108158514 [Drosophila
>gi|1036724124|ref|XP_017146371.1| PREDICTED: uncharacterized protein C1orf131 [Drosophila mira
>gi|1036724106|ref|XP_017146356.1| PREDICTED: alanine--glyoxylate aminotransferase 2-like [Dros
>gi|1036699276|ref|XP_017139662.1| PREDICTED: E3 ubiquitin-protein ligase RFWD3-like, partial
>gi|1036733820|ref|XP_017151829.1| PREDICTED: putative vitellogenin receptor isoform X2 [Drosop
>gi|1036733801|ref|XP_017151821.1| PREDICTED: putative vitellogenin receptor isoform X1 [Drosop
>gi|1036733890|ref|XP_017151883.1| PREDICTED: uncharacterized protein LOC108161948 [Drosophila
>gi|1036733947|ref|XP_017151921.1| PREDICTED: prisilkin-39 [Drosophila miranda]
>gi|1036733873|ref|XP_017151869.1| PREDICTED: probable ATP-dependent RNA helicase ddx17 [Drosop
>gi|1036733928|ref|XP_017151910.1| PREDICTED: uncharacterized protein LOC108161959 [Drosophila
>gi|1036733855|ref|XP_017151856.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
>gi|1036699291|ref|XP_017139786.1| PREDICTED: uncharacterized protein LOC108154124 [Drosophila
>gi|1036699305|ref|XP_017139917.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036699320|ref|XP_017140034.1| PREDICTED: proline-rich protein HaeIII subfamily 1 [Drosophi
>gi|1036733909|ref|XP_017151898.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
>gi|1036733837|ref|XP_017151842.1| PREDICTED: hyphal wall protein 1 [Drosophila miranda]
>gi|1036707092|ref|XP_017136749.1| PREDICTED: MICOS complex subunit Mic10 [Drosophila miranda]
>gi|1036704766|ref|XP_017135126.1| PREDICTED: probable mitochondrial import inner membrane tran
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>gi|1036705546|ref|XP_017135668.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036705072|ref|XP_017135358.1| PREDICTED: uncharacterized protein LOC108151319 [Drosophila
>gi|1036706211|ref|XP_017136146.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036705102|ref|XP_017135382.1| PREDICTED: sperm flagellar protein 1-like [Drosophila miran
>gi|1036704941|ref|XP_017135256.1| PREDICTED: uncharacterized protein LOC108151248 [Drosophila
>gi|1036704956|ref|XP_017135264.1| PREDICTED: uncharacterized protein LOC108151248 [Drosophila
>gi|1036699335|ref|XP_017140145.1| PREDICTED: uncharacterized protein LOC108154357 [Drosophila
>gi|1036744345|ref|XP_017155986.1| PREDICTED: ubiquitin-conjugating enzyme E2 N [Drosophila mi
>gi|1036744363|ref|XP_017155987.1| PREDICTED: ubiquitin-conjugating enzyme E2 N [Drosophila mi
>gi|1036700935|ref|XP_017152298.1| PREDICTED: uncharacterized protein LOC108162178 [Drosophila
>gi|1036701120|ref|XP_017153709.1| PREDICTED: uncharacterized protein LOC108163093 [Drosophila
>gi|1036700721|ref|XP_017150989.1| PREDICTED: uncharacterized protein LOC108161245 [Drosophila
>gi|1036700662|ref|XP_017150645.1| PREDICTED: AMP deaminase 2 isoform X2 [Drosophila miranda]
>gi|1036700645|ref|XP_017150564.1| PREDICTED: AMP deaminase 2 isoform X1 [Drosophila miranda]
>gi|1036700677|ref|XP_017150723.1| PREDICTED: AMP deaminase 2 isoform X3 [Drosophila miranda]
>gi|1036700707|ref|XP_017150886.1| PREDICTED: LIM domain-containing protein jub [Drosophila mi
>gi|1036700614|ref|XP_017150318.1| PREDICTED: nuclear pore complex protein Nup93 [Drosophila m
>gi|1036700983|ref|XP_017152620.1| PREDICTED: chloride intracellular channel exc-4 isoform X2
>gi|1036700969|ref|XP_017152540.1| PREDICTED: chloride intracellular channel exc-4 isoform X1
>gi|1036700954|ref|XP_017152418.1| PREDICTED: nucleoside diphosphate-linked moiety X motif 8 [
>gi|1036700600|ref|XP_017150194.1| PREDICTED: AP-1 complex subunit beta-1 [Drosophila miranda]
>gi|1036700868|ref|XP_017151767.1| PREDICTED: transmembrane protein 198 [Drosophila miranda]
>gi|1036701105|ref|XP_017153584.1| PREDICTED: cytochrome b-c1 complex subunit 10 [Drosophila m
>gi|1036701090|ref|XP_017153460.1| PREDICTED: cytochrome c oxidase subunit 6B1 [Drosophila mir
>gi|1036700882|ref|XP_017151897.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1036700895|ref|XP_017151980.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1036700907|ref|XP_017152065.1| PREDICTED: TRAF3-interacting protein 1 [Drosophila miranda]
>gi|1036701045|ref|XP_017153097.1| PREDICTED: ribonuclease P protein subunit p20 [Drosophila m
>gi|1036701075|ref|XP_017153340.1| PREDICTED: transmembrane protein 203 [Drosophila miranda]
>gi|1036700922|ref|XP_017152189.1| PREDICTED: beta-parvin [Drosophila miranda]
>gi|1036700998|ref|XP_017152732.1| PREDICTED: FAD-linked sulfhydryl oxidase ALR [Drosophila mi
>gi|1036701060|ref|XP_017153217.1| PREDICTED: uncharacterized protein LOC108162787 [Drosophila
>gi|1036700693|ref|XP_017150824.1| PREDICTED: uncharacterized protein LOC108161006 [Drosophila
>gi|1036700395|ref|XP_017148872.1| PREDICTED: nuclear pore complex protein Nup153 isoform X1 [
>gi|1036700409|ref|XP_017148950.1| PREDICTED: nuclear pore complex protein Nup153 isoform X1 [
>gi|1036700424|ref|XP_017149035.1| PREDICTED: nuclear pore complex protein Nup153 isoform X2 [
>gi|1036700453|ref|XP_017149204.1| PREDICTED: nuclear pore complex protein Nup153 isoform X4 [
>gi|1036700438|ref|XP_017149115.1| PREDICTED: nuclear pore complex protein Nup153 isoform X3 [
>gi|1036700736|ref|XP_017151097.1| PREDICTED: serine/threonine-protein kinase PAK mbt [Drosophi
>gi|1036700336|ref|XP_017148507.1| PREDICTED: homeotic protein female sterile isoform X5 [Droso
>gi|1036700307|ref|XP_017148328.1| PREDICTED: homeotic protein female sterile isoform X3 [Droso
>gi|1036700292|ref|XP_017148241.1| PREDICTED: homeotic protein female sterile isoform X2 [Droso
>gi|1036700321|ref|XP_017148412.1| PREDICTED: homeotic protein female sterile isoform X4 [Droso
>gi|1036700202|ref|XP_017147727.1| PREDICTED: homeotic protein female sterile isoform X1 [Droso
>gi|1036700216|ref|XP_017147815.1| PREDICTED: homeotic protein female sterile isoform X1 [Droso
>gi|1036700229|ref|XP_017147901.1| PREDICTED: homeotic protein female sterile isoform X1 [Droso
>gi|1036700242|ref|XP_017147986.1| PREDICTED: homeotic protein female sterile isoform X1 [Droso
>gi|1036700256|ref|XP_017148072.1| PREDICTED: homeotic protein female sterile isoform X1 [Droso
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>gi|1036700273|ref|XP_017148151.1| PREDICTED: homeotic protein female sterile isoform X1 [Dros
>gi|1036700353|ref|XP_017148586.1| PREDICTED: homeotic protein female sterile isoform X6 [Dros
>gi|1036700381|ref|XP_017148753.1| PREDICTED: homeotic protein female sterile isoform X8 [Dros
>gi|1036700367|ref|XP_017148671.1| PREDICTED: homeotic protein female sterile isoform X7 [Dros
>gi|1036700826|ref|XP_017151493.1| PREDICTED: multifunctional protein ADE2 [Drosophila miranda]
>gi|1036700630|ref|XP_017150441.1| PREDICTED: TBC domain-containing protein kinase-like protei
>gi|1036700811|ref|XP_017151391.1| PREDICTED: heparan-alpha-glucosaminide N-acetyltransferase
>gi|1036700750|ref|XP_017151204.1| PREDICTED: uncharacterized protein LOC108161422 [Drosophila
>gi|1036700787|ref|XP_017151279.1| PREDICTED: uncharacterized protein LOC108161422 [Drosophila
>gi|1036700844|ref|XP_017151607.1| PREDICTED: zinc finger BED domain-containing protein 4-like
>gi|1036700526|ref|XP_017149660.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X3 [Dros
>gi|1036700468|ref|XP_017149328.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036700483|ref|XP_017149411.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036700497|ref|XP_017149493.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036700511|ref|XP_017149577.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X2 [Dros
>gi|1036701028|ref|XP_017152973.1| PREDICTED: uncharacterized protein LOC108162621 [Drosophila
>gi|1036701013|ref|XP_017152853.1| PREDICTED: CTD nuclear envelope phosphatase 1A-like [Drosoph
>gi|1036700541|ref|XP_017149790.1| PREDICTED: protein ovo isoform X1 [Drosophila miranda]
>gi|1036700555|ref|XP_017149875.1| PREDICTED: protein ovo isoform X2 [Drosophila miranda]
>gi|1036700570|ref|XP_017149993.1| PREDICTED: zinc finger protein 236 [Drosophila miranda]
>gi|1036700585|ref|XP_017150075.1| PREDICTED: zinc finger protein 236 [Drosophila miranda]
>gi|1036739071|ref|XP_017154762.1| PREDICTED: innexin inx1 [Drosophila miranda]
>gi|1036739089|ref|XP_017154770.1| PREDICTED: innexin inx1 [Drosophila miranda]
>gi|1036753592|ref|XP_017156507.1| PREDICTED: uncharacterized protein LOC108164993 [Drosophila
>gi|1036755344|ref|XP_017156611.1| PREDICTED: innexin inx7 [Drosophila miranda]
>gi|1036732044|ref|XP_017150958.1| PREDICTED: innexin inx2 [Drosophila miranda]
>gi|1036736115|ref|XP_017153134.1| PREDICTED: mitochondrial import receptor subunit TOM7 homolo
>gi|1036736151|ref|XP_017153160.1| PREDICTED: AP-3 complex subunit mu-2 [Drosophila miranda]
>gi|1036736174|ref|XP_017153171.1| PREDICTED: small integral membrane protein 4 [Drosophila mi
>gi|1036736190|ref|XP_017153183.1| PREDICTED: uncharacterized protein C12orf73 homolog [Drosoph
>gi|1036736133|ref|XP_017153146.1| PREDICTED: zinc finger protein 782 [Drosophila miranda]
>gi|1036748513|ref|XP_017156216.1| PREDICTED: uncharacterized protein LOC108164807 [Drosophila
>gi|1036748551|ref|XP_017156218.1| PREDICTED: protein nullo [Drosophila miranda]
>gi|1036748532|ref|XP_017156217.1| PREDICTED: uncharacterized protein LOC108164808 [Drosophila
>gi|1036748570|ref|XP_017156219.1| PREDICTED: uncharacterized protein LOC108164810 [Drosophila
>gi|1036731237|ref|XP_017150618.1| PREDICTED: uncharacterized protein LOC108160864 isoform X1
>gi|1036731292|ref|XP_017150646.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036731307|ref|XP_017150653.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036731273|ref|XP_017150635.1| PREDICTED: uncharacterized protein LOC108160864 isoform X3
>gi|1036731254|ref|XP_017150626.1| PREDICTED: uncharacterized protein LOC108160864 isoform X2
>gi|1036722404|ref|XP_017145521.1| PREDICTED: peptide transporter family 1 [Drosophila miranda]
>gi|1036699352|ref|XP_017140257.1| PREDICTED: uncharacterized protein LOC108154456 [Drosophila
>gi|1036722387|ref|XP_017145509.1| PREDICTED: peptide transporter family 1 isoform X4 [Drosophi
>gi|1036722349|ref|XP_017145494.1| PREDICTED: peptide transporter family 1 isoform X2 [Drosophi
>gi|1036722368|ref|XP_017145502.1| PREDICTED: peptide transporter family 1 isoform X3 [Drosophi
>gi|1036722328|ref|XP_017145485.1| PREDICTED: peptide transporter family 1 isoform X1 [Drosophi
>gi|1036741634|ref|XP_017155841.1| PREDICTED: zinc finger protein ZPR1 [Drosophila miranda]
>gi|1036741672|ref|XP_017155844.1| PREDICTED: uncharacterized protein LOC108164546 [Drosophila
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>gi|1036741653|ref|XP_017155843.1| PREDICTED: uncharacterized protein LOC108164545 [Drosophila
>gi|1036722164|ref|XP_017145379.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1036722185|ref|XP_017145391.1| PREDICTED: histone H3.3 [Drosophila miranda]
>gi|1036722145|ref|XP_017145369.1| PREDICTED: basic proline-rich protein [Drosophila miranda]
>gi|1036729638|ref|XP_017149692.1| PREDICTED: frataxin homolog, mitochondrial [Drosophila miran
>gi|1036729619|ref|XP_017149678.1| PREDICTED: uncharacterized protein CG7065 [Drosophila mirand
>gi|1036754237|ref|XP_017156546.1| PREDICTED: probable sodium/metabolite cotransporter BASS5, c
>gi|1036739338|ref|XP_017154934.1| PREDICTED: transmembrane protein fend [Drosophila miranda]
>gi|1036722204|ref|XP_017145403.1| PREDICTED: LOW QUALITY PROTEIN: AP-1 complex subunit gamma-1
>gi|1036722241|ref|XP_017145426.1| PREDICTED: ADP-ribosylation factor-related protein 1 [Drosop
>gi|1036722260|ref|XP_017145438.1| PREDICTED: protein CutA homolog [Drosophila miranda]
>gi|1036722222|ref|XP_017145415.1| PREDICTED: puff-specific protein Bx42 [Drosophila miranda]
>gi|1036731362|ref|XP_017150681.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731447|ref|XP_017150719.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731483|ref|XP_017150735.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731326|ref|XP_017150665.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731344|ref|XP_017150673.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731394|ref|XP_017150696.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731431|ref|XP_017150711.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731377|ref|XP_017150687.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731412|ref|XP_017150704.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731466|ref|XP_017150727.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731518|ref|XP_017150751.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036731500|ref|XP_017150743.1| PREDICTED: probable phosphorylase b kinase regulatory subuni
>gi|1036743858|ref|XP_017155961.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036743839|ref|XP_017155960.1| PREDICTED: T-complex protein 1 subunit beta [Drosophila mira
>gi|1036757646|ref|XP_017156735.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036753388|ref|XP_017156495.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036753426|ref|XP_017156498.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036753443|ref|XP_017156499.1| PREDICTED: sporulation-specific protein 15-like isoform X4
>gi|1036753459|ref|XP_017156500.1| PREDICTED: uncharacterized protein LOC108164985 isoform X5
>gi|1036753407|ref|XP_017156496.1| PREDICTED: golgin subfamily A member 6-like protein 6 isofor
>gi|1036753479|ref|XP_017156501.1| PREDICTED: golgin subfamily A member 6-like protein 10 [Dros
>gi|1036704479|ref|XP_017134919.1| PREDICTED: uncharacterized protein LOC108151052 [Drosophila
>gi|1036705342|ref|XP_017135541.1| PREDICTED: uncharacterized protein LOC108151439 isoform X1
>gi|1036705360|ref|XP_017135550.1| PREDICTED: uncharacterized protein LOC108151439 isoform X2
>gi|1036705060|ref|XP_017135345.1| PREDICTED: uncharacterized protein LOC108151308 isoform X2
>gi|1036705046|ref|XP_017135338.1| PREDICTED: uncharacterized protein LOC108151308 isoform X1
>gi|1036707523|ref|XP_017137022.1| PREDICTED: chondroadherin [Drosophila miranda]
>gi|1036756414|ref|XP_017156669.1| PREDICTED: innexin inx6 [Drosophila miranda]
>gi|1036756432|ref|XP_017156670.1| PREDICTED: innexin inx6 [Drosophila miranda]
>gi|1036756450|ref|XP_017156671.1| PREDICTED: uncharacterized protein LOC108165120 [Drosophila
>gi|1036740839|ref|XP_017155773.1| PREDICTED: N-acetylglucosamine-6-phosphate deacetylase [Dros
>gi|1036740856|ref|XP_017155781.1| PREDICTED: ras-related protein Rab-10 [Drosophila miranda]
>gi|1036741107|ref|XP_017155808.1| PREDICTED: uncharacterized protein LOC108164527 [Drosophila
>gi|1036741123|ref|XP_017155809.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036755363|ref|XP_017156612.1| PREDICTED: peritrophin-1 [Drosophila miranda]
>gi|1036713844|ref|XP_017140937.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform
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>gi|1036713861|ref|XP_017140946.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713876|ref|XP_017140954.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713891|ref|XP_017140964.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713913|ref|XP_017140971.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713964|ref|XP_017140998.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713979|ref|XP_017141008.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713930|ref|XP_017140979.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713947|ref|XP_017140987.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714030|ref|XP_017141030.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714071|ref|XP_017141046.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714088|ref|XP_017141052.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036713996|ref|XP_017141014.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714013|ref|XP_017141022.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714044|ref|XP_017141037.1| PREDICTED: cytoplasmic dynein 1 intermediate chain isoform X
>gi|1036714117|ref|XP_017141076.1| PREDICTED: annexin B10 isoform X1 [Drosophila miranda]
>gi|1036714131|ref|XP_017141086.1| PREDICTED: annexin B10 isoform X2 [Drosophila miranda]
>gi|1036714180|ref|XP_017141120.1| PREDICTED: TIP41-like protein isoform X2 [Drosophila miranda
>gi|1036714162|ref|XP_017141111.1| PREDICTED: TIP41-like protein isoform X1 [Drosophila miranda
>gi|1036714103|ref|XP_017141064.1| PREDICTED: probable Xaa-Pro aminopeptidase 3 [Drosophila mir
>gi|1036714146|ref|XP_017141099.1| PREDICTED: delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mito
>gi|1036714216|ref|XP_017141140.1| PREDICTED: PHD finger protein 7 [Drosophila miranda]
>gi|1036742216|ref|XP_017155872.1| PREDICTED: ras-related protein Rab-35 [Drosophila miranda]
>gi|1036754076|ref|XP_017156536.1| PREDICTED: neurogenic locus notch homolog protein 3 isoform
>gi|1036754057|ref|XP_017156535.1| PREDICTED: neurogenic locus notch homolog protein 3 isoform
>gi|1036754039|ref|XP_017156534.1| PREDICTED: neurogenic locus notch homolog protein 3 isoform
>gi|1036699367|ref|XP_017140365.1| PREDICTED: putative odorant receptor 19b [Drosophila miranda
>gi|1036699381|ref|XP_017140474.1| PREDICTED: serine/threonine-protein kinase Warts, partial [
>gi|1036704989|ref|XP_017135290.1| PREDICTED: uncharacterized protein LOC108151272 [Drosophila
>gi|1036699395|ref|XP_017140596.1| PREDICTED: uncharacterized protein LOC108154714 [Drosophila
>gi|1036707438|ref|XP_017136960.1| PREDICTED: uncharacterized protein LOC108152259 [Drosophila
>gi|1036699410|ref|XP_017140835.1| PREDICTED: uncharacterized protein LOC108154869 [Drosophila
>gi|1036752500|ref|XP_017156440.1| PREDICTED: dopamine D2-like receptor isoform X1 [Drosophila
>gi|1036752593|ref|XP_017156445.1| PREDICTED: dopamine D2-like receptor isoform X6 [Drosophila
>gi|1036752538|ref|XP_017156442.1| PREDICTED: dopamine D2-like receptor isoform X3 [Drosophila
>gi|1036752558|ref|XP_017156443.1| PREDICTED: dopamine D2-like receptor isoform X4 [Drosophila
>gi|1036752519|ref|XP_017156441.1| PREDICTED: dopamine D2-like receptor isoform X2 [Drosophila
>gi|1036752577|ref|XP_017156444.1| PREDICTED: dopamine D2-like receptor isoform X5 [Drosophila
>gi|1036749517|ref|XP_017156270.1| PREDICTED: endothelin-converting enzyme 1 isoform X2 [Drosop
>gi|1036749500|ref|XP_017156269.1| PREDICTED: endothelin-converting enzyme 1 isoform X1 [Drosop
>gi|1036703827|ref|XP_017134536.1| PREDICTED: glycine-rich cell wall structural protein [Drosop
>gi|1036723465|ref|XP_017145989.1| PREDICTED: uncharacterized protein LOC108158260 [Drosophila
>gi|1036744574|ref|XP_017155997.1| PREDICTED: uncharacterized protein LOC108164656 [Drosophila
>gi|1036711778|ref|XP_017139678.1| PREDICTED: nuclear pore complex protein Nup205 isoform X1 [
>gi|1036711795|ref|XP_017139686.1| PREDICTED: nuclear pore complex protein Nup205 isoform X2 [
>gi|1036711812|ref|XP_017139694.1| PREDICTED: ras-associated and pleckstrin homology domains-c
>gi|1036749550|ref|XP_017156272.1| PREDICTED: uncharacterized protein LOC108164854 isoform X1
>gi|1036749569|ref|XP_017156273.1| PREDICTED: uncharacterized protein LOC108164854 isoform X2
>gi|1036731861|ref|XP_017150873.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
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>gi|1036726653|ref|XP_017147924.1| PREDICTED: uncharacterized protein LOC108159272 [Drosophila
>gi|1036726671|ref|XP_017147935.1| PREDICTED: ATP-binding cassette sub-family D member 3 [Droso
>gi|1036726689|ref|XP_017147944.1| PREDICTED: ATP-binding cassette sub-family D member 3 [Droso
>gi|1036742734|ref|XP_017155902.1| PREDICTED: bifunctional lysine-specific demethylase and hist
>gi|1036742752|ref|XP_017155903.1| PREDICTED: 39S ribosomal protein L33, mitochondrial [Drosoph
>gi|1036721355|ref|XP_017145072.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721297|ref|XP_017145045.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721333|ref|XP_017145062.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721278|ref|XP_017145036.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036721314|ref|XP_017145053.1| PREDICTED: insulin-like growth factor 2 mRNA-binding protei
>gi|1036716143|ref|XP_017142245.1| PREDICTED: protein transport protein sec31 [Drosophila mira
>gi|1036716158|ref|XP_017142259.1| PREDICTED: 60S ribosomal protein L22 [Drosophila miranda]
>gi|1036716232|ref|XP_017142306.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036716251|ref|XP_017142315.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036716212|ref|XP_017142293.1| PREDICTED: histone-lysine N-methyltransferase SMYD3 [Drosoph
>gi|1036716194|ref|XP_017142280.1| PREDICTED: centromere/kinetochore protein zw10 [Drosophila r
>gi|1036735957|ref|XP_017153045.1| PREDICTED: chromosome-associated kinesin KIF4 [Drosophila mi
>gi|1036734580|ref|XP_017152284.1| PREDICTED: beta-1,4-mannosyltransferase egh [Drosophila mira
>gi|1036734598|ref|XP_017152293.1| PREDICTED: beta-1,4-mannosyltransferase egh [Drosophila mira
>gi|1036729655|ref|XP_017149704.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036729674|ref|XP_017149717.1| PREDICTED: dynein light chain 1, cytoplasmic [Drosophila mir
>gi|1036729693|ref|XP_017149725.1| PREDICTED: dynein light chain 1, cytoplasmic [Drosophila mir
>gi|1036729713|ref|XP_017149733.1| PREDICTED: dynein light chain 1, cytoplasmic [Drosophila mir
>gi|1036729187|ref|XP_017149419.1| PREDICTED: uncharacterized protein LOC108160132 [Drosophila
>gi|1036734331|ref|XP_017152134.1| PREDICTED: uncharacterized protein LOC108162040 isoform X10
>gi|1036734278|ref|XP_017152117.1| PREDICTED: uncharacterized protein LOC108162040 isoform X8
>gi|1036734255|ref|XP_017152109.1| PREDICTED: uncharacterized protein LOC108162040 isoform X7
>gi|1036734222|ref|XP_017152092.1| PREDICTED: uncharacterized protein LOC108162040 isoform X5
>gi|1036734168|ref|XP_017152066.1| PREDICTED: uncharacterized protein LOC108162040 isoform X2
>gi|1036734241|ref|XP_017152100.1| PREDICTED: uncharacterized protein LOC108162040 isoform X6
>gi|1036734185|ref|XP_017152075.1| PREDICTED: uncharacterized protein LOC108162040 isoform X3
>gi|1036734296|ref|XP_017152125.1| PREDICTED: uncharacterized protein LOC108162040 isoform X9
>gi|1036734204|ref|XP_017152084.1| PREDICTED: uncharacterized protein LOC108162040 isoform X4
>gi|1036734150|ref|XP_017152057.1| PREDICTED: uncharacterized protein LOC108162040 isoform X1
>gi|1036743208|ref|XP_017155931.1| PREDICTED: uncharacterized protein DDB_G0271670 [Drosophila
>gi|1036743192|ref|XP_017155929.1| PREDICTED: uncharacterized protein LOC108164602 [Drosophila
>gi|1036740539|ref|XP_017155570.1| PREDICTED: anoctamin-10 [Drosophila miranda]
>gi|1036740556|ref|XP_017155584.1| PREDICTED: uncharacterized protein LOC108164415 [Drosophila
>gi|1036740575|ref|XP_017155594.1| PREDICTED: endochitinase A-like [Drosophila miranda]
>gi|1036740592|ref|XP_017155607.1| PREDICTED: uncharacterized protein LOC108164430 [Drosophila
>gi|1036740608|ref|XP_017155617.1| PREDICTED: uncharacterized protein LOC108164430 [Drosophila
>gi|1036718667|ref|XP_017143662.1| PREDICTED: DENN domain-containing protein 4C isoform X1 [Dr
>gi|1036718684|ref|XP_017143669.1| PREDICTED: DENN domain-containing protein 4C isoform X2 [Dr
>gi|1036718700|ref|XP_017143684.1| PREDICTED: chromatin assembly factor 1 subunit A-B [Drosoph
>gi|1036706368|ref|XP_017136253.1| PREDICTED: rab-like protein 3 [Drosophila miranda]
>gi|1036746562|ref|XP_017156115.1| PREDICTED: homeotic protein ocelliless [Drosophila miranda]
>gi|1036745865|ref|XP_017156071.1| PREDICTED: uncharacterized protein LOC108164709 isoform X1
>gi|1036745884|ref|XP_017156072.1| PREDICTED: uncharacterized protein LOC108164709 isoform X2
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>gi|1036736644|ref|XP_017153461.1| PREDICTED: carbonic anhydrase 2 [Drosophila miranda]
>gi|1036736626|ref|XP_017153450.1| PREDICTED: palmitoyl-protein thioesterase 1 [Drosophila mira
>gi|1036705706|ref|XP_017135771.1| PREDICTED: paired mesoderm homeobox protein 2 [Drosophila mi
>gi|1036740507|ref|XP_017155546.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosami
>gi|1036740523|ref|XP_017155558.1| PREDICTED: uncharacterized protein LOC108164404 [Drosophila
>gi|1036731792|ref|XP_017150837.1| PREDICTED: uncharacterized protein CG1785 [Drosophila miran
>gi|1036731754|ref|XP_017150829.1| PREDICTED: RNA cytidine acetyltransferase isoform X1 [Drosp
>gi|1036731773|ref|XP_017150830.1| PREDICTED: RNA cytidine acetyltransferase isoform X2 [Drosp
>gi|1036702255|ref|XP_017156245.1| PREDICTED: probable U3 small nucleolar RNA-associated prote
>gi|1036705880|ref|XP_017135899.1| PREDICTED: uncharacterized protein LOC108151668 [Drosophila
>gi|1036747447|ref|XP_017156169.1| PREDICTED: LIM/homeobox protein Lhx1 [Drosophila miranda]
>gi|1036706758|ref|XP_017136542.1| PREDICTED: uncharacterized protein LOC108152047 [Drosophila
>gi|1036706774|ref|XP_017136550.1| PREDICTED: uncharacterized protein LOC108152047 [Drosophila
>gi|1036749202|ref|XP_017156253.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D
>gi|1036749220|ref|XP_017156254.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D
>gi|1036729431|ref|XP_017149561.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036729452|ref|XP_017149569.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036729471|ref|XP_017149578.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036729491|ref|XP_017149586.1| PREDICTED: moesin/ezrin/radixin homolog 1 [Drosophila miran
>gi|1036753967|ref|XP_017156530.1| PREDICTED: uroporphyrinogen-III synthase [Drosophila miranda
>gi|1036753984|ref|XP_017156531.1| PREDICTED: uncharacterized protein LOC108165008 [Drosophila
>gi|1036756564|ref|XP_017156677.1| PREDICTED: calumenin-like [Drosophila miranda]
>gi|1036745439|ref|XP_017156047.1| PREDICTED: uncharacterized protein LOC108164688 [Drosophila
>gi|1036745457|ref|XP_017156048.1| PREDICTED: uncharacterized protein LOC108164688 [Drosophila
>gi|1036735406|ref|XP_017152756.1| PREDICTED: protein MAK16 homolog [Drosophila miranda]
>gi|1036735424|ref|XP_017152769.1| PREDICTED: protein enhancer of rudimentary [Drosophila mira
>gi|1036712278|ref|XP_017139951.1| PREDICTED: eye-specific diacylglycerol kinase isoform X14 [
>gi|1036712430|ref|XP_017140039.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036712445|ref|XP_017140051.1| PREDICTED: uncharacterized protein LOC108154304 isoform X1
>gi|1036712462|ref|XP_017140059.1| PREDICTED: uncharacterized protein LOC108154304 isoform X2
>gi|1036712115|ref|XP_017139878.1| PREDICTED: eye-specific diacylglycerol kinase isoform X5 [D
>gi|1036712165|ref|XP_017139904.1| PREDICTED: eye-specific diacylglycerol kinase isoform X8 [D
>gi|1036712049|ref|XP_017139841.1| PREDICTED: eye-specific diacylglycerol kinase isoform X1 [D
>gi|1036712132|ref|XP_017139887.1| PREDICTED: eye-specific diacylglycerol kinase isoform X6 [D
>gi|1036712082|ref|XP_017139856.1| PREDICTED: eye-specific diacylglycerol kinase isoform X3 [D
>gi|1036712099|ref|XP_017139867.1| PREDICTED: eye-specific diacylglycerol kinase isoform X4 [D
>gi|1036712067|ref|XP_017139847.1| PREDICTED: eye-specific diacylglycerol kinase isoform X2 [D
>gi|1036712148|ref|XP_017139897.1| PREDICTED: eye-specific diacylglycerol kinase isoform X7 [D
>gi|1036712182|ref|XP_017139913.1| PREDICTED: eye-specific diacylglycerol kinase isoform X9 [D
>gi|1036712247|ref|XP_017139937.1| PREDICTED: eye-specific diacylglycerol kinase isoform X12 [
>gi|1036712199|ref|XP_017139922.1| PREDICTED: eye-specific diacylglycerol kinase isoform X10 [
>gi|1036712216|ref|XP_017139930.1| PREDICTED: eye-specific diacylglycerol kinase isoform X11 [
>gi|1036712396|ref|XP_017140016.1| PREDICTED: nucleosome assembly protein 1-like 1 [Drosophila
>gi|1036712412|ref|XP_017140027.1| PREDICTED: nucleosome assembly protein 1-like 1 [Drosophila
>gi|1036712261|ref|XP_017139944.1| PREDICTED: eye-specific diacylglycerol kinase isoform X13 [
>gi|1036712330|ref|XP_017139978.1| PREDICTED: nostrin isoform X3 [Drosophila miranda]
>gi|1036712364|ref|XP_017139993.1| PREDICTED: nostrin isoform X5 [Drosophila miranda]
>gi|1036712347|ref|XP_017139985.1| PREDICTED: nostrin isoform X4 [Drosophila miranda]
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>gi|1036712381|ref|XP_017140002.1| PREDICTED: nostrin isoform X6 [Drosophila miranda]
>gi|1036712295|ref|XP_017139965.1| PREDICTED: uncharacterized protein LOC108154240 isoform X1
>gi|1036712313|ref|XP_017139973.1| PREDICTED: uncharacterized protein LOC108154240 isoform X2
>gi|1036704189|ref|XP_017134744.1| PREDICTED: uncharacterized protein DDB_G0271670-like [Droso
>gi|1036753292|ref|XP_017156490.1| PREDICTED: transcription factor SPT20 homolog isoform X2 [D
>gi|1036753274|ref|XP_017156489.1| PREDICTED: transcription factor SPT20 homolog isoform X1 [D
>gi|1036749623|ref|XP_017156275.1| PREDICTED: acylphosphatase-1 [Drosophila miranda]
>gi|1036749640|ref|XP_017156276.1| PREDICTED: acylphosphatase-1 [Drosophila miranda]
>gi|1036749604|ref|XP_017156274.1| PREDICTED: uncharacterized protein LOC108164856 [Drosophila
>gi|1036749657|ref|XP_017156277.1| PREDICTED: uncharacterized protein LOC108164859 [Drosophila
>gi|1036724197|ref|XP_017146421.1| PREDICTED: U3 small nucleolar RNA-associated protein 6 homol
>gi|1036724233|ref|XP_017146446.1| PREDICTED: chromobox protein homolog 3-like [Drosophila mira
>gi|1036724160|ref|XP_017146396.1| PREDICTED: anaphase-promoting complex subunit 4 isoform X1
>gi|1036724179|ref|XP_017146406.1| PREDICTED: anaphase-promoting complex subunit 4 isoform X2
>gi|1036724212|ref|XP_017146435.1| PREDICTED: uncharacterized protein F54F2.9 [Drosophila mira
>gi|1036746844|ref|XP_017156133.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036749076|ref|XP_017156246.1| PREDICTED: band 7 protein AGAP004871 isoform X1 [Drosophila
>gi|1036749095|ref|XP_017156247.1| PREDICTED: band 7 protein AGAP004871 isoform X2 [Drosophila
>gi|1036749059|ref|XP_017156244.1| PREDICTED: band 7 protein AGAP004871 [Drosophila miranda]
>gi|1036749114|ref|XP_017156248.1| PREDICTED: NADH-cytochrome b5 reductase-like [Drosophila mi
>gi|1036732084|ref|XP_017150975.1| PREDICTED: transmembrane protein 185B [Drosophila miranda]
>gi|1036732121|ref|XP_017150995.1| PREDICTED: barrier-to-autointegration factor-like [Drosophil
>gi|1036732103|ref|XP_017150985.1| PREDICTED: LOW QUALITY PROTEIN: post-GPI attachment to prote
>gi|1036732161|ref|XP_017151012.1| PREDICTED: barrier-to-autointegration factor-like isoform X
>gi|1036732182|ref|XP_017151019.1| PREDICTED: barrier-to-autointegration factor-like isoform X
>gi|1036732142|ref|XP_017151005.1| PREDICTED: barrier-to-autointegration factor-like isoform X
>gi|1036745683|ref|XP_017156062.1| PREDICTED: post-GPI attachment to proteins factor 2-like iso
>gi|1036745702|ref|XP_017156064.1| PREDICTED: post-GPI attachment to proteins factor 2-like iso
>gi|1036728207|ref|XP_017148840.1| PREDICTED: uncharacterized protein LOC108159778 isoform X2
>gi|1036728190|ref|XP_017148831.1| PREDICTED: uncharacterized protein LOC108159778 isoform X1
>gi|1036756844|ref|XP_017156693.1| PREDICTED: uncharacterized protein LOC108165139 [Drosophila
>gi|1036755160|ref|XP_017156600.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036754693|ref|XP_017156572.1| PREDICTED: dentin sialophosphoprotein [Drosophila miranda]
>gi|1036744419|ref|XP_017155991.1| PREDICTED: uncharacterized protein LOC108164650 [Drosophila
>gi|1036744162|ref|XP_017155976.1| PREDICTED: uncharacterized protein LOC108164641 isoform X1
>gi|1036744181|ref|XP_017155977.1| PREDICTED: uncharacterized protein LOC108164641 isoform X2
>gi|1036744200|ref|XP_017155978.1| PREDICTED: uncharacterized protein LOC108164641 isoform X3
>gi|1036744219|ref|XP_017155979.1| PREDICTED: uncharacterized protein LOC108164641 isoform X3
>gi|1036744236|ref|XP_017155980.1| PREDICTED: uncharacterized protein LOC108164641 isoform X4
>gi|1036744137|ref|XP_017155975.1| PREDICTED: uncharacterized protein LOC108164640 [Drosophila
>gi|1036751381|ref|XP_017156374.1| PREDICTED: uncharacterized protein LOC108164907 [Drosophila
>gi|1036751398|ref|XP_017156375.1| PREDICTED: uncharacterized protein LOC108164907 [Drosophila
>gi|1036751417|ref|XP_017156376.1| PREDICTED: uncharacterized protein LOC108164907 [Drosophila
>gi|1036732516|ref|XP_017151179.1| PREDICTED: uncharacterized protein LOC108161426 [Drosophila
>gi|1036755749|ref|XP_017156631.1| PREDICTED: monocarboxylate transporter 7 isoform X1 [Drosop
>gi|1036755768|ref|XP_017156632.1| PREDICTED: monocarboxylate transporter 7 isoform X2 [Drosop
>gi|1036746524|ref|XP_017156113.1| PREDICTED: receptor-binding cancer antigen expressed on SiSo
>gi|1036746505|ref|XP_017156112.1| PREDICTED: uncharacterized protein LOC108164730 [Drosophila
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>gi|1036736451|ref|XP_017153348.1| PREDICTED: protein vav isoform X2 [Drosophila miranda]
>gi|1036736433|ref|XP_017153341.1| PREDICTED: protein vav isoform X1 [Drosophila miranda]
>gi|1036736470|ref|XP_017153360.1| PREDICTED: uncharacterized protein LOC108162898 [Drosophila
>gi|1036731700|ref|XP_017150826.1| PREDICTED: rapamycin-insensitive companion of mTOR [Drosophi
>gi|1036731718|ref|XP_017150827.1| PREDICTED: rapamycin-insensitive companion of mTOR [Drosophi
>gi|1036754832|ref|XP_017156581.1| PREDICTED: heparan sulfate glucosamine 3-O-sulfotransferase
>gi|1036720563|ref|XP_017144695.1| PREDICTED: uncharacterized protein LOC108157237 [Drosophila
>gi|1036720585|ref|XP_017144701.1| PREDICTED: uncharacterized protein LOC108157237 [Drosophila
>gi|1036720621|ref|XP_017144726.1| PREDICTED: protein-lysine N-methyltransferase EEF2KMT [Droso
>gi|1036720545|ref|XP_017144680.1| PREDICTED: pre-mRNA-splicing factor ATP-dependent RNA helica
>gi|1036720639|ref|XP_017144739.1| PREDICTED: uncharacterized protein LOC108157265 isoform X1
>gi|1036720658|ref|XP_017144748.1| PREDICTED: uncharacterized protein LOC108157265 isoform X2
>gi|1036720676|ref|XP_017144757.1| PREDICTED: uncharacterized protein LOC108157265 isoform X2
>gi|1036720605|ref|XP_017144714.1| PREDICTED: tenascin [Drosophila miranda]
>gi|1036730325|ref|XP_017150131.1| PREDICTED: 39S ribosomal protein L38, mitochondrial [Drosoph
>gi|1036730268|ref|XP_017150097.1| PREDICTED: type I inositol 3,4-bisphosphate 4-phosphatase [D
>gi|1036730287|ref|XP_017150106.1| PREDICTED: type I inositol 3,4-bisphosphate 4-phosphatase [D
>gi|1036730306|ref|XP_017150119.1| PREDICTED: F-box/LRR-repeat protein 4 [Drosophila miranda]
>gi|1036713343|ref|XP_017140558.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036713360|ref|XP_017140568.1| PREDICTED: neurogenic protein mastermind isoform X3 [Drosoph
>gi|1036713327|ref|XP_017140549.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036713376|ref|XP_017140576.1| PREDICTED: neurogenic protein mastermind isoform X4 [Drosoph
>gi|1036713393|ref|XP_017140588.1| PREDICTED: UMP-CMP kinase 2-like [Drosophila miranda]
>gi|1036713410|ref|XP_017140600.1| PREDICTED: protein Turandot Z [Drosophila miranda]
>gi|1036738073|ref|XP_017154255.1| PREDICTED: ras-related protein Ral-a isoform X2 [Drosophila
>gi|1036738037|ref|XP_017154241.1| PREDICTED: ras-related protein Ral-a isoform X1 [Drosophila
>gi|1036738055|ref|XP_017154249.1| PREDICTED: ras-related protein Ral-a isoform X1 [Drosophila
>gi|1036742862|ref|XP_017155910.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X2
>gi|1036742843|ref|XP_017155909.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036743462|ref|XP_017155945.1| PREDICTED: salivary glue protein Sgs-4-like [Drosophila mira
>gi|1036742339|ref|XP_017155880.1| PREDICTED: poly(ADP-ribose) glycohydrolase [Drosophila miran
>gi|1036742355|ref|XP_017155881.1| PREDICTED: probable insulin-like peptide 7 [Drosophila miran
>gi|1036706078|ref|XP_017136061.1| PREDICTED: uncharacterized protein LOC108151746 [Drosophila
>gi|1036754294|ref|XP_017156549.1| PREDICTED: allatostatin-A receptor [Drosophila miranda]
>gi|1036744015|ref|XP_017155971.1| PREDICTED: nucleolar complex protein 4 homolog B [Drosophila
>gi|1036743998|ref|XP_017155970.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1036750244|ref|XP_017156310.1| PREDICTED: limbic system-associated membrane protein [Drosop
>gi|1036750279|ref|XP_017156312.1| PREDICTED: uncharacterized protein LOC108164877 isoform X2
>gi|1036750295|ref|XP_017156313.1| PREDICTED: uncharacterized protein LOC108164877 isoform X2
>gi|1036750314|ref|XP_017156314.1| PREDICTED: uncharacterized protein LOC108164877 isoform X3
>gi|1036750263|ref|XP_017156311.1| PREDICTED: uncharacterized protein LOC108164877 isoform X1
>gi|1036739123|ref|XP_017154793.1| PREDICTED: growth arrest-specific protein 8 homolog [Drosoph
>gi|1036739158|ref|XP_017154806.1| PREDICTED: uncharacterized protein LOC108163820 isoform X2
>gi|1036739141|ref|XP_017154799.1| PREDICTED: uncharacterized protein LOC108163820 isoform X1
>gi|1036746952|ref|XP_017156140.1| PREDICTED: neurogenic protein mastermind [Drosophila miranda
>gi|1036746916|ref|XP_017156138.1| PREDICTED: uncharacterized protein LOC108164750 isoform X2
>gi|1036746899|ref|XP_017156136.1| PREDICTED: uncharacterized protein LOC108164750 isoform X1
>gi|1036746933|ref|XP_017156139.1| PREDICTED: voltage-gated potassium channel subunit beta-2-li
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>gi|1036756342|ref|XP_017156664.1| PREDICTED: uncharacterized protein LOC108165117 [Drosophila
>gi|1036719606|ref|XP_017144204.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719625|ref|XP_017144213.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719645|ref|XP_017144222.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719586|ref|XP_017144197.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719665|ref|XP_017144230.1| PREDICTED: mannosyl-oligosaccharide alpha-1,2-mannosidase IA
>gi|1036719704|ref|XP_017144256.1| PREDICTED: activating signal cointegrator 1 [Drosophila mira
>gi|1036719749|ref|XP_017144265.1| PREDICTED: migration and invasion enhancer 1 [Drosophila mi
>gi|1036719685|ref|XP_017144244.1| PREDICTED: cactin [Drosophila miranda]
>gi|1036742500|ref|XP_017155889.1| PREDICTED: molybdenum cofactor sulfurase isoform X2 [Drosop
>gi|1036742482|ref|XP_017155888.1| PREDICTED: molybdenum cofactor sulfurase isoform X1 [Drosop
>gi|1036754003|ref|XP_017156532.1| PREDICTED: small G protein signaling modulator 1 isoform X1
>gi|1036754019|ref|XP_017156533.1| PREDICTED: small G protein signaling modulator 1 isoform X2
>gi|1036752077|ref|XP_017156414.1| PREDICTED: LOW QUALITY PROTEIN: small G protein signaling mo
>gi|1036752096|ref|XP_017156415.1| PREDICTED: glutathione S-transferase theta-1 [Drosophila mi
>gi|1036720782|ref|XP_017144813.1| PREDICTED: uncharacterized protein LOC108157307 isoform X2
>gi|1036720800|ref|XP_017144822.1| PREDICTED: uncharacterized protein LOC108157307 isoform X3
>gi|1036720766|ref|XP_017144804.1| PREDICTED: uncharacterized protein LOC108157307 isoform X1
>gi|1036720817|ref|XP_017144830.1| PREDICTED: uncharacterized protein LOC108157307 isoform X4
>gi|1036725719|ref|XP_017147323.1| PREDICTED: kelch-like protein 22 [Drosophila miranda]
>gi|1036725702|ref|XP_017147311.1| PREDICTED: mitogen-activated protein kinase kinase kinase 7
>gi|1036725736|ref|XP_017147337.1| PREDICTED: uncharacterized protein LOC108158994 [Drosophila
>gi|1036752249|ref|XP_017156424.1| PREDICTED: uncharacterized protein LOC108164946 [Drosophila
>gi|1036757833|ref|XP_017156744.1| PREDICTED: uncharacterized protein LOC108165185 [Drosophila
>gi|1036757852|ref|XP_017156745.1| PREDICTED: uncharacterized protein LOC108165186 [Drosophila
>gi|1036757774|ref|XP_017156741.1| PREDICTED: uncharacterized protein LOC108165182 [Drosophila
>gi|1036704333|ref|XP_017134824.1| PREDICTED: runt-related transcription factor 3 isoform X2 [
>gi|1036704316|ref|XP_017134815.1| PREDICTED: protein lozenge isoform X1 [Drosophila miranda]
>gi|1036745942|ref|XP_017156076.1| PREDICTED: probable cytochrome P450 6v1 isoform X1 [Drosoph
>gi|1036745960|ref|XP_017156077.1| PREDICTED: probable cytochrome P450 6v1 isoform X2 [Drosoph
>gi|1036750556|ref|XP_017156326.1| PREDICTED: segmentation protein Runt isoform X1 [Drosophila
>gi|1036750575|ref|XP_017156327.1| PREDICTED: segmentation protein Runt isoform X2 [Drosophila
>gi|1036756015|ref|XP_017156645.1| PREDICTED: uncharacterized protein LOC108165101 [Drosophila
>gi|1036756034|ref|XP_017156646.1| PREDICTED: uncharacterized protein LOC108165101 [Drosophila
>gi|1036703603|ref|XP_017157044.1| PREDICTED: uncharacterized protein LOC108165494 [Drosophila
>gi|1036739559|ref|XP_017155056.1| PREDICTED: innexin shaking-B isoform X1 [Drosophila miranda]
>gi|1036739595|ref|XP_017155073.1| PREDICTED: innexin shaking-B isoform X3 [Drosophila miranda]
>gi|1036739577|ref|XP_017155065.1| PREDICTED: innexin shaking-B isoform X2 [Drosophila miranda]
>gi|1036739528|ref|XP_017155040.1| PREDICTED: glycerol-3-phosphate acyltransferase 3 [Drosophil
>gi|1036739543|ref|XP_017155048.1| PREDICTED: glycerol-3-phosphate acyltransferase 3 [Drosophil
>gi|1036739508|ref|XP_017155028.1| PREDICTED: uncharacterized protein LOC108163978 [Drosophila
>gi|1036739612|ref|XP_017155084.1| PREDICTED: protein phosphatase 1 regulatory subunit 11 [Dros
>gi|1036699425|ref|XP_017141085.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036755626|ref|XP_017156623.1| PREDICTED: Niemann-Pick C1 protein [Drosophila miranda]
>gi|1036703095|ref|XP_017156755.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036757682|ref|XP_017156737.1| PREDICTED: uncharacterized protein LOC108165177 [Drosophila
>gi|1036748966|ref|XP_017156239.1| PREDICTED: uncharacterized protein LOC108164827 [Drosophila
>gi|1036748983|ref|XP_017156240.1| PREDICTED: glyoxalase 1 [Drosophila miranda]
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>gi|1036730613|ref|XP_017150288.1| PREDICTED: protein Cep78 homolog isoform X3 [Drosophila mira
>gi|1036730594|ref|XP_017150281.1| PREDICTED: protein Cep78 homolog isoform X2 [Drosophila mira
>gi|1036730575|ref|XP_017150272.1| PREDICTED: protein Cep78 homolog isoform X1 [Drosophila mira
>gi|1036730743|ref|XP_017150372.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036730651|ref|XP_017150312.1| PREDICTED: uncharacterized protein LOC108160674 [Drosophila
>gi|1036730632|ref|XP_017150300.1| PREDICTED: uncharacterized protein LOC108160665 [Drosophila
>gi|1036699442|ref|XP_017141213.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosopl
>gi|1036730669|ref|XP_017150326.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosopl
>gi|1036730707|ref|XP_017150348.1| PREDICTED: protein NipSnap [Drosophila miranda]
>gi|1036730726|ref|XP_017150359.1| PREDICTED: protein NipSnap [Drosophila miranda]
>gi|1036730688|ref|XP_017150336.1| PREDICTED: zinc finger matrin-type protein 3 [Drosophila mi
>gi|1036752936|ref|XP_017156468.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miranc
>gi|1036702288|ref|XP_017156267.1| PREDICTED: UPF0488 protein CG14286 [Drosophila miranda]
>gi|1036721260|ref|XP_017145024.1| PREDICTED: protein Tob1 isoform X2 [Drosophila miranda]
>gi|1036721133|ref|XP_017144968.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721153|ref|XP_017144976.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721171|ref|XP_017144983.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721187|ref|XP_017144990.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721203|ref|XP_017144998.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721222|ref|XP_017145007.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721241|ref|XP_017145016.1| PREDICTED: protein Tob1 isoform X1 [Drosophila miranda]
>gi|1036721078|ref|XP_017144940.1| PREDICTED: uncharacterized protein LOC108157404 [Drosophila
>gi|1036721095|ref|XP_017144952.1| PREDICTED: uncharacterized protein LOC108157410 [Drosophila
>gi|1036721114|ref|XP_017144959.1| PREDICTED: uncharacterized protein LOC108157410 [Drosophila
>gi|1036721063|ref|XP_017144928.1| PREDICTED: putative uncharacterized protein DDB_G0291608 [D
>gi|1036757373|ref|XP_017156722.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosopl
>gi|1036704496|ref|XP_017134931.1| PREDICTED: fork head domain transcription factor slp1, part
>gi|1036747428|ref|XP_017156168.1| PREDICTED: uncharacterized protein LOC108164767 [Drosophila
>gi|1036749002|ref|XP_017156241.1| PREDICTED: protein disconnected [Drosophila miranda]
>gi|1036705986|ref|XP_017135981.1| PREDICTED: uncharacterized protein LOC108151716 isoform X1
>gi|1036706007|ref|XP_017135989.1| PREDICTED: uncharacterized protein LOC108151716 isoform X2
>gi|1036699457|ref|XP_017141336.1| PREDICTED: uncharacterized protein LOC108155149 [Drosophila
>gi|1036732797|ref|XP_017151323.1| PREDICTED: katanin p80 WD40 repeat-containing subunit B1 iso
>gi|1036732778|ref|XP_017151317.1| PREDICTED: katanin p80 WD40 repeat-containing subunit B1 iso
>gi|1036732815|ref|XP_017151335.1| PREDICTED: peroxisomal multifunctional enzyme type 2 [Drosop
>gi|1036732834|ref|XP_017151343.1| PREDICTED: peroxisomal multifunctional enzyme type 2 [Drosop
>gi|1036746489|ref|XP_017156111.1| PREDICTED: calpain-C [Drosophila miranda]
>gi|1036744677|ref|XP_017156003.1| PREDICTED: uncharacterized protein LOC108164661 [Drosophila
>gi|1036744695|ref|XP_017156004.1| PREDICTED: HIG1 domain family member 2A, mitochondrial [Dros
>gi|1036741037|ref|XP_017155803.1| PREDICTED: high mobility group protein DSP1 isoform X1 [Dros
>gi|1036741055|ref|XP_017155804.1| PREDICTED: high mobility group protein DSP1 isoform X2 [Dros
>gi|1036741072|ref|XP_017155805.1| PREDICTED: high mobility group protein DSP1 isoform X3 [Dros
>gi|1036741089|ref|XP_017155807.1| PREDICTED: high mobility group protein DSP1 isoform X4 [Dros
>gi|1036717210|ref|XP_017142880.1| PREDICTED: acyl-CoA Delta(11) desaturase-like [Drosophila mi
>gi|1036717122|ref|XP_017142829.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036717194|ref|XP_017142868.1| PREDICTED: splicing factor U2AF 50 kDa subunit [Drosophila n
>gi|1036717138|ref|XP_017142840.1| PREDICTED: uncharacterized protein LOC108156065 [Drosophila
>gi|1036717154|ref|XP_017142848.1| PREDICTED: uncharacterized protein LOC108156065 [Drosophila
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>gi|1036717171|ref|XP_017142856.1| PREDICTED: uncharacterized protein LOC108156065 [Drosophila
>gi|1036729166|ref|XP_017149407.1| PREDICTED: uncharacterized protein LOC108160122 [Drosophila
>gi|1036729148|ref|XP_017149396.1| PREDICTED: protein no-on-transient A [Drosophila miranda]
>gi|1036723090|ref|XP_017145799.1| PREDICTED: furin-like protease 2 isoform X1 [Drosophila mira
>gi|1036723112|ref|XP_017145808.1| PREDICTED: furin-like protease 2 isoform X1 [Drosophila mira
>gi|1036723130|ref|XP_017145820.1| PREDICTED: furin-like protease 2 isoform X2 [Drosophila mira
>gi|1036723148|ref|XP_017145831.1| PREDICTED: furin-like protease 2 isoform X3 [Drosophila mira
>gi|1036723167|ref|XP_017145840.1| PREDICTED: furin-like protease 2 isoform X4 [Drosophila mira
>gi|1036723203|ref|XP_017145861.1| PREDICTED: uncharacterized protein LOC108158142 isoform X2
>gi|1036723186|ref|XP_017145853.1| PREDICTED: uncharacterized protein LOC108158142 isoform X1
>gi|1036723214|ref|XP_017145869.1| PREDICTED: uncharacterized protein LOC108158142 isoform X3
>gi|1036736680|ref|XP_017153482.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736698|ref|XP_017153491.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736714|ref|XP_017153499.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736733|ref|XP_017153508.1| PREDICTED: trimeric intracellular cation channel type B-B [
>gi|1036736663|ref|XP_017153471.1| PREDICTED: negative elongation factor D [Drosophila miranda]
>gi|1036756892|ref|XP_017156698.1| PREDICTED: probable phospholipid-transporting ATPase IF [Dro
>gi|1036751992|ref|XP_017156409.1| PREDICTED: probable phospholipid-transporting ATPase IF [Dro
>gi|1036747113|ref|XP_017156149.1| PREDICTED: TNF receptor-associated factor 3 isoform X1 [Dros
>gi|1036747131|ref|XP_017156151.1| PREDICTED: TNF receptor-associated factor 3 isoform X2 [Dros
>gi|1036747097|ref|XP_017156148.1| PREDICTED: annexin B11 isoform X2 [Drosophila miranda]
>gi|1036747062|ref|XP_017156146.1| PREDICTED: annexin B11 isoform X1 [Drosophila miranda]
>gi|1036747078|ref|XP_017156147.1| PREDICTED: annexin B11 isoform X1 [Drosophila miranda]
>gi|1036733599|ref|XP_017151723.1| PREDICTED: gamma-glutamyltranspeptidase 1 [Drosophila miranc
>gi|1036733650|ref|XP_017151741.1| PREDICTED: gamma-glutamyltranspeptidase 1 [Drosophila miranc
>gi|1036733690|ref|XP_017151759.1| PREDICTED: uncharacterized protein LOC108161868 isoform X2
>gi|1036733669|ref|XP_017151752.1| PREDICTED: uncharacterized protein LOC108161868 isoform X1
>gi|1036733709|ref|XP_017151771.1| PREDICTED: uncharacterized protein LOC108161884 [Drosophila
>gi|1036733561|ref|XP_017151703.1| PREDICTED: sushi, von Willebrand factor type A, EGF and pent
>gi|1036733581|ref|XP_017151710.1| PREDICTED: sushi, von Willebrand factor type A, EGF and pent
>gi|1036756308|ref|XP_017156662.1| PREDICTED: fibrous sheath CABYR-binding protein [Drosophila
>gi|1036746809|ref|XP_017156131.1| PREDICTED: serine protease gd [Drosophila miranda]
>gi|1036746826|ref|XP_017156132.1| PREDICTED: protein twisted gastrulation [Drosophila miranda]
>gi|1036755307|ref|XP_017156608.1| PREDICTED: uncharacterized protein LOC108165071 [Drosophila
>gi|1036755325|ref|XP_017156610.1| PREDICTED: uncharacterized protein LOC108165071 [Drosophila
>gi|1036717982|ref|XP_017143303.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718002|ref|XP_017143313.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718021|ref|XP_017143323.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718059|ref|XP_017143341.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718040|ref|XP_017143331.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718076|ref|XP_017143349.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718113|ref|XP_017143364.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718095|ref|XP_017143356.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718128|ref|XP_017143372.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718147|ref|XP_017143378.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718166|ref|XP_017143387.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718188|ref|XP_017143393.1| PREDICTED: voltage-dependent calcium channel type A subunit
>gi|1036718207|ref|XP_017143400.1| PREDICTED: voltage-dependent calcium channel type A subunit
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>gi|1036754527|ref|XP_017156563.1| PREDICTED: PDZ domain-containing protein 8 [Drosophila mira
>gi|1036756053|ref|XP_017156647.1| PREDICTED: leucine-rich repeat-containing protein 20 isoform
>gi|1036756072|ref|XP_017156648.1| PREDICTED: leucine-rich repeat-containing protein 20 isoform
>gi|1036703879|ref|XP_017134567.1| PREDICTED: leucine-rich repeat-containing protein 69-like is
>gi|1036703930|ref|XP_017134588.1| PREDICTED: protein lap1-like isoform X4 [Drosophila miranda]
>gi|1036703913|ref|XP_017134582.1| PREDICTED: leucine-rich repeat-containing protein 69-like is
>gi|1036703896|ref|XP_017134574.1| PREDICTED: leucine-rich repeat-containing protein 69-like is
>gi|1036748107|ref|XP_017156204.1| PREDICTED: uncharacterized protein LOC108164792 [Drosophila
>gi|1036748088|ref|XP_017156203.1| PREDICTED: uncharacterized protein LOC108164791 [Drosophila
>gi|1036754275|ref|XP_017156548.1| PREDICTED: probable cytochrome P450 28a5 [Drosophila miranda
>gi|1036732315|ref|XP_017151082.1| PREDICTED: protein xmas-2 [Drosophila miranda]
>gi|1036732332|ref|XP_017151094.1| PREDICTED: uncharacterized protein LOC108161359 [Drosophila
>gi|1036699471|ref|XP_017141454.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036729976|ref|XP_017149886.1| PREDICTED: uncharacterized protein LOC108160402 [Drosophila
>gi|1036699488|ref|XP_017141582.1| PREDICTED: uncharacterized protein LOC108155322 [Drosophila
>gi|1036740746|ref|XP_017155712.1| PREDICTED: histone H1.3-like [Drosophila miranda]
>gi|1036740775|ref|XP_017155735.1| PREDICTED: histone H2B-like [Drosophila miranda]
>gi|1036740760|ref|XP_017155723.1| PREDICTED: coiled-coil-helix-coiled-coil-helix domain-contai
>gi|1036743427|ref|XP_017155943.1| PREDICTED: hexokinase type 2 [Drosophila miranda]
>gi|1036699502|ref|XP_017141705.1| PREDICTED: LOW QUALITY PROTEIN: protein vreteno-like [Drosop
>gi|1036727344|ref|XP_017148347.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036727361|ref|XP_017148355.1| PREDICTED: serine-rich adhesin for platelets isoform X3 [Dro
>gi|1036727295|ref|XP_017148325.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036727325|ref|XP_017148337.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036723885|ref|XP_017146206.1| PREDICTED: probable cytochrome P450 318a1 [Drosophila miran
>gi|1036723848|ref|XP_017146187.1| PREDICTED: protein strawberry notch isoform X1 [Drosophila
>gi|1036723867|ref|XP_017146196.1| PREDICTED: protein strawberry notch isoform X2 [Drosophila
>gi|1036723919|ref|XP_017146228.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036723902|ref|XP_017146217.1| PREDICTED: 2-methoxy-6-polyprenyl-1,4-benzoquinol methylase
>gi|1036699514|ref|XP_017141828.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036723936|ref|XP_017146241.1| PREDICTED: GTP-binding nuclear protein Ran-like [Drosophila
>gi|1036727144|ref|XP_017148209.1| PREDICTED: relaxin receptor 2 [Drosophila miranda]
>gi|1036727215|ref|XP_017148264.1| PREDICTED: uncharacterized protein LOC108159462 [Drosophila
>gi|1036727198|ref|XP_017148250.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 5 [
>gi|1036727181|ref|XP_017148235.1| PREDICTED: brahma-associated protein of 60 kDa [Drosophila
>gi|1036727162|ref|XP_017148223.1| PREDICTED: cleft lip and palate transmembrane protein 1-like
>gi|1036756184|ref|XP_017156655.1| PREDICTED: uncharacterized protein LOC108165108 [Drosophila
>gi|1036745049|ref|XP_017156024.1| PREDICTED: voltage-dependent T-type calcium channel subunit
>gi|1036745032|ref|XP_017156022.1| PREDICTED: voltage-dependent T-type calcium channel subunit
>gi|1036754602|ref|XP_017156567.1| PREDICTED: vanin-like protein 3 [Drosophila miranda]
>gi|1036754584|ref|XP_017156566.1| PREDICTED: vanin-like protein 1 [Drosophila miranda]
>gi|1036757265|ref|XP_017156715.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036756797|ref|XP_017156691.1| PREDICTED: vanin-like protein 2 [Drosophila miranda]
>gi|1036740626|ref|XP_017155631.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740645|ref|XP_017155638.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740661|ref|XP_017155648.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740677|ref|XP_017155656.1| PREDICTED: UDP-xylose and UDP-N-acetylglucosamine transporte
>gi|1036740693|ref|XP_017155669.1| PREDICTED: myosin regulatory light chain sqh [Drosophila mi
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>gi|1036727653|ref|XP_017148524.1| PREDICTED: transmembrane protein 132B [Drosophila miranda]
>gi|1036703295|ref|XP_017156876.1| PREDICTED: uncharacterized protein LOC108165351 isoform X1
>gi|1036703334|ref|XP_017156891.1| PREDICTED: uncharacterized protein LOC108165351 isoform X3
>gi|1036703369|ref|XP_017156906.1| PREDICTED: uncharacterized protein LOC108165351 isoform X5
>gi|1036703315|ref|XP_017156883.1| PREDICTED: uncharacterized protein LOC108165351 isoform X2
>gi|1036703354|ref|XP_017156899.1| PREDICTED: uncharacterized protein LOC108165351 isoform X4
>gi|1036729305|ref|XP_017149482.1| PREDICTED: dmX-like protein 2 [Drosophila miranda]
>gi|1036751755|ref|XP_017156394.1| PREDICTED: splicing factor 3B subunit 4 isoform X2 [Drosophi
>gi|1036751736|ref|XP_017156393.1| PREDICTED: splicing factor 3B subunit 4 isoform X1 [Drosophi
>gi|1036746971|ref|XP_017156141.1| PREDICTED: ceramide synthase 6 [Drosophila miranda]
>gi|1036746990|ref|XP_017156142.1| PREDICTED: josephin-like protein [Drosophila miranda]
>gi|1036724726|ref|XP_017146751.1| PREDICTED: transcription elongation factor SPT6 [Drosophila
>gi|1036747479|ref|XP_017156171.1| PREDICTED: uncharacterized protein LOC108164770 [Drosophila
>gi|1036747496|ref|XP_017156172.1| PREDICTED: uncharacterized protein LOC108164770 [Drosophila
>gi|1036747534|ref|XP_017156174.1| PREDICTED: cytochrome b5 isoform X2 [Drosophila miranda]
>gi|1036747515|ref|XP_017156173.1| PREDICTED: cytochrome b5 isoform X1 [Drosophila miranda]
>gi|1036747555|ref|XP_017156175.1| PREDICTED: probable cytochrome b5 isoform X3 [Drosophila mi
>gi|1036710068|ref|XP_017138615.1| PREDICTED: uncharacterized protein LOC108153261 [Drosophila
>gi|1036710166|ref|XP_017138682.1| PREDICTED: uncharacterized protein LOC108153299 [Drosophila
>gi|1036710183|ref|XP_017138692.1| PREDICTED: uncharacterized protein LOC108153299 [Drosophila
>gi|1036710113|ref|XP_017138650.1| PREDICTED: U4/U6.U5 tri-snRNP-associated protein 2 isoform
>gi|1036710130|ref|XP_017138660.1| PREDICTED: U4/U6.U5 tri-snRNP-associated protein 2 isoform
>gi|1036710098|ref|XP_017138642.1| PREDICTED: U4/U6.U5 tri-snRNP-associated protein 2 isoform
>gi|1036710215|ref|XP_017138717.1| PREDICTED: L-xylulose reductase [Drosophila miranda]
>gi|1036710083|ref|XP_017138628.1| PREDICTED: mucin-12 [Drosophila miranda]
>gi|1036710147|ref|XP_017138672.1| PREDICTED: probable sulfite oxidase, mitochondrial [Drosophi
>gi|1036710200|ref|XP_017138706.1| PREDICTED: proteasome subunit alpha type-7-1 [Drosophila mi
>gi|1036749844|ref|XP_017156288.1| PREDICTED: XK-related protein 6 isoform X2 [Drosophila mira
>gi|1036749826|ref|XP_017156287.1| PREDICTED: XK-related protein 6 isoform X1 [Drosophila mira
>gi|1036749375|ref|XP_017156263.1| PREDICTED: 27 kDa hemolymph protein [Drosophila miranda]
>gi|1036726872|ref|XP_017148064.1| PREDICTED: uncharacterized protein LOC108159354 [Drosophila
>gi|1036726909|ref|XP_017148080.1| PREDICTED: lambda-crystallin homolog [Drosophila miranda]
>gi|1036726852|ref|XP_017148050.1| PREDICTED: protein IWS1 homolog [Drosophila miranda]
>gi|1036749770|ref|XP_017156284.1| PREDICTED: WASH complex subunit 7 homolog isoform X1 [Droso
>gi|1036749789|ref|XP_017156285.1| PREDICTED: WASH complex subunit 7 homolog isoform X2 [Droso
>gi|1036745580|ref|XP_017156056.1| PREDICTED: peptidyl-prolyl cis-trans isomerase [Drosophila
>gi|1036738801|ref|XP_017154611.1| PREDICTED: myotubularin-related protein 4 isoform X1 [Droso
>gi|1036738817|ref|XP_017154619.1| PREDICTED: myotubularin-related protein 4 isoform X2 [Droso
>gi|1036738832|ref|XP_017154628.1| PREDICTED: myotubularin-related protein 4 isoform X2 [Droso
>gi|1036738849|ref|XP_017154637.1| PREDICTED: myotubularin-related protein 4 isoform X2 [Droso
>gi|1036738865|ref|XP_017154644.1| PREDICTED: myotubularin-related protein 4 isoform X3 [Droso
>gi|1036747408|ref|XP_017156166.1| PREDICTED: endoplasmic reticulum resident protein 44 [Droso
>gi|1036744291|ref|XP_017155983.1| PREDICTED: RNA-binding protein cabeza isoform X1 [Drosophila
>gi|1036744308|ref|XP_017155984.1| PREDICTED: RNA-binding protein cabeza isoform X2 [Drosophila
>gi|1036749023|ref|XP_017156242.1| PREDICTED: vesicle transport protein GOT1B [Drosophila mira
>gi|1036749041|ref|XP_017156243.1| PREDICTED: cytochrome b-c1 complex subunit 7 [Drosophila mi
>gi|1036747007|ref|XP_017156143.1| PREDICTED: ethanolamine kinase [Drosophila miranda]
>gi|1036747025|ref|XP_017156144.1| PREDICTED: ethanolamine kinase [Drosophila miranda]
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>gi|1036747044|ref|XP_017156145.1| PREDICTED: ethanolamine kinase [Drosophila miranda]
>gi|1036699531|ref|XP_017141956.1| PREDICTED: uncharacterized protein LOC108155546 [Drosophila
>gi|1036729267|ref|XP_017149466.1| PREDICTED: vascular endothelial growth factor C [Drosophila
>gi|1036729240|ref|XP_017149454.1| PREDICTED: zinc finger protein 596 [Drosophila miranda]
>gi|1036729203|ref|XP_017149433.1| PREDICTED: protein FAM188A homolog isoform X1 [Drosophila m:
>gi|1036729222|ref|XP_017149441.1| PREDICTED: protein FAM188A homolog isoform X2 [Drosophila m:
>gi|1036740872|ref|XP_017155796.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D:
>gi|1036740890|ref|XP_017155797.1| PREDICTED: zinc finger CCCH domain-containing protein 13 [D:
>gi|1036756671|ref|XP_017156684.1| PREDICTED: succinate dehydrogenase [ubiquinone] iron-sulfur
>gi|1036753001|ref|XP_017156472.1| PREDICTED: dual specificity protein phosphatase 3 isoform X:
>gi|1036753018|ref|XP_017156473.1| PREDICTED: dual specificity protein phosphatase 3 isoform X:
>gi|1036753037|ref|XP_017156474.1| PREDICTED: dual specificity protein phosphatase 3 isoform X:
>gi|1036753054|ref|XP_017156475.1| PREDICTED: dual specificity protein phosphatase 3 isoform X4
>gi|1036720526|ref|XP_017144670.1| PREDICTED: putative uncharacterized protein DDB_G0277255 [D:
>gi|1036706051|ref|XP_017136033.1| PREDICTED: uncharacterized protein LOC108151737 [Drosophila
>gi|1036755195|ref|XP_017156602.1| PREDICTED: uncharacterized protein LOC108165065 [Drosophila
>gi|1036705477|ref|XP_017135628.1| PREDICTED: uncharacterized protein LOC108151496 [Drosophila
>gi|1036705493|ref|XP_017135637.1| PREDICTED: uncharacterized protein LOC108151496 [Drosophila
>gi|1036699547|ref|XP_017142209.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036754112|ref|XP_017156538.1| PREDICTED: uncharacterized protein LOC108165013 [Drosophila
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>gi|1036754150|ref|XP_017156541.1| PREDICTED: uncharacterized protein LOC108165014 [Drosophila
>gi|1036703695|ref|XP_017157096.1| PREDICTED: myotrophin [Drosophila miranda]
>gi|1036703654|ref|XP_017157074.1| PREDICTED: uncharacterized protein LOC108165523 isoform X1
>gi|1036703678|ref|XP_017157084.1| PREDICTED: uncharacterized protein LOC108165523 isoform X2
>gi|1036706280|ref|XP_017136196.1| PREDICTED: cAMP-regulated phosphoprotein 19-A-like [Drosoph:
>gi|1036704972|ref|XP_017135276.1| PREDICTED: nuclear pore complex protein Nup50-like [Drosoph:
>gi|1036703843|ref|XP_017134547.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galact
>gi|1036703862|ref|XP_017134555.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galact
>gi|1036732909|ref|XP_017151377.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
>gi|1036732928|ref|XP_017151386.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
>gi|1036732951|ref|XP_017151396.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
>gi|1036733006|ref|XP_017151403.1| PREDICTED: uncharacterized protein LOC108161591 isoform X2
>gi|1036732890|ref|XP_017151370.1| PREDICTED: uncharacterized protein LOC108161591 isoform X1
>gi|1036733044|ref|XP_017151421.1| PREDICTED: vacuolar ATPase assembly integral membrane prote
>gi|1036733025|ref|XP_017151411.1| PREDICTED: innexin inx5 [Drosophila miranda]
>gi|1036704047|ref|XP_017134656.1| PREDICTED: uncharacterized protein LOC108150861 [Drosophila
>gi|1036704109|ref|XP_017134691.1| PREDICTED: uncharacterized protein LOC108150889 [Drosophila
>gi|1036735457|ref|XP_017152788.1| PREDICTED: uncharacterized protein LOC108162511 isoform X2
>gi|1036735440|ref|XP_017152781.1| PREDICTED: uncharacterized protein LOC108162511 isoform X1
>gi|1036725417|ref|XP_017147153.1| PREDICTED: uncharacterized protein LOC108158886 isoform X1
>gi|1036725435|ref|XP_017147164.1| PREDICTED: uncharacterized protein LOC108158886 isoform X1
>gi|1036725472|ref|XP_017147180.1| PREDICTED: uncharacterized protein LOC108158886 isoform X3
>gi|1036725491|ref|XP_017147189.1| PREDICTED: uncharacterized protein LOC108158886 isoform X4
>gi|1036725510|ref|XP_017147197.1| PREDICTED: uncharacterized protein LOC108158886 isoform X5
>gi|1036725452|ref|XP_017147172.1| PREDICTED: uncharacterized protein LOC108158886 isoform X2
>gi|1036725528|ref|XP_017147210.1| PREDICTED: lectin subunit alpha [Drosophila miranda]
>gi|1036739293|ref|XP_017154907.1| PREDICTED: serine protease snake [Drosophila miranda]
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>gi|1036739309|ref|XP_017154916.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1036739323|ref|XP_017154924.1| PREDICTED: serine protease snake [Drosophila miranda]
>gi|1036739278|ref|XP_017154896.1| PREDICTED: DNA replication licensing factor Mcm3 [Drosophil
>gi|1036743982|ref|XP_017155969.1| PREDICTED: protein DENND6B [Drosophila miranda]
>gi|1036741869|ref|XP_017155853.1| PREDICTED: DNA repair protein XRCC1 [Drosophila miranda]
>gi|1036741887|ref|XP_017155854.1| PREDICTED: calcineurin subunit B type 1 [Drosophila miranda]
>gi|1036741905|ref|XP_017155855.1| PREDICTED: calcineurin subunit B type 1 [Drosophila miranda]
>gi|1036741924|ref|XP_017155856.1| PREDICTED: calcineurin subunit B type 1 [Drosophila miranda]
>gi|1036742951|ref|XP_017155915.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036742969|ref|XP_017155917.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036742988|ref|XP_017155918.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743007|ref|XP_017155919.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743026|ref|XP_017155920.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743045|ref|XP_017155921.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743064|ref|XP_017155922.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743079|ref|XP_017155923.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743098|ref|XP_017155924.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036743117|ref|XP_017155925.1| PREDICTED: small conductance calcium-activated potassium cha
>gi|1036754489|ref|XP_017156560.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036723016|ref|XP_017145754.1| PREDICTED: A-kinase anchor protein 1, mitochondrial [Drosoph
>gi|1036723059|ref|XP_017145780.1| PREDICTED: 60S ribosomal protein L36 [Drosophila miranda]
>gi|1036723075|ref|XP_017145788.1| PREDICTED: 60S ribosomal protein L36 [Drosophila miranda]
>gi|1036723040|ref|XP_017145767.1| PREDICTED: histone H4 transcription factor [Drosophila mira
>gi|1036733112|ref|XP_017151453.1| PREDICTED: uncharacterized protein LOC108161665 [Drosophila
>gi|1036733131|ref|XP_017151467.1| PREDICTED: uncharacterized protein LOC108161671 [Drosophila
>gi|1036733095|ref|XP_017151447.1| PREDICTED: frizzled-4 [Drosophila miranda]
>gi|1036741962|ref|XP_017155858.1| PREDICTED: uncharacterized protein LOC108164555 isoform X1
>gi|1036741998|ref|XP_017155860.1| PREDICTED: uncharacterized protein LOC108164555 isoform X3
>gi|1036741981|ref|XP_017155859.1| PREDICTED: uncharacterized protein LOC108164555 isoform X2
>gi|1036757097|ref|XP_017156705.1| PREDICTED: uncharacterized protein LOC108165149 [Drosophila
>gi|1036703496|ref|XP_017156979.1| PREDICTED: thioredoxin domain-containing protein 9-like [Dr
>gi|1036731843|ref|XP_017150862.1| PREDICTED: zinc finger CCCH domain-containing protein 18 iso
>gi|1036731825|ref|XP_017150856.1| PREDICTED: zinc finger CCCH domain-containing protein 18 iso
>gi|1036731807|ref|XP_017150848.1| PREDICTED: zinc finger CCCH domain-containing protein 18 iso
>gi|1036753574|ref|XP_017156506.1| PREDICTED: protein ABHD11 [Drosophila miranda]
>gi|1036747291|ref|XP_017156161.1| PREDICTED: protein unc-119 homolog [Drosophila miranda]
>gi|1036743512|ref|XP_017155946.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1036744830|ref|XP_017156010.1| PREDICTED: autophagy protein 5 [Drosophila miranda]
>gi|1036736920|ref|XP_017153624.1| PREDICTED: rho GTPase-activating protein gacZ [Drosophila mi
>gi|1036752137|ref|XP_017156417.1| PREDICTED: acyl-CoA synthetase family member 3, mitochondria
>gi|1036752155|ref|XP_017156418.1| PREDICTED: venom allergen 5 [Drosophila miranda]
>gi|1036742807|ref|XP_017155906.1| PREDICTED: sarcoplasmic calcium-binding protein [Drosophila
>gi|1036742826|ref|XP_017155908.1| PREDICTED: uncharacterized protein LOC108164593 [Drosophila
>gi|1036705562|ref|XP_017135680.1| PREDICTED: uncharacterized protein LOC108151534 isoform X1
>gi|1036705592|ref|XP_017135689.1| PREDICTED: uncharacterized protein LOC108151534 isoform X2
>gi|1036705241|ref|XP_017135477.1| PREDICTED: uncharacterized protein LOC108151397 [Drosophila
>gi|1036705258|ref|XP_017135483.1| PREDICTED: uncharacterized protein LOC108151397 [Drosophila
>gi|1036703620|ref|XP_017157054.1| PREDICTED: uncharacterized protein LOC108165505 [Drosophila
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>gi|1036703637|ref|XP_017157061.1| PREDICTED: uncharacterized protein LOC108165505 [Drosophila
>gi|1036753884|ref|XP_017156525.1| PREDICTED: condensin complex subunit 2-like isoform X1 [Dros
>gi|1036753900|ref|XP_017156526.1| PREDICTED: condensin complex subunit 2-like isoform X2 [Dros
>gi|1036753916|ref|XP_017156527.1| PREDICTED: uncharacterized protein LOC108165003 isoform X3
>gi|1036705646|ref|XP_017135726.1| PREDICTED: uncharacterized protein LOC108151562 [Drosophila
>gi|1036706387|ref|XP_017136266.1| PREDICTED: uncharacterized protein LOC108151877 [Drosophila
>gi|1036705526|ref|XP_017135657.1| PREDICTED: uncharacterized protein LOC108151511 isoform X2
>gi|1036705509|ref|XP_017135649.1| PREDICTED: uncharacterized protein LOC108151511 isoform X1
>gi|1036757737|ref|XP_017156739.1| PREDICTED: uncharacterized protein LOC108165180 [Drosophila
>gi|1036705392|ref|XP_017135575.1| PREDICTED: uncharacterized protein LOC108151461 [Drosophila
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>gi|1036707574|ref|XP_017137057.1| PREDICTED: uncharacterized protein LOC108152322 [Drosophila
>gi|1036705003|ref|XP_017135302.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1036699562|ref|XP_017142462.1| PREDICTED: F-box/WD repeat-containing protein A [Drosophila
>gi|1036707369|ref|XP_017136915.1| PREDICTED: uncharacterized protein LOC108152233 [Drosophila
>gi|1036707163|ref|XP_017136794.1| PREDICTED: small nuclear ribonucleoprotein-associated prote
>gi|1036707420|ref|XP_017136949.1| PREDICTED: uncharacterized protein LOC108152250 [Drosophila
>gi|1036699577|ref|XP_017142588.1| PREDICTED: uncharacterized protein LOC108155912 [Drosophila
>gi|1036753311|ref|XP_017156491.1| PREDICTED: transcription factor GTE3, chloroplastic-like [D
>gi|1036753369|ref|XP_017156494.1| PREDICTED: uncharacterized protein LOC108164984 [Drosophila
>gi|1036753351|ref|XP_017156493.1| PREDICTED: uncharacterized protein LOC108164983 [Drosophila
>gi|1036753332|ref|XP_017156492.1| PREDICTED: uncharacterized protein LOC108164981 [Drosophila
>gi|1036699591|ref|XP_017142714.1| PREDICTED: uncharacterized protein LOC108155989 [Drosophila
>gi|1036756128|ref|XP_017156651.1| PREDICTED: uncharacterized protein LOC108165105 [Drosophila
>gi|1036705800|ref|XP_017135843.1| PREDICTED: uncharacterized protein LOC108151644 [Drosophila
>gi|1036705815|ref|XP_017135854.1| PREDICTED: uncharacterized protein LOC108151644 [Drosophila
>gi|1036735122|ref|XP_017152591.1| PREDICTED: homeobox protein cut isoform X1 [Drosophila mira
>gi|1036735140|ref|XP_017152600.1| PREDICTED: homeobox protein cut isoform X2 [Drosophila mira
>gi|1036735155|ref|XP_017152612.1| PREDICTED: LOW QUALITY PROTEIN: serine/threonine-protein ph
>gi|1036757227|ref|XP_017156712.1| PREDICTED: uncharacterized protein LOC108165157 [Drosophila
>gi|1036730870|ref|XP_017150446.1| PREDICTED: uncharacterized protein LOC108160754 [Drosophila
>gi|1036730889|ref|XP_017150453.1| PREDICTED: uncharacterized protein LOC108160754 [Drosophila
>gi|1036706590|ref|XP_017136397.1| PREDICTED: uncharacterized protein LOC108151962 [Drosophila
>gi|1036704207|ref|XP_017134751.1| PREDICTED: LOW QUALITY PROTEIN: AF4/FMR2 family member 4 [D
>gi|1036715798|ref|XP_017142050.1| PREDICTED: protein HIRA homolog [Drosophila miranda]
>gi|1036715815|ref|XP_017142063.1| PREDICTED: negative elongation factor B [Drosophila miranda]
>gi|1036715831|ref|XP_017142077.1| PREDICTED: uncharacterized protein LOC108155639 [Drosophila
>gi|1036715851|ref|XP_017142089.1| PREDICTED: pyruvate dehydrogenase [acetyl-transferring]-pho
>gi|1036715871|ref|XP_017142102.1| PREDICTED: ras-related protein Rab-39B [Drosophila miranda]
>gi|1036742374|ref|XP_017155882.1| PREDICTED: mitochondrial import receptor subunit TOM40 homo
>gi|1036758069|ref|XP_017156754.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036699606|ref|XP_017142847.1| PREDICTED: uncharacterized protein LOC108156047 [Drosophila
>gi|1036706649|ref|XP_017136459.1| PREDICTED: uncharacterized protein LOC108151998 [Drosophila
>gi|1036699621|ref|XP_017142968.1| PREDICTED: uncharacterized protein LOC108156128 [Drosophila
>gi|1036753796|ref|XP_017156519.1| PREDICTED: venom carboxylesterase-6 [Drosophila miranda]
>gi|1036749861|ref|XP_017156289.1| PREDICTED: L-asparaginase [Drosophila miranda]
>gi|1036737180|ref|XP_017153765.1| PREDICTED: vesicle-associated membrane protein-associated p
>gi|1036737128|ref|XP_017153742.1| PREDICTED: vesicle-associated membrane protein-associated p
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>gi|1036737146|ref|XP_017153748.1| PREDICTED: vesicle-associated membrane protein-associated p
>gi|1036737164|ref|XP_017153757.1| PREDICTED: vesicle-associated membrane protein-associated p
>gi|1036725887|ref|XP_017147442.1| PREDICTED: uncharacterized protein LOC108159029 [Drosophila
>gi|1036725905|ref|XP_017147454.1| PREDICTED: CAS1 domain-containing protein 1 [Drosophila mira
>gi|1036699638|ref|XP_017143087.1| PREDICTED: uncharacterized protein LOC108156219 [Drosophila
>gi|1036757946|ref|XP_017156751.1| PREDICTED: uncharacterized protein ZK546.14 [Drosophila mira
>gi|1036757963|ref|XP_017156752.1| PREDICTED: uncharacterized protein ZK546.14 [Drosophila mira
>gi|1036745490|ref|XP_017156050.1| PREDICTED: uncharacterized protein LOC108164691 [Drosophila
>gi|1036745474|ref|XP_017156049.1| PREDICTED: V-type proton ATPase subunit d 1 [Drosophila mira
>gi|1036734365|ref|XP_017152153.1| PREDICTED: fizzy-related protein homolog [Drosophila miranda
>gi|1036734383|ref|XP_017152162.1| PREDICTED: fizzy-related protein homolog [Drosophila miranda
>gi|1036751528|ref|XP_017156383.1| PREDICTED: uncharacterized protein LOC108164911 isoform X1
>gi|1036751546|ref|XP_017156384.1| PREDICTED: acidic phospholipase A2 PA4 isoform X2 [Drosophil
>gi|1036751567|ref|XP_017156385.1| PREDICTED: uncharacterized protein LOC108164911 isoform X3
>gi|1036751588|ref|XP_017156386.1| PREDICTED: uncharacterized protein LOC108164911 isoform X3
>gi|1036734545|ref|XP_017152252.1| PREDICTED: nuclear cap-binding protein subunit 1 [Drosophila
>gi|1036734562|ref|XP_017152263.1| PREDICTED: mastermind-like domain-containing protein 1 [Dros
>gi|1036746577|ref|XP_017156116.1| PREDICTED: torsin-like protein [Drosophila miranda]
>gi|1036746594|ref|XP_017156118.1| PREDICTED: 39S ribosomal protein L30, mitochondrial [Drosoph
>gi|1036706790|ref|XP_017136564.1| PREDICTED: helix-loop-helix protein 1 [Drosophila miranda]
>gi|1036705672|ref|XP_017135748.1| PREDICTED: protein Flattop homolog [Drosophila miranda]
>gi|1036708216|ref|XP_017137496.1| PREDICTED: uncharacterized protein LOC108152568 [Drosophila
>gi|1036747833|ref|XP_017156192.1| PREDICTED: ras-interacting protein RIP3 [Drosophila miranda]
>gi|1036739472|ref|XP_017155007.1| PREDICTED: COPII coat assembly protein SEC16 isoform X1 [Dr
>gi|1036739490|ref|XP_017155016.1| PREDICTED: uncharacterized protein LOC108163962 isoform X2
>gi|1036721467|ref|XP_017145136.1| PREDICTED: uncharacterized abhydrolase domain-containing pro
>gi|1036721430|ref|XP_017145114.1| PREDICTED: serine/threonine-protein kinase minibrain isoform
>gi|1036721411|ref|XP_017145106.1| PREDICTED: serine/threonine-protein kinase minibrain isoform
>gi|1036721448|ref|XP_017145124.1| PREDICTED: zinc finger protein 622 [Drosophila miranda]
>gi|1036742270|ref|XP_017155875.1| PREDICTED: actin-related protein 8 [Drosophila miranda]
>gi|1036742285|ref|XP_017155876.1| PREDICTED: putative sulfiredoxin isoform X1 [Drosophila mira
>gi|1036742303|ref|XP_017155877.1| PREDICTED: putative sulfiredoxin isoform X2 [Drosophila mira
>gi|1036713586|ref|XP_017140747.1| PREDICTED: laccase-3 [Drosophila miranda]
>gi|1036713719|ref|XP_017140839.1| PREDICTED: uncharacterized protein LOC108154897 [Drosophila
>gi|1036713619|ref|XP_017140768.1| PREDICTED: uncharacterized protein LOC108154855 [Drosophila
>gi|1036713671|ref|XP_017140803.1| PREDICTED: RING finger protein 121 [Drosophila miranda]
>gi|1036713636|ref|XP_017140781.1| PREDICTED: mitochondrial uncoupling protein 4 [Drosophila mi
>gi|1036713655|ref|XP_017140791.1| PREDICTED: mitochondrial uncoupling protein 4 [Drosophila mi
>gi|1036713603|ref|XP_017140755.1| PREDICTED: replication factor C subunit 4 [Drosophila miranc
>gi|1036713704|ref|XP_017140827.1| PREDICTED: transcription initiation factor TFIID subunit 9
>gi|1036713688|ref|XP_017140815.1| PREDICTED: uncharacterized protein LOC108154880 [Drosophila
>gi|1036739732|ref|XP_017155160.1| PREDICTED: uncharacterized protein LOC108164082 [Drosophila
>gi|1036705782|ref|XP_017135831.1| PREDICTED: uncharacterized protein LOC108151634 [Drosophila
>gi|1036754620|ref|XP_017156568.1| PREDICTED: homeobox protein unc-4 isoform X1 [Drosophila mir
>gi|1036754637|ref|XP_017156569.1| PREDICTED: homeobox protein unc-4 isoform X2 [Drosophila mir
>gi|1036704299|ref|XP_017134806.1| PREDICTED: homeobox protein unc-4 [Drosophila miranda]
>gi|1036706421|ref|XP_017136290.1| PREDICTED: uncharacterized protein LOC108151894 [Drosophila
>gi|1036709987|ref|XP_017138564.1| PREDICTED: probable Ras GTPase-activating protein isoform X4
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>gi|1036709970|ref|XP_017138558.1| PREDICTED: probable Ras GTPase-activating protein isoform X3
>gi|1036709954|ref|XP_017138549.1| PREDICTED: probable Ras GTPase-activating protein isoform X2
>gi|1036709938|ref|XP_017138541.1| PREDICTED: probable Ras GTPase-activating protein isoform X1
>gi|1036710004|ref|XP_017138577.1| PREDICTED: DNA topoisomerase 1 isoform X1 [Drosophila miranc
>gi|1036710021|ref|XP_017138586.1| PREDICTED: DNA topoisomerase 1 isoform X2 [Drosophila miranc
>gi|1036710036|ref|XP_017138596.1| PREDICTED: DNA topoisomerase 1 isoform X3 [Drosophila miranc
>gi|1036710053|ref|XP_017138606.1| PREDICTED: DNA topoisomerase 1 isoform X4 [Drosophila miranc
>gi|1036739696|ref|XP_017155141.1| PREDICTED: dystrotelin [Drosophila miranda]
>gi|1036739714|ref|XP_017155149.1| PREDICTED: uncharacterized protein LOC108164074 [Drosophila
>gi|1036724829|ref|XP_017146830.1| PREDICTED: histone deacetylase 6 isoform X1 [Drosophila mira
>gi|1036724896|ref|XP_017146861.1| PREDICTED: histone deacetylase 6 isoform X4 [Drosophila mira
>gi|1036724914|ref|XP_017146869.1| PREDICTED: histone deacetylase 6 isoform X5 [Drosophila mira
>gi|1036724877|ref|XP_017146852.1| PREDICTED: histone deacetylase 6 isoform X3 [Drosophila mira
>gi|1036724845|ref|XP_017146839.1| PREDICTED: histone deacetylase 6 isoform X2 [Drosophila mira
>gi|1036724859|ref|XP_017146845.1| PREDICTED: histone deacetylase 6 isoform X2 [Drosophila mira
>gi|1036724932|ref|XP_017146882.1| PREDICTED: LOW QUALITY PROTEIN: p21-activated protein kinase
>gi|1036735729|ref|XP_017152924.1| PREDICTED: probable ATP-dependent RNA helicase DDX46 [Drosop
>gi|1036752409|ref|XP_017156434.1| PREDICTED: inhibitory POU protein isoform X4 [Drosophila mir
>gi|1036752390|ref|XP_017156433.1| PREDICTED: inhibitory POU protein isoform X3 [Drosophila mir
>gi|1036752372|ref|XP_017156432.1| PREDICTED: inhibitory POU protein isoform X2 [Drosophila mir
>gi|1036752356|ref|XP_017156431.1| PREDICTED: inhibitory POU protein isoform X1 [Drosophila mir
>gi|1036752427|ref|XP_017156435.1| PREDICTED: serine/threonine-protein phosphatase alpha-3 isor
>gi|1036752444|ref|XP_017156437.1| PREDICTED: uncharacterized protein LOC108164952 [Drosophila
>gi|1036752463|ref|XP_017156438.1| PREDICTED: uncharacterized protein LOC108164952 [Drosophila
>gi|1036757464|ref|XP_017156727.1| PREDICTED: WSCD family member GA21586 isoform X2 [Drosophila
>gi|1036757447|ref|XP_017156726.1| PREDICTED: WSCD family member GA21586 isoform X1 [Drosophila
>gi|1036748644|ref|XP_017156223.1| PREDICTED: uncharacterized protein LOC108164814 isoform X2
>gi|1036748625|ref|XP_017156221.1| PREDICTED: uncharacterized protein LOC108164814 isoform X1
>gi|1036748663|ref|XP_017156224.1| PREDICTED: uncharacterized protein LOC108164814 isoform X3
>gi|1036699658|ref|XP_017143210.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036748682|ref|XP_017156225.1| PREDICTED: uncharacterized protein LOC108164815 [Drosophila
>gi|1036736261|ref|XP_017153228.1| PREDICTED: secretory carrier-associated membrane protein 1
>gi|1036736246|ref|XP_017153218.1| PREDICTED: adenosylhomocysteinase [Drosophila miranda]
>gi|1036716747|ref|XP_017142604.1| PREDICTED: P3 protein [Drosophila miranda]
>gi|1036716708|ref|XP_017142580.1| PREDICTED: anaphase-promoting complex subunit 1 [Drosophila
>gi|1036716785|ref|XP_017142630.1| PREDICTED: uncharacterized protein LOC108155963 [Drosophila
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>gi|1036716824|ref|XP_017142647.1| PREDICTED: uncharacterized protein LOC108155963 [Drosophila
>gi|1036716847|ref|XP_017142659.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036716726|ref|XP_017142593.1| PREDICTED: serine/threonine-protein kinase Nek11 [Drosophila
>gi|1036716766|ref|XP_017142616.1| PREDICTED: acetyl-CoA acetyltransferase, mitochondrial [Dros
>gi|1036757191|ref|XP_017156710.1| PREDICTED: uncharacterized protein LOC108165155 [Drosophila
>gi|1036704800|ref|XP_017135157.1| PREDICTED: selenide, water dikinase-like [Drosophila miranda
>gi|1036707217|ref|XP_017136828.1| PREDICTED: uncharacterized protein LOC108152189 [Drosophila
>gi|1036699672|ref|XP_017143336.1| PREDICTED: uncharacterized protein LOC108156386 [Drosophila
>gi|1036699685|ref|XP_017143453.1| PREDICTED: uncharacterized protein LOC108156460 [Drosophila
>gi|1036751014|ref|XP_017156353.1| PREDICTED: uncharacterized protein LOC108164897 [Drosophila
>gi|1036751032|ref|XP_017156354.1| PREDICTED: defective chorion-1 protein, FC106 isoform [Droso
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>gi|1036757627|ref|XP_017156734.1| PREDICTED: golgin subfamily A member 4-like [Drosophila mira
>gi|1036699701|ref|XP_017143575.1| PREDICTED: uncharacterized protein LOC108156538 [Drosophila
>gi|1036699714|ref|XP_017143700.1| PREDICTED: mitochondrial carrier homolog 2-like [Drosophila
>gi|1036699731|ref|XP_017143948.1| PREDICTED: mitochondrial carrier homolog 2-like [Drosophila
>gi|1036699746|ref|XP_017144072.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036757755|ref|XP_017156740.1| PREDICTED: carbonic anhydrase-related protein 10 [Drosophila
>gi|1036755734|ref|XP_017156630.1| PREDICTED: AT-rich interactive domain-containing protein 1B
>gi|1036753867|ref|XP_017156524.1| PREDICTED: uncharacterized protein LOC108165002 [Drosophila
>gi|1036743371|ref|XP_017155940.1| PREDICTED: cap-specific mRNA (nucleoside-2'-O-)-methyltrans
>gi|1036747312|ref|XP_017156162.1| PREDICTED: uncharacterized protein LOC108164762 [Drosophila
>gi|1036747331|ref|XP_017156163.1| PREDICTED: uncharacterized protein LOC108164763 [Drosophila
>gi|1036757135|ref|XP_017156707.1| PREDICTED: LOW QUALITY PROTEIN: polycystic kidney disease 2-
>gi|1036756376|ref|XP_017156667.1| PREDICTED: polycystic kidney disease 2-like 1 protein isofor
>gi|1036756359|ref|XP_017156666.1| PREDICTED: polycystic kidney disease 2-like 2 protein isofor
>gi|1036756395|ref|XP_017156668.1| PREDICTED: uncharacterized protein LOC108165118 isoform X3
>gi|1036755228|ref|XP_017156604.1| PREDICTED: collagen alpha-1(II) chain-like isoform X1 [Dros
>gi|1036755244|ref|XP_017156605.1| PREDICTED: collagen alpha-1(III) chain-like isoform X2 [Dro
>gi|1036737545|ref|XP_017153972.1| PREDICTED: uncharacterized protein LOC108163279 [Drosophila
>gi|1036723410|ref|XP_017145959.1| PREDICTED: serine/threonine-protein kinase TAO3 isoform X1
>gi|1036723429|ref|XP_017145967.1| PREDICTED: serine/threonine-protein kinase TAO3 isoform X2
>gi|1036723446|ref|XP_017145979.1| PREDICTED: vacuolar protein sorting-associated protein 33A
>gi|1036735745|ref|XP_017152937.1| PREDICTED: gamma-tubulin complex component 2 homolog isoform
>gi|1036735764|ref|XP_017152944.1| PREDICTED: gamma-tubulin complex component 2 homolog isoform
>gi|1036699760|ref|XP_017144189.1| PREDICTED: homeobox protein ARX [Drosophila miranda]
>gi|1036730153|ref|XP_017150023.1| PREDICTED: REST corepressor isoform X2 [Drosophila miranda]
>gi|1036730134|ref|XP_017150015.1| PREDICTED: REST corepressor isoform X1 [Drosophila miranda]
>gi|1036730172|ref|XP_017150031.1| PREDICTED: REST corepressor isoform X3 [Drosophila miranda]
>gi|1036745649|ref|XP_017156060.1| PREDICTED: probable isocitrate dehydrogenase [NAD] subunit a
>gi|1036745667|ref|XP_017156061.1| PREDICTED: probable isocitrate dehydrogenase [NAD] subunit a
>gi|1036718720|ref|XP_017143701.1| PREDICTED: uncharacterized protein LOC108156629 [Drosophila
>gi|1036718740|ref|XP_017143712.1| PREDICTED: uncharacterized protein LOC108156629 [Drosophila
>gi|1036718769|ref|XP_017143733.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036718754|ref|XP_017143724.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036718788|ref|XP_017143741.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036718807|ref|XP_017143751.1| PREDICTED: 6-phosphofructo-2-kinase/fructose-2,6-bisphosphat
>gi|1036702925|ref|XP_017156678.1| PREDICTED: protein kish [Drosophila miranda]
>gi|1036702323|ref|XP_017156292.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prote
>gi|1036721393|ref|XP_017145094.1| PREDICTED: LOW QUALITY PROTEIN: 5'-3' exoribonuclease 1 [Dro
>gi|1036721374|ref|XP_017145083.1| PREDICTED: probable ATP-dependent RNA helicase DHX34 [Drosop
>gi|1036734653|ref|XP_017152326.1| PREDICTED: 28S ribosomal protein S14, mitochondrial [Drosoph
>gi|1036734615|ref|XP_017152306.1| PREDICTED: N-alpha-acetyltransferase 16, NatA auxiliary subu
>gi|1036734634|ref|XP_017152314.1| PREDICTED: N-alpha-acetyltransferase 16, NatA auxiliary subu
>gi|1036699775|ref|XP_017144311.1| PREDICTED: uncharacterized protein LOC108156988 [Drosophila
>gi|1036722703|ref|XP_017145617.1| PREDICTED: RING-box protein 1-like [Drosophila miranda]
>gi|1036757889|ref|XP_017156748.1| PREDICTED: uncharacterized protein LOC108165188 [Drosophila
>gi|1036704819|ref|XP_017135169.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036757392|ref|XP_017156723.1| PREDICTED: nose resistant to fluoxetine protein 6-like isofo
>gi|1036757410|ref|XP_017156724.1| PREDICTED: nose resistant to fluoxetine protein 6-like isofo
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>gi|1036757429|ref|XP_017156725.1| PREDICTED: nose resistant to fluoxetine protein 6-like isof
>gi|1036757588|ref|XP_017156732.1| PREDICTED: nardilysin-like isoform X1 [Drosophila miranda]
>gi|1036757607|ref|XP_017156733.1| PREDICTED: nardilysin-like isoform X2 [Drosophila miranda]
>gi|1036756780|ref|XP_017156690.1| PREDICTED: nose resistant to fluoxetine protein 6-like [Dros
>gi|1036708492|ref|XP_017137675.1| PREDICTED: 40S ribosomal protein S10b [Drosophila miranda]
>gi|1036708380|ref|XP_017137603.1| PREDICTED: sin3 histone deacetylase corepressor complex comp
>gi|1036708431|ref|XP_017137642.1| PREDICTED: heat shock protein beta-6 isoform X1 [Drosophila
>gi|1036708447|ref|XP_017137648.1| PREDICTED: heat shock protein beta-1 isoform X2 [Drosophila
>gi|1036708318|ref|XP_017137562.1| PREDICTED: solute carrier family 25 member 40 isoform X1 [D:
>gi|1036708331|ref|XP_017137571.1| PREDICTED: solute carrier family 25 member 40 isoform X2 [D:
>gi|1036708284|ref|XP_017137542.1| PREDICTED: solute carrier family 25 member 40 isoform X1 [D:
>gi|1036708301|ref|XP_017137550.1| PREDICTED: solute carrier family 25 member 40 isoform X2 [D:
>gi|1036708509|ref|XP_017137687.1| PREDICTED: coiled-coil domain-containing protein 86 [Drosoph
>gi|1036708397|ref|XP_017137618.1| PREDICTED: L-seryl-tRNA(Sec) kinase [Drosophila miranda]
>gi|1036708525|ref|XP_017137701.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036699792|ref|XP_017144438.1| PREDICTED: uncharacterized protein LOC108157068 [Drosophila
>gi|1036708478|ref|XP_017137663.1| PREDICTED: N-alpha-acetyltransferase 20 [Drosophila miranda]
>gi|1036708346|ref|XP_017137583.1| PREDICTED: dual specificity protein phosphatase 12 [Drosophi
>gi|1036708364|ref|XP_017137591.1| PREDICTED: dual specificity protein phosphatase 12 [Drosophi
>gi|1036708414|ref|XP_017137630.1| PREDICTED: pyridoxal phosphate phosphatase PHOSPHO2 [Drosoph
>gi|1036708250|ref|XP_017137521.1| PREDICTED: supporter of activation of yellow protein [Drosop
>gi|1036708266|ref|XP_017137529.1| PREDICTED: supporter of activation of yellow protein [Drosop
>gi|1036714845|ref|XP_017141500.1| PREDICTED: pyridoxal phosphate phosphatase PHOSPHO2 [Drosoph
>gi|1036714809|ref|XP_017141480.1| PREDICTED: cell differentiation protein RCD1 homolog [Drosop
>gi|1036714828|ref|XP_017141489.1| PREDICTED: cell differentiation protein RCD1 homolog [Drosop
>gi|1036714863|ref|XP_017141510.1| PREDICTED: protein transport protein Sec61 gamma-2 subunit
>gi|1036714772|ref|XP_017141455.1| PREDICTED: signal recognition particle receptor subunit alph
>gi|1036714790|ref|XP_017141468.1| PREDICTED: 26S protease regulatory subunit 6B [Drosophila m:
>gi|1036714735|ref|XP_017141434.1| PREDICTED: uncharacterized protein LOC108155248 isoform X1
>gi|1036714753|ref|XP_017141442.1| PREDICTED: uncharacterized protein LOC108155248 isoform X2
>gi|1036742320|ref|XP_017155878.1| PREDICTED: 60 kDa heat shock protein, mitochondrial [Drosoph
>gi|1036753814|ref|XP_017156520.1| PREDICTED: uncharacterized protein LOC108165001 [Drosophila
>gi|1036753831|ref|XP_017156521.1| PREDICTED: uncharacterized protein LOC108165001 [Drosophila
>gi|1036753850|ref|XP_017156523.1| PREDICTED: uncharacterized protein LOC108165001 [Drosophila
>gi|1036726131|ref|XP_017147590.1| PREDICTED: protein ELYS homolog [Drosophila miranda]
>gi|1036739261|ref|XP_017154884.1| PREDICTED: exportin-5 [Drosophila miranda]
>gi|1036732555|ref|XP_017151198.1| PREDICTED: actin-related protein 10 [Drosophila miranda]
>gi|1036732537|ref|XP_017151189.1| PREDICTED: CWF19-like protein 2 homolog [Drosophila miranda]
>gi|1036732572|ref|XP_017151208.1| PREDICTED: 40S ribosomal protein S19a [Drosophila miranda]
>gi|1036726596|ref|XP_017147892.1| PREDICTED: rho-associated protein kinase 1 isoform X1 [Dros
>gi|1036726614|ref|XP_017147902.1| PREDICTED: rho-associated protein kinase 1 isoform X2 [Dros
>gi|1036726635|ref|XP_017147912.1| PREDICTED: rho-associated protein kinase 1 isoform X3 [Dros
>gi|1036730558|ref|XP_017150260.1| PREDICTED: brain acid soluble protein 1 homolog [Drosophila
>gi|1036730522|ref|XP_017150243.1| PREDICTED: nardilysin-like isoform X1 [Drosophila miranda]
>gi|1036730542|ref|XP_017150248.1| PREDICTED: nardilysin-like isoform X2 [Drosophila miranda]
>gi|1036751918|ref|XP_017156404.1| PREDICTED: 40S ribosomal protein S19a-like [Drosophila mirar
>gi|1036741362|ref|XP_017155825.1| PREDICTED: myotubularin-related protein 10-B [Drosophila mi:
>gi|1036699809|ref|XP_017144563.1| PREDICTED: uncharacterized protein LOC108157134 [Drosophila
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>gi|1036752338|ref|XP_017156430.1| PREDICTED: glutaredoxin-related protein 5, mitochondrial [Di
>gi|1036716487|ref|XP_017142475.1| PREDICTED: flotillin-2 isoform X1 [Drosophila miranda]
>gi|1036716503|ref|XP_017142484.1| PREDICTED: flotillin-2 isoform X1 [Drosophila miranda]
>gi|1036716557|ref|XP_017142492.1| PREDICTED: flotillin-2 isoform X1 [Drosophila miranda]
>gi|1036716576|ref|XP_017142501.1| PREDICTED: flotillin-2 isoform X2 [Drosophila miranda]
>gi|1036716595|ref|XP_017142511.1| PREDICTED: flotillin-2 isoform X3 [Drosophila miranda]
>gi|1036716453|ref|XP_017142450.1| PREDICTED: glucose dehydrogenase [FAD, quinone] isoform X2
>gi|1036716381|ref|XP_017142404.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716434|ref|XP_017142444.1| PREDICTED: glucose dehydrogenase [FAD, quinone] isoform X1
>gi|1036716400|ref|XP_017142420.1| PREDICTED: LOW QUALITY PROTEIN: glucose dehydrogenase [FAD,
>gi|1036716362|ref|XP_017142392.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716651|ref|XP_017142542.1| PREDICTED: uncharacterized protein LOC108155905 [Drosophila
>gi|1036716288|ref|XP_017142339.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716690|ref|XP_017142567.1| PREDICTED: glucose dehydrogenase [FAD, quinone]-like [Drosop
>gi|1036716268|ref|XP_017142328.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716470|ref|XP_017142463.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716325|ref|XP_017142368.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716670|ref|XP_017142555.1| PREDICTED: uncharacterized protein LOC108155915 [Drosophila
>gi|1036716343|ref|XP_017142379.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716306|ref|XP_017142354.1| PREDICTED: LOW QUALITY PROTEIN: glucose dehydrogenase [FAD,
>gi|1036716416|ref|XP_017142433.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1036716630|ref|XP_017142530.1| PREDICTED: uncharacterized protein LOC108155896 [Drosophila
>gi|1036716611|ref|XP_017142519.1| PREDICTED: flotillin-2 isoform X4 [Drosophila miranda]
>gi|1036737512|ref|XP_017153952.1| PREDICTED: probable 4-coumarate--CoA ligase 1 [Drosophila mi
>gi|1036750922|ref|XP_017156348.1| PREDICTED: chromobox protein homolog 5-like [Drosophila mira
>gi|1036750848|ref|XP_017156344.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036750904|ref|XP_017156347.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036750866|ref|XP_017156345.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036750885|ref|XP_017156346.1| PREDICTED: potassium voltage-gated channel protein eag isofo
>gi|1036755473|ref|XP_017156618.1| PREDICTED: cathepsin E-B-like [Drosophila miranda]
>gi|1036743389|ref|XP_017155941.1| PREDICTED: uncharacterized protein LOC108164612 [Drosophila
>gi|1036743408|ref|XP_017155942.1| PREDICTED: branched-chain-amino-acid aminotransferase, cytos
>gi|1036723989|ref|XP_017146275.1| PREDICTED: mpv17-like protein 2 [Drosophila miranda]
>gi|1036724008|ref|XP_017146288.1| PREDICTED: protein yippee [Drosophila miranda]
>gi|1036724029|ref|XP_017146296.1| PREDICTED: protein yippee [Drosophila miranda]
>gi|1036724048|ref|XP_017146309.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036723970|ref|XP_017146263.1| PREDICTED: RNA-binding protein 1 isoform X2 [Drosophila mira
>gi|1036723953|ref|XP_017146252.1| PREDICTED: RNA-binding protein 1 isoform X1 [Drosophila mira
>gi|1036743339|ref|XP_017155938.1| PREDICTED: alanine aminotransferase 1 [Drosophila miranda]
>gi|1036743355|ref|XP_017155939.1| PREDICTED: alanine aminotransferase 1 [Drosophila miranda]
>gi|1036706093|ref|XP_017136076.1| PREDICTED: doublesex- and mab-3-related transcription facto
>gi|1036706108|ref|XP_017136083.1| PREDICTED: uncharacterized protein LOC108151748 isoform X2
>gi|1036757699|ref|XP_017156738.1| PREDICTED: uncharacterized protein LOC108165178 [Drosophila
>gi|1036738883|ref|XP_017154657.1| PREDICTED: nuclear speckle splicing regulatory protein 1 [Di
>gi|1036738901|ref|XP_017154668.1| PREDICTED: 40S ribosomal protein S15Aa [Drosophila miranda]
>gi|1036715760|ref|XP_017142027.1| PREDICTED: peroxiredoxin 1 [Drosophila miranda]
>gi|1036715712|ref|XP_017141989.1| PREDICTED: uncharacterized protein LOC108155578 isoform X2
>gi|1036715674|ref|XP_017141973.1| PREDICTED: uncharacterized protein LOC108155578 isoform X1
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>gi|1036715693|ref|XP_017141982.1| PREDICTED: uncharacterized protein LOC108155578 isoform X1
>gi|1036715730|ref|XP_017142002.1| PREDICTED: uncharacterized protein LOC108155592 [Drosophila
>gi|1036715747|ref|XP_017142015.1| PREDICTED: pre-mRNA-splicing factor 38B [Drosophila miranda]
>gi|1036715640|ref|XP_017141951.1| PREDICTED: raf homolog serine/threonine-protein kinase phl
>gi|1036715657|ref|XP_017141961.1| PREDICTED: raf homolog serine/threonine-protein kinase phl
>gi|1036715779|ref|XP_017142039.1| PREDICTED: 39S ribosomal protein L14, mitochondrial [Drosoph
>gi|1036737020|ref|XP_017153684.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036737038|ref|XP_017153692.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036737057|ref|XP_017153701.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036737076|ref|XP_017153710.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036737094|ref|XP_017153719.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036757793|ref|XP_017156742.1| PREDICTED: sporozoite surface protein 2-like [Drosophila mi
>gi|1036757814|ref|XP_017156743.1| PREDICTED: sporozoite surface protein 2-like [Drosophila mi
>gi|1036704513|ref|XP_017134942.1| PREDICTED: putative uncharacterized protein DDB_G0268364 [Dr
>gi|1036744504|ref|XP_017155993.1| PREDICTED: uncharacterized protein LOC108164652 isoform X1
>gi|1036744523|ref|XP_017155994.1| PREDICTED: uncharacterized protein LOC108164652 isoform X2
>gi|1036704226|ref|XP_017134763.1| PREDICTED: uncharacterized protein LOC108150940 [Drosophila
>gi|1036726833|ref|XP_017148037.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036743927|ref|XP_017155965.1| PREDICTED: inosine-5'-monophosphate dehydrogenase [Drosophi
>gi|1036743946|ref|XP_017155966.1| PREDICTED: inosine-5'-monophosphate dehydrogenase [Drosophi
>gi|1036755210|ref|XP_017156603.1| PREDICTED: rabphilin-3A [Drosophila miranda]
>gi|1036744063|ref|XP_017155972.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036744082|ref|XP_017155973.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036744101|ref|XP_017155974.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036748183|ref|XP_017156209.1| PREDICTED: gamma-aminobutyric acid receptor-associated prote
>gi|1036752829|ref|XP_017156460.1| PREDICTED: transcription initiation factor IIA subunit 2 [Dr
>gi|1036719530|ref|XP_017144168.1| PREDICTED: integrin beta-PS isoform X1 [Drosophila miranda]
>gi|1036719547|ref|XP_017144176.1| PREDICTED: integrin beta-PS isoform X1 [Drosophila miranda]
>gi|1036719567|ref|XP_017144184.1| PREDICTED: integrin beta-PS isoform X2 [Drosophila miranda]
>gi|1036719511|ref|XP_017144156.1| PREDICTED: regulator of nonsense transcripts 2 [Drosophila m
>gi|1036749467|ref|XP_017156266.1| PREDICTED: dynein intermediate chain 3, ciliary [Drosophila
>gi|1036749481|ref|XP_017156268.1| PREDICTED: 17-beta-hydroxysteroid dehydrogenase 13 [Drosoph
>gi|1036699824|ref|XP_017144691.1| PREDICTED: EF-hand calcium-binding domain-containing protei
>gi|1036725087|ref|XP_017146972.1| PREDICTED: ubiquitin-conjugating enzyme E2 H [Drosophila mi
>gi|1036725103|ref|XP_017146979.1| PREDICTED: ubiquitin-conjugating enzyme E2 H [Drosophila mi
>gi|1036725070|ref|XP_017146959.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036725116|ref|XP_017146990.1| PREDICTED: uncharacterized protein LOC108158809 [Drosophila
>gi|1036725134|ref|XP_017146999.1| PREDICTED: uncharacterized protein LOC108158809 [Drosophila
>gi|1036725152|ref|XP_017147006.1| PREDICTED: uncharacterized protein LOC108158809 [Drosophila
>gi|1036749533|ref|XP_017156271.1| PREDICTED: uncharacterized protein LOC108164853 [Drosophila
>gi|1036699841|ref|XP_017144812.1| PREDICTED: inosine-5'-monophosphate dehydrogenase-like [Dros
>gi|1036730503|ref|XP_017150229.1| PREDICTED: putative folylpolyglutamate synthase [Drosophila
>gi|1036730484|ref|XP_017150217.1| PREDICTED: uncharacterized protein LOC108160603 isoform X2
>gi|1036730450|ref|XP_017150200.1| PREDICTED: uncharacterized protein LOC108160603 isoform X1
>gi|1036730467|ref|XP_017150208.1| PREDICTED: uncharacterized protein LOC108160603 isoform X1
>gi|1036756545|ref|XP_017156676.1| PREDICTED: YEATS domain-containing protein 4 [Drosophila mi
>gi|1036756524|ref|XP_017156675.1| PREDICTED: uncharacterized protein LOC108165123 [Drosophila
>gi|1036717066|ref|XP_017142799.1| PREDICTED: protein kinase shaggy isoform X2 [Drosophila mira
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>gi|1036717086|ref|XP_017142807.1| PREDICTED: protein kinase shaggy isoform X3 [Drosophila mira
>gi|1036717047|ref|XP_017142789.1| PREDICTED: protein kinase shaggy isoform X1 [Drosophila mira
>gi|1036732065|ref|XP_017150966.1| PREDICTED: phosphatidylinositol 4-kinase alpha [Drosophila m
>gi|1036732615|ref|XP_017151230.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036732633|ref|XP_017151238.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036732651|ref|XP_017151246.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036732669|ref|XP_017151253.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036737668|ref|XP_017154048.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036737686|ref|XP_017154058.1| PREDICTED: enhancer of yellow 2 transcription factor isoform
>gi|1036737703|ref|XP_017154062.1| PREDICTED: enhancer of yellow 2 transcription factor isoform
>gi|1036737652|ref|XP_017154034.1| PREDICTED: transcription factor grauzone [Drosophila miranda
>gi|1036744992|ref|XP_017156020.1| PREDICTED: uncharacterized protein LOC108164676 [Drosophila
>gi|1036745011|ref|XP_017156021.1| PREDICTED: transmembrane protein 242 [Drosophila miranda]
>gi|1036744937|ref|XP_017156017.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036744954|ref|XP_017156018.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036744973|ref|XP_017156019.1| PREDICTED: zinc finger protein 239-like [Drosophila miranda]
>gi|1036723507|ref|XP_017146006.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB1 [Dros
>gi|1036722753|ref|XP_017145636.1| PREDICTED: putative ATP-dependent RNA helicase BoYb [Drosoph
>gi|1036722997|ref|XP_017145742.1| PREDICTED: LOW QUALITY PROTEIN: peptidoglycan-recognition pr
>gi|1036722770|ref|XP_017145646.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722791|ref|XP_017145653.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722812|ref|XP_017145661.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722855|ref|XP_017145669.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722874|ref|XP_017145676.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722889|ref|XP_017145683.1| PREDICTED: homeobox protein 5 isoform X1 [Drosophila miranda
>gi|1036722925|ref|XP_017145698.1| PREDICTED: homeobox protein 5 isoform X3 [Drosophila miranda
>gi|1036722906|ref|XP_017145691.1| PREDICTED: homeobox protein 5 isoform X2 [Drosophila miranda
>gi|1036722943|ref|XP_017145706.1| PREDICTED: homeobox protein 5 isoform X4 [Drosophila miranda
>gi|1036722980|ref|XP_017145727.1| PREDICTED: coiled-coil domain-containing protein 25 [Drosoph
>gi|1036722962|ref|XP_017145716.1| PREDICTED: facilitated trehalose transporter Tret1 [Drosoph
>gi|1036720091|ref|XP_017144450.1| PREDICTED: putative uncharacterized protein DDB_G0286901 [D
>gi|1036720053|ref|XP_017144428.1| PREDICTED: protein sex-lethal isoform X10 [Drosophila miranc
>gi|1036719889|ref|XP_017144358.1| PREDICTED: protein sex-lethal isoform X2 [Drosophila miranda
>gi|1036719972|ref|XP_017144392.1| PREDICTED: protein sex-lethal isoform X6 [Drosophila miranda
>gi|1036719869|ref|XP_017144350.1| PREDICTED: protein sex-lethal isoform X1 [Drosophila miranda
>gi|1036720073|ref|XP_017144439.1| PREDICTED: protein sex-lethal isoform X11 [Drosophila miranc
>gi|1036719908|ref|XP_017144368.1| PREDICTED: protein sex-lethal isoform X3 [Drosophila miranda
>gi|1036720033|ref|XP_017144419.1| PREDICTED: protein sex-lethal isoform X9 [Drosophila miranda
>gi|1036720014|ref|XP_017144408.1| PREDICTED: protein sex-lethal isoform X8 [Drosophila miranda
>gi|1036719928|ref|XP_017144375.1| PREDICTED: protein sex-lethal isoform X4 [Drosophila miranda
>gi|1036719952|ref|XP_017144383.1| PREDICTED: protein sex-lethal isoform X5 [Drosophila miranda
>gi|1036719994|ref|XP_017144400.1| PREDICTED: protein sex-lethal isoform X7 [Drosophila miranda
>gi|1036710707|ref|XP_017139059.1| PREDICTED: protein suppressor of sable isoform X2 [Drosophil
>gi|1036710692|ref|XP_017139050.1| PREDICTED: protein suppressor of sable isoform X1 [Drosophil
>gi|1036710744|ref|XP_017139082.1| PREDICTED: caspase-8 [Drosophila miranda]
>gi|1036710726|ref|XP_017139070.1| PREDICTED: uncharacterized protein LOC108153519 [Drosophila
>gi|1036710776|ref|XP_017139105.1| PREDICTED: tetraspanin-33 [Drosophila miranda]
>gi|1036710793|ref|XP_017139114.1| PREDICTED: tetraspanin-33 [Drosophila miranda]
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>gi|1036710760|ref|XP_017139093.1| PREDICTED: seipin [Drosophila miranda]
>gi|1036748948|ref|XP_017156238.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X2 [Dros
>gi|1036748907|ref|XP_017156236.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X1 [Dros
>gi|1036748929|ref|XP_017156237.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X1 [Dros
>gi|1036745978|ref|XP_017156078.1| PREDICTED: 60S ribosomal protein L17 [Drosophila miranda]
>gi|1036744920|ref|XP_017156016.1| PREDICTED: putative metabolite transport protein HI_1104 [Di
>gi|1036711151|ref|XP_017139364.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711167|ref|XP_017139372.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711184|ref|XP_017139381.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711199|ref|XP_017139389.1| PREDICTED: uncharacterized protein LOC108153742 [Drosophila
>gi|1036711324|ref|XP_017139426.1| PREDICTED: uncharacterized protein LOC108153835 [Drosophila
>gi|1036711290|ref|XP_017139421.1| PREDICTED: uncharacterized protein LOC108153801 [Drosophila
>gi|1036711307|ref|XP_017139422.1| PREDICTED: 5-demethoxyubiquinone hydroxylase, mitochondrial
>gi|1036711273|ref|XP_017139420.1| PREDICTED: protein shifted isoform X4 [Drosophila miranda]
>gi|1036711256|ref|XP_017139415.1| PREDICTED: protein shifted isoform X3 [Drosophila miranda]
>gi|1036711235|ref|XP_017139408.1| PREDICTED: protein shifted isoform X2 [Drosophila miranda]
>gi|1036711216|ref|XP_017139399.1| PREDICTED: protein shifted isoform X1 [Drosophila miranda]
>gi|1036726982|ref|XP_017148114.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X4
>gi|1036727021|ref|XP_017148131.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X6
>gi|1036726963|ref|XP_017148107.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X3
>gi|1036726945|ref|XP_017148099.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X2
>gi|1036726927|ref|XP_017148092.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X1
>gi|1036726999|ref|XP_017148122.1| PREDICTED: guanine nucleotide-releasing factor 2 isoform X5
>gi|1036725173|ref|XP_017147019.1| PREDICTED: coronin-7 isoform X1 [Drosophila miranda]
>gi|1036725191|ref|XP_017147027.1| PREDICTED: coronin-7 isoform X2 [Drosophila miranda]
>gi|1036725210|ref|XP_017147038.1| PREDICTED: coronin-7 isoform X3 [Drosophila miranda]
>gi|1036725226|ref|XP_017147047.1| PREDICTED: coronin-7 isoform X4 [Drosophila miranda]
>gi|1036737198|ref|XP_017153777.1| PREDICTED: uncharacterized protein LOC108163163 [Drosophila
>gi|1036737233|ref|XP_017153800.1| PREDICTED: putative uncharacterized protein DDB_G0268364 iso
>gi|1036737213|ref|XP_017153791.1| PREDICTED: putative uncharacterized protein DDB_G0268364 iso
>gi|1036744540|ref|XP_017155995.1| PREDICTED: probable dolichyl pyrophosphate Glc1Man9GlcNAc2 a
>gi|1036744556|ref|XP_017155996.1| PREDICTED: ataxin-1 [Drosophila miranda]
>gi|1036742682|ref|XP_017155900.1| PREDICTED: frizzled-3 [Drosophila miranda]
>gi|1036749239|ref|XP_017156255.1| PREDICTED: protein O-mannosyl-transferase 2 [Drosophila mira
>gi|1036749258|ref|XP_017156256.1| PREDICTED: uncharacterized protein LOC108164841 [Drosophila
>gi|1036749297|ref|XP_017156259.1| PREDICTED: uncharacterized protein LOC108164842 isoform X2
>gi|1036749278|ref|XP_017156258.1| PREDICTED: uncharacterized protein LOC108164842 isoform X1
>gi|1036749327|ref|XP_017156260.1| PREDICTED: uncharacterized protein LOC108164842 isoform X3
>gi|1036699856|ref|XP_017144927.1| PREDICTED: uncharacterized protein LOC108157372 [Drosophila
>gi|1036749342|ref|XP_017156261.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036749360|ref|XP_017156262.1| PREDICTED: uncharacterized protein LOC108164844 [Drosophila
>gi|1036717849|ref|XP_017143223.1| PREDICTED: histone deacetylase complex subunit SAP30 homolog
>gi|1036717731|ref|XP_017143155.1| PREDICTED: exosome complex component RRP45 [Drosophila mirar
>gi|1036717772|ref|XP_017143180.1| PREDICTED: uncharacterized protein LOC108156303 isoform X1
>gi|1036717791|ref|XP_017143187.1| PREDICTED: UPF0472 protein C16orf72 isoform X2 [Drosophila r
>gi|1036717809|ref|XP_017143198.1| PREDICTED: NECAP-like protein CG9132 [Drosophila miranda]
>gi|1036717753|ref|XP_017143167.1| PREDICTED: zinc finger protein 845 [Drosophila miranda]
>gi|1036717830|ref|XP_017143211.1| PREDICTED: 39S ribosomal protein L22, mitochondrial [Drosopl
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>gi|1036744327|ref|XP_017155985.1| PREDICTED: uncharacterized protein LOC108164646 [Drosophila
>gi|1036730907|ref|XP_017150464.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730926|ref|XP_017150472.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730940|ref|XP_017150481.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730959|ref|XP_017150488.1| PREDICTED: integrin alpha-PS2 isoform X1 [Drosophila miranda
>gi|1036730979|ref|XP_017150498.1| PREDICTED: integrin alpha-PS2 isoform X2 [Drosophila miranda
>gi|1036705657|ref|XP_017135737.1| PREDICTED: transmembrane protein 256 homolog [Drosophila mir
>gi|1036728452|ref|XP_017148976.1| PREDICTED: endothelin-converting enzyme 1 isoform X1 [Drosop
>gi|1036728471|ref|XP_017148984.1| PREDICTED: endothelin-converting enzyme 1 isoform X2 [Drosop
>gi|1036728532|ref|XP_017149017.1| PREDICTED: keratin, type I cytoskeletal 10 [Drosophila miran
>gi|1036728585|ref|XP_017149055.1| PREDICTED: uncharacterized protein LOC108159922 [Drosophila
>gi|1036728568|ref|XP_017149042.1| PREDICTED: uncharacterized protein LOC108159916 [Drosophila
>gi|1036728550|ref|XP_017149031.1| PREDICTED: uncharacterized protein LOC108159907 [Drosophila
>gi|1036728509|ref|XP_017149006.1| PREDICTED: uncharacterized protein LOC108159887 isoform X2
>gi|1036728490|ref|XP_017148997.1| PREDICTED: uncharacterized protein LOC108159887 isoform X1
>gi|1036728604|ref|XP_017149068.1| PREDICTED: transforming growth factor beta-1-induced transcr
>gi|1036728621|ref|XP_017149076.1| PREDICTED: transforming growth factor beta-1-induced transcr
>gi|1036747573|ref|XP_017156177.1| PREDICTED: uncharacterized protein LOC108164772 [Drosophila
>gi|1036747631|ref|XP_017156180.1| PREDICTED: carboxypeptidase D-like isoform X3 [Drosophila mi
>gi|1036747612|ref|XP_017156179.1| PREDICTED: carboxypeptidase D-like isoform X2 [Drosophila mi
>gi|1036747591|ref|XP_017156178.1| PREDICTED: carboxypeptidase D-like isoform X1 [Drosophila mi
>gi|1036747666|ref|XP_017156182.1| PREDICTED: carboxypeptidase D-like isoform X5 [Drosophila mi
>gi|1036747649|ref|XP_017156181.1| PREDICTED: carboxypeptidase D-like isoform X4 [Drosophila mi
>gi|1036747722|ref|XP_017156186.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036747760|ref|XP_017156188.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036747685|ref|XP_017156183.1| PREDICTED: trypsin beta-like [Drosophila miranda]
>gi|1036747741|ref|XP_017156187.1| PREDICTED: chymotrypsin-1-like [Drosophila miranda]
>gi|1036747704|ref|XP_017156184.1| PREDICTED: serine protease SP24D-like [Drosophila miranda]
>gi|1036717598|ref|XP_017143079.1| PREDICTED: transmembrane GTPase Marf isoform X3 [Drosophila
>gi|1036717579|ref|XP_017143072.1| PREDICTED: transmembrane GTPase Marf isoform X2 [Drosophila
>gi|1036717554|ref|XP_017143062.1| PREDICTED: transmembrane GTPase Marf isoform X1 [Drosophila
>gi|1036717617|ref|XP_017143092.1| PREDICTED: probable citrate synthase, mitochondrial isoform
>gi|1036717628|ref|XP_017143101.1| PREDICTED: probable citrate synthase, mitochondrial isoform
>gi|1036717647|ref|XP_017143110.1| PREDICTED: probable citrate synthase, mitochondrial isoform
>gi|1036717694|ref|XP_017143135.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036717712|ref|XP_017143143.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036717676|ref|XP_017143123.1| PREDICTED: zinc finger protein 624 [Drosophila miranda]
>gi|1036734492|ref|XP_017152219.1| PREDICTED: retinol dehydrogenase 12 [Drosophila miranda]
>gi|1036734510|ref|XP_017152232.1| PREDICTED: LOW QUALITY PROTEIN: probable myosin light chain
>gi|1036734427|ref|XP_017152198.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036734445|ref|XP_017152206.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036734526|ref|XP_017152246.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036755381|ref|XP_017156613.1| PREDICTED: alpha-tocopherol transfer protein [Drosophila mir
>gi|1036713245|ref|XP_017140494.1| PREDICTED: E3 ubiquitin-protein ligase highwire isoform X2
>gi|1036713228|ref|XP_017140487.1| PREDICTED: E3 ubiquitin-protein ligase highwire isoform X1
>gi|1036713262|ref|XP_017140506.1| PREDICTED: uncharacterized protein LOC108154673 [Drosophila
>gi|1036713278|ref|XP_017140517.1| PREDICTED: uncharacterized protein LOC108154684 [Drosophila
>gi|1036713295|ref|XP_017140527.1| PREDICTED: uncharacterized protein LOC108154684 [Drosophila
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>gi|1036746647|ref|XP_017156121.1| PREDICTED: transmembrane emp24 domain-containing protein 5
>gi|1036746666|ref|XP_017156122.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomple
>gi|1036728171|ref|XP_017148819.1| PREDICTED: probable DNA replication complex GINS protein PSI
>gi|1036728149|ref|XP_017148809.1| PREDICTED: protein FAM57B [Drosophila miranda]
>gi|1036698255|ref|XP_017137402.1| PREDICTED: uncharacterized protein LOC108152276 [Drosophila
>gi|1036728131|ref|XP_017148795.1| PREDICTED: proline-rich protein 36 [Drosophila miranda]
>gi|1036729098|ref|XP_017149359.1| PREDICTED: plexin domain-containing protein 2 [Drosophila mi
>gi|1036729135|ref|XP_017149386.1| PREDICTED: perlucin-like [Drosophila miranda]
>gi|1036729116|ref|XP_017149373.1| PREDICTED: accessory gland protein Acp29AB-like [Drosophila
>gi|1036754428|ref|XP_017156557.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036745181|ref|XP_017156032.1| PREDICTED: putative hydroxypyruvate isomerase [Drosophila mi
>gi|1036745162|ref|XP_017156031.1| PREDICTED: uncharacterized protein LOC108164680 [Drosophila
>gi|1036709629|ref|XP_017138340.1| PREDICTED: serine/threonine-protein phosphatase 4 catalytic
>gi|1036709870|ref|XP_017138492.1| PREDICTED: general odorant-binding protein 19d [Drosophila m
>gi|1036709817|ref|XP_017138456.1| PREDICTED: general odorant-binding protein 56a [Drosophila m
>gi|1036709851|ref|XP_017138479.1| PREDICTED: uncharacterized protein LOC108153179 [Drosophila
>gi|1036709887|ref|XP_017138504.1| PREDICTED: general odorant-binding protein 19a [Drosophila m
>gi|1036709921|ref|XP_017138528.1| PREDICTED: putative 60S ribosomal protein L33 [Drosophila mi
>gi|1036709733|ref|XP_017138401.1| PREDICTED: uncharacterized protein LOC108153120 [Drosophila
>gi|1036709646|ref|XP_017138352.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC2 [Droso
>gi|1036709716|ref|XP_017138388.1| PREDICTED: syntaxin-16 [Drosophila miranda]
>gi|1036709750|ref|XP_017138412.1| PREDICTED: RNA-binding protein pno1 [Drosophila miranda]
>gi|1036709767|ref|XP_017138426.1| PREDICTED: pre-rRNA-processing protein esf2-like [Drosophila
>gi|1036709834|ref|XP_017138467.1| PREDICTED: AN1-type zinc finger protein 6 [Drosophila mirand
>gi|1036709784|ref|XP_017138438.1| PREDICTED: uncharacterized protein LOC108153144 isoform X1
>gi|1036709800|ref|XP_017138445.1| PREDICTED: uncharacterized protein LOC108153144 isoform X2
>gi|1036709904|ref|XP_017138517.1| PREDICTED: sarcocystatin-A [Drosophila miranda]
>gi|1036709698|ref|XP_017138378.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036709665|ref|XP_017138358.1| PREDICTED: dihydropyrimidine dehydrogenase [NADP(+)] isoform
>gi|1036709682|ref|XP_017138366.1| PREDICTED: dihydropyrimidine dehydrogenase [NADP(+)] isoform
>gi|1036737111|ref|XP_017153732.1| PREDICTED: NCK-interacting protein with SH3 domain [Drosophi
>gi|1036730102|ref|XP_017149981.1| PREDICTED: protein will die slowly [Drosophila miranda]
>gi|1036730116|ref|XP_017149989.1| PREDICTED: protein will die slowly [Drosophila miranda]
>gi|1036730083|ref|XP_017149968.1| PREDICTED: pyruvate dehydrogenase E1 component subunit alpha
>gi|1036730062|ref|XP_017149958.1| PREDICTED: probable pyruvate dehydrogenase E1 component subu
>gi|1036730009|ref|XP_017149929.1| PREDICTED: uncharacterized protein LOC108160432 [Drosophila
>gi|1036730024|ref|XP_017149937.1| PREDICTED: uncharacterized protein LOC108160432 [Drosophila
>gi|1036730043|ref|XP_017149945.1| PREDICTED: uncharacterized protein LOC108160432 [Drosophila
>gi|1036706663|ref|XP_017136472.1| PREDICTED: uncharacterized protein LOC108152006 [Drosophila
>gi|1036703804|ref|XP_017157172.1| PREDICTED: E3 ubiquitin-protein ligase RNF185-like [Drosophi
>gi|1036757871|ref|XP_017156746.1| PREDICTED: putative inorganic phosphate cotransporter [Droso
>gi|1036706143|ref|XP_017136103.1| PREDICTED: uncharacterized protein LOC108151766 [Drosophila
>gi|1036706436|ref|XP_017136302.1| PREDICTED: Krueppel-like factor 6 [Drosophila miranda]
>gi|1036747885|ref|XP_017156195.1| PREDICTED: general transcription factor IIH subunit 3-like
>gi|1036747864|ref|XP_017156194.1| PREDICTED: uncharacterized protein LOC108164786 [Drosophila
>gi|1036747848|ref|XP_017156193.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1036699871|ref|XP_017145044.1| PREDICTED: uncharacterized protein LOC108157453 [Drosophila
>gi|1036739368|ref|XP_017154956.1| PREDICTED: glycine receptor subunit alpha-4 [Drosophila mira
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>gi|1036739384|ref|XP_017154969.1| PREDICTED: zinc finger protein 391 isoform X1 [Drosophila m
>gi|1036739400|ref|XP_017154977.1| PREDICTED: gastrula zinc finger protein XlCGF8.2DB isoform
>gi|1036699886|ref|XP_017145164.1| PREDICTED: nardilysin [Drosophila miranda]
>gi|1036719418|ref|XP_017144118.1| PREDICTED: nardilysin-like isoform X1 [Drosophila miranda]
>gi|1036719436|ref|XP_017144126.1| PREDICTED: nardilysin-like isoform X2 [Drosophila miranda]
>gi|1036719399|ref|XP_017144106.1| PREDICTED: copper-transporting ATPase 1 isoform X2 [Drosoph
>gi|1036719380|ref|XP_017144097.1| PREDICTED: copper-transporting ATPase 1 isoform X1 [Drosoph
>gi|1036741195|ref|XP_017155813.1| PREDICTED: replication protein A 14 kDa subunit A [Drosophil
>gi|1036741177|ref|XP_017155812.1| PREDICTED: ATP-binding cassette sub-family F member 1 [Droso
>gi|1036712680|ref|XP_017140170.1| PREDICTED: circadian locomoter output cycles protein kaput
>gi|1036712646|ref|XP_017140150.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X2 [Dros
>gi|1036712663|ref|XP_017140158.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X3 [Dros
>gi|1036712629|ref|XP_017140142.1| PREDICTED: tyrosine-protein phosphatase 10D isoform X1 [Dros
>gi|1036712714|ref|XP_017140190.1| PREDICTED: uncharacterized protein LOC108154429 [Drosophila
>gi|1036712697|ref|XP_017140180.1| PREDICTED: lysophosphatidylcholine acyltransferase [Drosoph
>gi|1036740908|ref|XP_017155798.1| PREDICTED: cubilin homolog [Drosophila miranda]
>gi|1036740924|ref|XP_017155799.1| PREDICTED: lon protease homolog, mitochondrial-like [Drosopl
>gi|1036703949|ref|XP_017134599.1| PREDICTED: uncharacterized protein LOC108150803 [Drosophila
>gi|1036707589|ref|XP_017137082.1| PREDICTED: LOW QUALITY PROTEIN: glycine-rich cell wall stru
>gi|1036725363|ref|XP_017147111.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725306|ref|XP_017147084.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725287|ref|XP_017147076.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725269|ref|XP_017147066.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725251|ref|XP_017147058.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725327|ref|XP_017147093.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725346|ref|XP_017147102.1| PREDICTED: ral GTPase-activating protein subunit beta isofo
>gi|1036725399|ref|XP_017147141.1| PREDICTED: uncharacterized protein LOC108158881 [Drosophila
>gi|1036699905|ref|XP_017145282.1| PREDICTED: skin secretory protein xP2 [Drosophila miranda]
>gi|1036725382|ref|XP_017147125.1| PREDICTED: ATP synthase subunit delta, mitochondrial [Drosop
>gi|1036723542|ref|XP_017146018.1| PREDICTED: LOW QUALITY PROTEIN: nuclear envelope integral me
>gi|1036723561|ref|XP_017146029.1| PREDICTED: uncharacterized protein LOC108158297 [Drosophila
>gi|1036723580|ref|XP_017146039.1| PREDICTED: uncharacterized protein LOC108158307 [Drosophila
>gi|1036707251|ref|XP_017136854.1| PREDICTED: kunitz-type serine protease inhibitor textilinin-
>gi|1036732685|ref|XP_017151265.1| PREDICTED: myc protein [Drosophila miranda]
>gi|1036699920|ref|XP_017145517.1| PREDICTED: uncharacterized protein LOC108157795 [Drosophila
>gi|1036747236|ref|XP_017156157.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X3 [Dr
>gi|1036747251|ref|XP_017156158.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X3 [Dr
>gi|1036747167|ref|XP_017156153.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X1 [Dr
>gi|1036747184|ref|XP_017156154.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X1 [Dr
>gi|1036747203|ref|XP_017156155.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X1 [Dr
>gi|1036747272|ref|XP_017156159.1| PREDICTED: uncharacterized protein LOC108164760 [Drosophila
>gi|1036747220|ref|XP_017156156.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 isoform X2 [Dr
>gi|1036755573|ref|XP_017156620.1| PREDICTED: glutaredoxin domain-containing cysteine-rich prot
>gi|1036705085|ref|XP_017135370.1| PREDICTED: calumenin-B-like [Drosophila miranda]
>gi|1036750995|ref|XP_017156352.1| PREDICTED: sodium-dependent neutral amino acid transporter
>gi|1036726707|ref|XP_017147957.1| PREDICTED: alanyl-tRNA editing protein Aarsd1-B [Drosophila
>gi|1036726757|ref|XP_017147996.1| PREDICTED: uncharacterized protein C19orf52 [Drosophila mira
>gi|1036726776|ref|XP_017148004.1| PREDICTED: uncharacterized protein C19orf52 [Drosophila mira
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>gi|1036726795|ref|XP_017148013.1| PREDICTED: uncharacterized protein C19orf52 [Drosophila mira
>gi|1036726740|ref|XP_017147981.1| PREDICTED: uncharacterized protein LOC108159314 [Drosophila
>gi|1036726814|ref|XP_017148024.1| PREDICTED: 40S ribosomal protein S28 [Drosophila miranda]
>gi|1036726723|ref|XP_017147969.1| PREDICTED: translocon-associated protein subunit alpha [Dros
>gi|1036699936|ref|XP_017145583.1| PREDICTED: LOW QUALITY PROTEIN: ran GTPase-activating protei
>gi|1036737740|ref|XP_017154082.1| PREDICTED: uncharacterized protein LOC108163349 [Drosophila
>gi|1036737720|ref|XP_017154072.1| PREDICTED: uncharacterized protein LOC108163342 [Drosophila
>gi|1036706351|ref|XP_017136241.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosoph
>gi|1036713311|ref|XP_017140538.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1036745613|ref|XP_017156058.1| PREDICTED: B-cell CLL/lymphoma 7 protein family member A [Dr
>gi|1036720303|ref|XP_017144564.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
>gi|1036720350|ref|XP_017144580.1| PREDICTED: palmitoyltransferase ZDHHC5 isoform X3 [Drosophil
>gi|1036720322|ref|XP_017144572.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036720369|ref|XP_017144589.1| PREDICTED: palmitoyltransferase ZDHHC5 isoform X4 [Drosophil
>gi|1036720407|ref|XP_017144605.1| PREDICTED: probable protein S-acyltransferase 2 isoform X6
>gi|1036720426|ref|XP_017144613.1| PREDICTED: palmitoyltransferase ZDHHC9 isoform X7 [Drosophil
>gi|1036720390|ref|XP_017144597.1| PREDICTED: neurogenic protein mastermind isoform X5 [Drosoph
>gi|1036720285|ref|XP_017144552.1| PREDICTED: von Willebrand factor A domain-containing protei
>gi|1036720507|ref|XP_017144658.1| PREDICTED: protein CWC15 homolog [Drosophila miranda]
>gi|1036720445|ref|XP_017144625.1| PREDICTED: protein lozenge [Drosophila miranda]
>gi|1036720465|ref|XP_017144636.1| PREDICTED: protein lozenge [Drosophila miranda]
>gi|1036720485|ref|XP_017144645.1| PREDICTED: protein lozenge [Drosophila miranda]
>gi|1036731878|ref|XP_017150882.1| PREDICTED: maestro heat-like repeat-containing protein fami
>gi|1036731897|ref|XP_017150893.1| PREDICTED: aladin [Drosophila miranda]
>gi|1036703130|ref|XP_017156793.1| PREDICTED: uncharacterized protein LOC108165265 isoform X1
>gi|1036703149|ref|XP_017156798.1| PREDICTED: uncharacterized protein LOC108165265 isoform X2
>gi|1036702601|ref|XP_017156471.1| PREDICTED: CXXC-type zinc finger protein 1-like [Drosophila
>gi|1036702454|ref|XP_017156392.1| PREDICTED: uncharacterized protein LOC108164915 isoform X1
>gi|1036702470|ref|XP_017156400.1| PREDICTED: uncharacterized protein LOC108164915 isoform X1
>gi|1036702487|ref|XP_017156408.1| PREDICTED: uncharacterized protein LOC108164915 isoform X2
>gi|1036702617|ref|XP_017156484.1| PREDICTED: synembryn [Drosophila miranda]
>gi|1036702567|ref|XP_017156450.1| PREDICTED: kelch-like protein 5 isoform X1 [Drosophila miran
>gi|1036702584|ref|XP_017156459.1| PREDICTED: kelch-like protein 5 isoform X2 [Drosophila miran
>gi|1036702649|ref|XP_017156509.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702633|ref|XP_017156497.1| PREDICTED: uncharacterized protein LOC108164982 [Drosophila
>gi|1036699953|ref|XP_017145690.1| PREDICTED: LOW QUALITY PROTEIN: pickpocket protein 28 [Dros
>gi|1036702506|ref|XP_017156419.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X1 [Drosop
>gi|1036702552|ref|XP_017156436.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X3 [Drosop
>gi|1036702535|ref|XP_017156427.1| PREDICTED: gamma-glutamyltranspeptidase 1 isoform X2 [Drosop
>gi|1036702666|ref|XP_017156522.1| PREDICTED: uncharacterized protein LOC108164998 [Drosophila
>gi|1036702437|ref|XP_017156381.1| PREDICTED: sodium/potassium-transporting ATPase subunit alp
>gi|1036702405|ref|XP_017156358.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702390|ref|XP_017156350.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702421|ref|XP_017156368.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702753|ref|XP_017156573.1| PREDICTED: dual specificity mitogen-activated protein kinase
>gi|1036702772|ref|XP_017156585.1| PREDICTED: probable alpha-aspartyl dipeptidase [Drosophila
>gi|1036702698|ref|XP_017156540.1| PREDICTED: chondroitin sulfate glucuronyltransferase [Droso
>gi|1036702787|ref|XP_017156599.1| PREDICTED: soma ferritin [Drosophila miranda]
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>gi|1036702717|ref|XP_017156550.1| PREDICTED: actin-related protein 6 [Drosophila miranda]
>gi|1036702838|ref|XP_017156627.1| PREDICTED: iron-sulfur cluster assembly 1 homolog, mitochond
>gi|1036702821|ref|XP_017156619.1| PREDICTED: iron-sulfur cluster assembly 1 homolog, mitochond
>gi|1036702804|ref|XP_017156609.1| PREDICTED: DNL-type zinc finger protein [Drosophila miranda]
>gi|1036702734|ref|XP_017156562.1| PREDICTED: uncharacterized protein LOC108165033 [Drosophila
>gi|1036717282|ref|XP_017142918.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X4 [Dro
>gi|1036717242|ref|XP_017142900.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X2 [Dro
>gi|1036717301|ref|XP_017142927.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X5 [Dro
>gi|1036717263|ref|XP_017142909.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X3 [Dro
>gi|1036717224|ref|XP_017142892.1| PREDICTED: E3 ubiquitin-protein ligase HUWE1 isoform X1 [Dro
>gi|1036717322|ref|XP_017142938.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subur
>gi|1036753514|ref|XP_017156503.1| PREDICTED: uncharacterized protein LOC108164988 [Drosophila
>gi|1036736826|ref|XP_017153559.1| PREDICTED: ATP-binding cassette sub-family F member 2 [Dros
>gi|1036748164|ref|XP_017156208.1| PREDICTED: uncharacterized protein LOC108164795 [Drosophila
>gi|1036734022|ref|XP_017151984.1| PREDICTED: CDP-diacylglycerol--inositol 3-phosphatidyltrans
>gi|1036734041|ref|XP_017151997.1| PREDICTED: mitochondrial inner membrane protease subunit 1
>gi|1036733966|ref|XP_017151952.1| PREDICTED: nudix hydrolase 8 [Drosophila miranda]
>gi|1036733985|ref|XP_017151962.1| PREDICTED: nudix hydrolase 8 [Drosophila miranda]
>gi|1036734003|ref|XP_017151971.1| PREDICTED: nudix hydrolase 8 [Drosophila miranda]
>gi|1036699966|ref|XP_017145814.1| PREDICTED: uncharacterized protein LOC108158109 [Drosophila
>gi|1036755682|ref|XP_017156626.1| PREDICTED: uncharacterized protein LOC108165086 [Drosophila
>gi|1036699987|ref|XP_017145930.1| PREDICTED: visual system homeobox 2, partial [Drosophila mi
>gi|1036750499|ref|XP_017156323.1| PREDICTED: homeotic protein ocelliless isoform X3 [Drosophil
>gi|1036750518|ref|XP_017156324.1| PREDICTED: homeotic protein ocelliless isoform X4 [Drosophil
>gi|1036750537|ref|XP_017156325.1| PREDICTED: uncharacterized protein LOC108164886 [Drosophila
>gi|1036750460|ref|XP_017156320.1| PREDICTED: diencephalon/mesencephalon homeobox protein 1 iso
>gi|1036750477|ref|XP_017156321.1| PREDICTED: diencephalon/mesencephalon homeobox protein 1 iso
>gi|1036750442|ref|XP_017156319.1| PREDICTED: uncharacterized protein LOC108164884 [Drosophila
>gi|1036750425|ref|XP_017156318.1| PREDICTED: N-chimaerin [Drosophila miranda]
>gi|1036736551|ref|XP_017153409.1| PREDICTED: myosin-2 essential light chain isoform X2 [Drosop
>gi|1036736538|ref|XP_017153400.1| PREDICTED: myosin-2 essential light chain isoform X1 [Drosop
>gi|1036736486|ref|XP_017153372.1| PREDICTED: ribonucleases P/MRP protein subunit POP1 [Drosoph
>gi|1036736505|ref|XP_017153381.1| PREDICTED: ribonucleases P/MRP protein subunit POP1 [Drosoph
>gi|1036736522|ref|XP_017153389.1| PREDICTED: ribonucleases P/MRP protein subunit POP1 [Drosoph
>gi|1036737952|ref|XP_017154196.1| PREDICTED: uncharacterized protein LOC108163366 isoform X12
>gi|1036737883|ref|XP_017154162.1| PREDICTED: uncharacterized protein LOC108163366 isoform X8
>gi|1036737936|ref|XP_017154186.1| PREDICTED: uncharacterized protein LOC108163366 isoform X11
>gi|1036737811|ref|XP_017154129.1| PREDICTED: uncharacterized protein LOC108163366 isoform X4
>gi|1036737849|ref|XP_017154146.1| PREDICTED: uncharacterized protein LOC108163366 isoform X6
>gi|1036737865|ref|XP_017154154.1| PREDICTED: uncharacterized protein LOC108163366 isoform X7
>gi|1036737793|ref|XP_017154122.1| PREDICTED: uncharacterized protein LOC108163366 isoform X3
>gi|1036737829|ref|XP_017154137.1| PREDICTED: uncharacterized protein LOC108163366 isoform X5
>gi|1036737775|ref|XP_017154117.1| PREDICTED: uncharacterized protein LOC108163366 isoform X2
>gi|1036737758|ref|XP_017154110.1| PREDICTED: uncharacterized protein LOC108163366 isoform X1
>gi|1036737970|ref|XP_017154204.1| PREDICTED: uncharacterized protein LOC108163366 isoform X13
>gi|1036737918|ref|XP_017154178.1| PREDICTED: uncharacterized protein LOC108163366 isoform X10
>gi|1036737901|ref|XP_017154170.1| PREDICTED: uncharacterized protein LOC108163366 isoform X9
>gi|1036749952|ref|XP_017156295.1| PREDICTED: uncharacterized protein LOC108164866 isoform X2
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>gi|1036749879|ref|XP_017156290.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749897|ref|XP_017156291.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749916|ref|XP_017156293.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749933|ref|XP_017156294.1| PREDICTED: uncharacterized protein LOC108164866 isoform X1
>gi|1036749971|ref|XP_017156296.1| PREDICTED: uncharacterized protein LOC108164866 isoform X3
>gi|1036753758|ref|XP_017156517.1| PREDICTED: ras-related protein Rab-21 [Drosophila miranda]
>gi|1036729882|ref|XP_017149826.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036729901|ref|XP_017149833.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036729924|ref|XP_017149842.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036729747|ref|XP_017149758.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729765|ref|XP_017149767.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729784|ref|XP_017149775.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729805|ref|XP_017149784.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729821|ref|XP_017149796.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729842|ref|XP_017149805.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729859|ref|XP_017149813.1| PREDICTED: protein stoned-A [Drosophila miranda]
>gi|1036729730|ref|XP_017149745.1| PREDICTED: protein stoned-B [Drosophila miranda]
>gi|1036756763|ref|XP_017156689.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036756746|ref|XP_017156688.1| PREDICTED: farnesol dehydrogenase [Drosophila miranda]
>gi|1036704064|ref|XP_017134668.1| PREDICTED: uncharacterized protein LOC108150871 [Drosophila
>gi|1036705831|ref|XP_017135865.1| PREDICTED: 39 kDa FK506-binding nuclear protein-like [Drosop
>gi|1036700002|ref|XP_017146043.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1036751789|ref|XP_017156396.1| PREDICTED: uncharacterized protein LOC108164921 [Drosophila
>gi|1036751771|ref|XP_017156395.1| PREDICTED: uncharacterized protein LOC108164920 [Drosophila
>gi|1036735990|ref|XP_017153071.1| PREDICTED: protein folded gastrulation [Drosophila miranda]
>gi|1036736007|ref|XP_017153079.1| PREDICTED: protein folded gastrulation [Drosophila miranda]
>gi|1036736023|ref|XP_017153087.1| PREDICTED: protein folded gastrulation [Drosophila miranda]
>gi|1036757551|ref|XP_017156731.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1036706245|ref|XP_017136170.1| PREDICTED: uncharacterized protein LOC108151818 [Drosophila
>gi|1036754546|ref|XP_017156564.1| PREDICTED: uncharacterized protein LOC108165038 [Drosophila
>gi|1036754565|ref|XP_017156565.1| PREDICTED: uncharacterized protein LOC108165038 [Drosophila
>gi|1036739929|ref|XP_017155271.1| PREDICTED: uncharacterized protein LOC108164168 isoform X1
>gi|1036739946|ref|XP_017155280.1| PREDICTED: uncharacterized protein LOC108164168 isoform X2
>gi|1036739964|ref|XP_017155291.1| PREDICTED: uncharacterized protein LOC108164181 isoform X1
>gi|1036739982|ref|XP_017155298.1| PREDICTED: uncharacterized protein LOC108164181 isoform X2
>gi|1036740000|ref|XP_017155307.1| PREDICTED: uncharacterized protein LOC108164181 isoform X2
>gi|1036740074|ref|XP_017155347.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
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>gi|1036740216|ref|XP_017155404.1| PREDICTED: uncharacterized protein LOC108164224 [Drosophila
>gi|1036740022|ref|XP_017155320.1| PREDICTED: probable isoaspartyl peptidase/L-asparaginase GA2
>gi|1036740056|ref|XP_017155339.1| PREDICTED: UPF0454 protein C12orf49 homolog isoform X2 [Dros
>gi|1036740040|ref|XP_017155331.1| PREDICTED: UPF0454 protein C12orf49 homolog isoform X1 [Dros
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>gi|1036715291|ref|XP_017141775.1| PREDICTED: dnaJ homolog subfamily C member 25 homolog [Droso
>gi|1036715255|ref|XP_017141751.1| PREDICTED: neurotrophin receptor-interacting factor homolog
>gi|1036715272|ref|XP_017141763.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036715165|ref|XP_017141706.1| PREDICTED: chondroitin sulfate synthase 1 isoform X1 [Droso
>gi|1036715184|ref|XP_017141716.1| PREDICTED: chondroitin sulfate synthase 1 isoform X1 [Droso
>gi|1036715203|ref|XP_017141723.1| PREDICTED: chondroitin sulfate synthase 1 isoform X1 [Droso
>gi|1036715219|ref|XP_017141732.1| PREDICTED: chondroitin sulfate synthase 3 isoform X2 [Droso
>gi|1036715237|ref|XP_017141740.1| PREDICTED: chondroitin sulfate synthase 3 isoform X3 [Droso
>gi|1036733508|ref|XP_017151672.1| PREDICTED: dorsal-ventral patterning protein Sog [Drosophila
>gi|1036754675|ref|XP_017156571.1| PREDICTED: uncharacterized protein LOC108165044 [Drosophila
>gi|1036706604|ref|XP_017136411.1| PREDICTED: uncharacterized protein LOC108151968 [Drosophila
>gi|1036700017|ref|XP_017146158.1| PREDICTED: uncharacterized protein LOC108158376 [Drosophila
>gi|1036700032|ref|XP_017146280.1| PREDICTED: endothelin-converting enzyme-like 1 [Drosophila
>gi|1036753950|ref|XP_017156529.1| PREDICTED: sex peptide receptor [Drosophila miranda]
>gi|1036700047|ref|XP_017146420.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036751051|ref|XP_017156355.1| PREDICTED: hornerin [Drosophila miranda]
>gi|1036751070|ref|XP_017156356.1| PREDICTED: lysozyme [Drosophila miranda]
>gi|1036733076|ref|XP_017151440.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 16 isoform
>gi|1036733060|ref|XP_017151433.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 16 isoform
>gi|1036711610|ref|XP_017139576.1| PREDICTED: serine/threonine-protein kinase Smg1 isoform X1
>gi|1036711629|ref|XP_017139583.1| PREDICTED: serine/threonine-protein kinase Smg1 isoform X2
>gi|1036711646|ref|XP_017139594.1| PREDICTED: beta-1 adrenergic receptor [Drosophila miranda]
>gi|1036711663|ref|XP_017139604.1| PREDICTED: 40S ribosomal protein S23-like [Drosophila miran
>gi|1036711733|ref|XP_017139642.1| PREDICTED: E3 ubiquitin-protein ligase ZNRF2 [Drosophila mi
>gi|1036711748|ref|XP_017139649.1| PREDICTED: E3 ubiquitin-protein ligase ZNRF2 [Drosophila mi
>gi|1036711680|ref|XP_017139615.1| PREDICTED: TATA element modulatory factor isoform X1 [Droso
>gi|1036711697|ref|XP_017139622.1| PREDICTED: TATA element modulatory factor isoform X2 [Droso
>gi|1036711716|ref|XP_017139632.1| PREDICTED: chitinase-3-like protein 1 [Drosophila miranda]
>gi|1036711761|ref|XP_017139663.1| PREDICTED: tRNA 2'-phosphotransferase 1 [Drosophila miranda]
>gi|1036742934|ref|XP_017155914.1| PREDICTED: zinc finger protein 236 [Drosophila miranda]
>gi|1036700064|ref|XP_017146550.1| PREDICTED: mitochondrial import receptor subunit TOM70-like
>gi|1036706469|ref|XP_017136323.1| PREDICTED: probable serine/threonine-protein kinase mps1 [D
>gi|1036727380|ref|XP_017148366.1| PREDICTED: glutathione synthetase-like isoform X1 [Drosophil
>gi|1036727398|ref|XP_017148376.1| PREDICTED: glutathione synthetase-like isoform X1 [Drosophil
>gi|1036727417|ref|XP_017148383.1| PREDICTED: glutathione synthetase-like isoform X2 [Drosophil
>gi|1036727436|ref|XP_017148392.1| PREDICTED: glutathione synthetase-like isoform X3 [Drosophil
>gi|1036727472|ref|XP_017148404.1| PREDICTED: glutathione synthetase-like isoform X4 [Drosophil
>gi|1036727489|ref|XP_017148418.1| PREDICTED: LOW QUALITY PROTEIN: mitochondrial fission proces
>gi|1036727507|ref|XP_017148434.1| PREDICTED: hsc70-interacting protein 1-like [Drosophila mira
>gi|1036754331|ref|XP_017156551.1| PREDICTED: uncharacterized protein LOC108165026 [Drosophila
>gi|1036754348|ref|XP_017156552.1| PREDICTED: uncharacterized protein LOC108165026 [Drosophila
>gi|1036723616|ref|XP_017146054.1| PREDICTED: uncharacterized protein LOC108158321 isoform X1
>gi|1036723632|ref|XP_017146061.1| PREDICTED: uncharacterized protein LOC108158321 isoform X2
>gi|1036723650|ref|XP_017146070.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036700080|ref|XP_017146676.1| PREDICTED: mucin-19-like [Drosophila miranda]
>gi|1036703230|ref|XP_017156829.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036718884|ref|XP_017143802.1| PREDICTED: malate dehydrogenase-like [Drosophila miranda]
>gi|1036718865|ref|XP_017143788.1| PREDICTED: malate dehydrogenase, mitochondrial-like [Drosoph
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>gi|1036718827|ref|XP_017143762.1| PREDICTED: uncharacterized protein LOC108156656 [Drosophila
>gi|1036718846|ref|XP_017143776.1| PREDICTED: mitoguardin [Drosophila miranda]
>gi|1036718904|ref|XP_017143816.1| PREDICTED: uncharacterized protein LOC108156683 [Drosophila
>gi|1036711076|ref|XP_017139312.1| PREDICTED: uncharacterized protein LOC108153694 [Drosophila
>gi|1036711106|ref|XP_017139334.1| PREDICTED: GPI transamidase component PIG-T [Drosophila mira
>gi|1036711085|ref|XP_017139323.1| PREDICTED: trithorax group protein osa [Drosophila miranda]
>gi|1036711121|ref|XP_017139346.1| PREDICTED: CTD nuclear envelope phosphatase 1 homolog isofor
>gi|1036711138|ref|XP_017139353.1| PREDICTED: CTD nuclear envelope phosphatase 1 homolog isofor
>gi|1036736902|ref|XP_017153613.1| PREDICTED: SRA stem-loop-interacting RNA-binding protein, m
>gi|1036736882|ref|XP_017153600.1| PREDICTED: acyl-CoA-binding domain-containing protein 6 [Dr
>gi|1036736864|ref|XP_017153585.1| PREDICTED: 26S protease regulatory subunit 8 [Drosophila mi
>gi|1036736845|ref|XP_017153572.1| PREDICTED: ATP-binding cassette sub-family A member 2 [Dros
>gi|1036737563|ref|XP_017153983.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofor
>gi|1036737579|ref|XP_017153992.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofor
>gi|1036737598|ref|XP_017154001.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofor
>gi|1036737618|ref|XP_017154008.1| PREDICTED: ATP-binding cassette sub-family A member 3 isofor
>gi|1036755855|ref|XP_017156637.1| PREDICTED: ATP-binding cassette sub-family A member 3 [Dros
>gi|1036739225|ref|XP_017154858.1| PREDICTED: loricrin isoform X1 [Drosophila miranda]
>gi|1036739243|ref|XP_017154866.1| PREDICTED: pro-resilin isoform X2 [Drosophila miranda]
>gi|1036705755|ref|XP_017135808.1| PREDICTED: uncharacterized protein LOC108151618 [Drosophila
>gi|1036730379|ref|XP_017150158.1| PREDICTED: ATP-binding cassette sub-family A member 7 isofor
>gi|1036730342|ref|XP_017150144.1| PREDICTED: ATP-binding cassette sub-family A member 13 isof
>gi|1036730360|ref|XP_017150150.1| PREDICTED: ATP-binding cassette sub-family A member 13 isof
>gi|1036730396|ref|XP_017150171.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1036730413|ref|XP_017150179.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1036730429|ref|XP_017150188.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1036748852|ref|XP_017156233.1| PREDICTED: uncharacterized protein LOC108164823 [Drosophila
>gi|1036748871|ref|XP_017156234.1| PREDICTED: uncharacterized protein LOC108164823 [Drosophila
>gi|1036700097|ref|XP_017146813.1| PREDICTED: uncharacterized protein LOC108158723 [Drosophila
>gi|1036700111|ref|XP_017146945.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036748890|ref|XP_017156235.1| PREDICTED: NEDD8-specific protease 2 [Drosophila miranda]
>gi|1036727782|ref|XP_017148596.1| PREDICTED: blood vessel epicardial substance isoform X3 [Dr
>gi|1036727764|ref|XP_017148587.1| PREDICTED: blood vessel epicardial substance isoform X2 [Dr
>gi|1036727725|ref|XP_017148570.1| PREDICTED: blood vessel epicardial substance isoform X1 [Dr
>gi|1036727747|ref|XP_017148578.1| PREDICTED: blood vessel epicardial substance isoform X1 [Dr
>gi|1036727707|ref|XP_017148558.1| PREDICTED: transcription factor kayak [Drosophila miranda]
>gi|1036735348|ref|XP_017152720.1| PREDICTED: dolichyl-diphosphooligosaccharide--protein glycos
>gi|1036735369|ref|XP_017152733.1| PREDICTED: uncharacterized protein LOC108162488 isoform X1
>gi|1036735388|ref|XP_017152741.1| PREDICTED: uncharacterized protein LOC108162488 isoform X2
>gi|1036744643|ref|XP_017156001.1| PREDICTED: cystathionine beta-synthase [Drosophila miranda]
>gi|1036734725|ref|XP_017152381.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X
>gi|1036734706|ref|XP_017152371.1| PREDICTED: microsomal glutathione S-transferase 1 isoform X
>gi|1036734670|ref|XP_017152351.1| PREDICTED: uncharacterized protein LOC108162231 isoform X1
>gi|1036734688|ref|XP_017152360.1| PREDICTED: uncharacterized protein LOC108162231 isoform X2
>gi|1036734742|ref|XP_017152393.1| PREDICTED: probable nuclear transport factor 2 [Drosophila
>gi|1036739193|ref|XP_017154835.1| PREDICTED: zinc finger protein 134 [Drosophila miranda]
>gi|1036739209|ref|XP_017154848.1| PREDICTED: NAD(P)H-hydrate epimerase [Drosophila miranda]
>gi|1036739173|ref|XP_017154822.1| PREDICTED: RNA-binding protein NOB1 [Drosophila miranda]
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>gi|1036720170|ref|XP_017144493.1| PREDICTED: trithorax group protein osa isoform X1 [Drosophil
>gi|1036720189|ref|XP_017144502.1| PREDICTED: collagen alpha-1(I) chain isoform X2 [Drosophila
>gi|1036720209|ref|XP_017144512.1| PREDICTED: collagen alpha-1(I) chain isoform X3 [Drosophila
>gi|1036720266|ref|XP_017144539.1| PREDICTED: uncharacterized protein LOC108157132 isoform X3
>gi|1036720229|ref|XP_017144523.1| PREDICTED: uncharacterized protein LOC108157132 isoform X1
>gi|1036720247|ref|XP_017144531.1| PREDICTED: uncharacterized protein LOC108157132 isoform X2
>gi|1036698274|ref|XP_017138640.1| PREDICTED: putative gustatory receptor 9a [Drosophila mirand
>gi|1036755801|ref|XP_017156634.1| PREDICTED: vitellogenin-1 [Drosophila miranda]
>gi|1036756202|ref|XP_017156656.1| PREDICTED: vitellogenin-2 [Drosophila miranda]
>gi|1036742628|ref|XP_017155897.1| PREDICTED: mucin-5AC isoform X7 [Drosophila miranda]
>gi|1036742610|ref|XP_017155895.1| PREDICTED: mucin-5AC isoform X6 [Drosophila miranda]
>gi|1036742538|ref|XP_017155891.1| PREDICTED: uncharacterized protein LOC108164581 isoform X2
>gi|1036742520|ref|XP_017155890.1| PREDICTED: uncharacterized protein LOC108164581 isoform X1
>gi|1036742592|ref|XP_017155894.1| PREDICTED: mucin-5AC isoform X5 [Drosophila miranda]
>gi|1036742557|ref|XP_017155892.1| PREDICTED: mucin-5AC isoform X3 [Drosophila miranda]
>gi|1036742644|ref|XP_017155898.1| PREDICTED: serine-rich adhesin for platelets isoform X8 [Dro
>gi|1036742576|ref|XP_017155893.1| PREDICTED: mucin-5AC isoform X4 [Drosophila miranda]
>gi|1036739678|ref|XP_017155131.1| PREDICTED: DNA replication ATP-dependent helicase/nuclease
>gi|1036705629|ref|XP_017135713.1| PREDICTED: basic helix-loop-helix transcription factor scle
>gi|1036752664|ref|XP_017156449.1| PREDICTED: keratin, type I cytoskeletal 10 isoform X3 [Droso
>gi|1036752628|ref|XP_017156447.1| PREDICTED: keratin, type I cytoskeletal 10 isoform X1 [Droso
>gi|1036752701|ref|XP_017156452.1| PREDICTED: uncharacterized protein LOC108164957 isoform X5
>gi|1036752719|ref|XP_017156453.1| PREDICTED: uncharacterized protein LOC108164957 isoform X5
>gi|1036752645|ref|XP_017156448.1| PREDICTED: uncharacterized protein DDB_G0280205 isoform X2
>gi|1036752682|ref|XP_017156451.1| PREDICTED: uncharacterized protein LOC108164957 isoform X4
>gi|1036752735|ref|XP_017156454.1| PREDICTED: uncharacterized protein LOC108164957 isoform X6
>gi|1036700126|ref|XP_017147072.1| PREDICTED: uncharacterized protein LOC108158833, partial [D
>gi|1036755125|ref|XP_017156598.1| PREDICTED: uncharacterized protein LOC108165061 isoform X2
>gi|1036755087|ref|XP_017156596.1| PREDICTED: uncharacterized protein LOC108165061 isoform X1
>gi|1036755106|ref|XP_017156597.1| PREDICTED: uncharacterized protein LOC108165061 isoform X1
>gi|1036755069|ref|XP_017156595.1| PREDICTED: microfibril-associated glycoprotein 4 isoform X2
>gi|1036755050|ref|XP_017156594.1| PREDICTED: microfibril-associated glycoprotein 4 isoform X1
>gi|1036755014|ref|XP_017156592.1| PREDICTED: fibrinogen-like protein A isoform X1 [Drosophila
>gi|1036755031|ref|XP_017156593.1| PREDICTED: fibrinogen-like protein A isoform X2 [Drosophila
>gi|1036746190|ref|XP_017156091.1| PREDICTED: chitinase-like protein Idgf4 [Drosophila miranda]
>gi|1036700141|ref|XP_017147205.1| PREDICTED: lipoprotein lipase [Drosophila miranda]
>gi|1036756109|ref|XP_017156650.1| PREDICTED: endonuclease G, mitochondrial-like isoform X2 [D
>gi|1036756090|ref|XP_017156649.1| PREDICTED: endonuclease G, mitochondrial-like isoform X1 [D
>gi|1036748145|ref|XP_017156207.1| PREDICTED: uncharacterized protein LOC108164794 [Drosophila
>gi|1036748126|ref|XP_017156205.1| PREDICTED: uncharacterized protein LOC108164793 [Drosophila
>gi|1036703416|ref|XP_017156936.1| PREDICTED: uncharacterized protein LOC108165405 [Drosophila
>gi|1036704401|ref|XP_017134870.1| PREDICTED: uncharacterized protein LOC108151018 [Drosophila
>gi|1036704907|ref|XP_017135229.1| PREDICTED: uncharacterized protein LOC108151237 [Drosophila
>gi|1036703114|ref|XP_017156773.1| PREDICTED: uncharacterized protein LOC108165243 [Drosophila
>gi|1036756583|ref|XP_017156679.1| PREDICTED: PERQ amino acid-rich with GYF domain-containing p
>gi|1036704530|ref|XP_017134954.1| PREDICTED: uncharacterized protein LOC108151079 [Drosophila
>gi|1036754395|ref|XP_017156555.1| PREDICTED: uncharacterized protein LOC108165029 [Drosophila
>gi|1036754412|ref|XP_017156556.1| PREDICTED: uncharacterized protein LOC108165030 [Drosophila
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>gi|1036754380|ref|XP_017156554.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1036702984|ref|XP_017156713.1| PREDICTED: uncharacterized protein LOC108165154 isoform X1
>gi|1036703002|ref|XP_017156721.1| PREDICTED: uncharacterized protein LOC108165154 isoform X1
>gi|1036703018|ref|XP_017156728.1| PREDICTED: uncharacterized protein LOC108165154 isoform X2
>gi|1036751844|ref|XP_017156399.1| PREDICTED: carbonic anhydrase-related protein 10 [Drosophila
>gi|1036751880|ref|XP_017156402.1| PREDICTED: N-alpha-acetyltransferase 38, NatC auxiliary subu
>gi|1036751863|ref|XP_017156401.1| PREDICTED: N-alpha-acetyltransferase 38, NatC auxiliary subu
>gi|1036741596|ref|XP_017155839.1| PREDICTED: obg-like ATPase 1 [Drosophila miranda]
>gi|1036741615|ref|XP_017155840.1| PREDICTED: obg-like ATPase 1 [Drosophila miranda]
>gi|1036749185|ref|XP_017156252.1| PREDICTED: transcription factor Sp9 isoform X2 [Drosophila m
>gi|1036749152|ref|XP_017156250.1| PREDICTED: transcription factor Sp9 isoform X1 [Drosophila m
>gi|1036749167|ref|XP_017156251.1| PREDICTED: transcription factor Sp9 isoform X1 [Drosophila m
>gi|1036745200|ref|XP_017156033.1| PREDICTED: transcription factor btd [Drosophila miranda]
>gi|1036720130|ref|XP_017144471.1| PREDICTED: histone acetyltransferase p300 isoform X1 [Drosop
>gi|1036720150|ref|XP_017144480.1| PREDICTED: histone acetyltransferase p300 isoform X2 [Drosop
>gi|1036754470|ref|XP_017156559.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036724778|ref|XP_017146786.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 9
>gi|1036724743|ref|XP_017146764.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 9
>gi|1036724761|ref|XP_017146775.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 9
>gi|1036731645|ref|XP_017150820.1| PREDICTED: E3 ubiquitin-protein ligase HECW2 isoform X1 [Dro
>gi|1036731681|ref|XP_017150825.1| PREDICTED: E3 ubiquitin-protein ligase HECW2 isoform X3 [Dro
>gi|1036731664|ref|XP_017150821.1| PREDICTED: E3 ubiquitin-protein ligase HECW2 isoform X2 [Dro
>gi|1036750209|ref|XP_017156308.1| PREDICTED: uncharacterized protein LOC108164874 [Drosophila
>gi|1036745509|ref|XP_017156052.1| PREDICTED: lysine--tRNA ligase isoform X1 [Drosophila miran
>gi|1036745525|ref|XP_017156053.1| PREDICTED: lysine--tRNA ligase isoform X2 [Drosophila miran
>gi|1036717368|ref|XP_017142962.1| PREDICTED: probable histone-lysine N-methyltransferase CG17
>gi|1036717387|ref|XP_017142973.1| PREDICTED: fatty acid hydroxylase domain-containing protein
>gi|1036717406|ref|XP_017142985.1| PREDICTED: glutathione S-transferase theta-1 [Drosophila mi
>gi|1036744381|ref|XP_017155989.1| PREDICTED: kinesin-like protein Nod [Drosophila miranda]
>gi|1036744400|ref|XP_017155990.1| PREDICTED: uncharacterized protein LOC108164649 [Drosophila
>gi|1036753741|ref|XP_017156516.1| PREDICTED: nuclease [Drosophila miranda]
>gi|1036753650|ref|XP_017156511.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753668|ref|XP_017156512.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753686|ref|XP_017156513.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753705|ref|XP_017156514.1| PREDICTED: uncharacterized protein LOC108164995 isoform X1
>gi|1036753724|ref|XP_017156515.1| PREDICTED: uncharacterized protein LOC108164995 isoform X2
>gi|1036724391|ref|XP_017146525.1| PREDICTED: uncharacterized protein LOC108158553 isoform X4
>gi|1036724356|ref|XP_017146508.1| PREDICTED: uncharacterized protein LOC108158553 isoform X2
>gi|1036724373|ref|XP_017146516.1| PREDICTED: uncharacterized protein LOC108158553 isoform X3
>gi|1036724250|ref|XP_017146459.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724270|ref|XP_017146469.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724286|ref|XP_017146477.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724305|ref|XP_017146487.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036724321|ref|XP_017146495.1| PREDICTED: uncharacterized protein LOC108158553 isoform X1
>gi|1036742663|ref|XP_017155899.1| PREDICTED: uncharacterized protein LOC108164583 [Drosophila
>gi|1036710962|ref|XP_017139235.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X2
>gi|1036710945|ref|XP_017139228.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X1
>gi|1036710996|ref|XP_017139251.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X4
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>gi|1036710979|ref|XP_017139243.1| PREDICTED: serine/threonine-protein kinase S6KL isoform X3
>gi|1036711030|ref|XP_017139276.1| PREDICTED: protein bangles and beads [Drosophila miranda]
>gi|1036711046|ref|XP_017139287.1| PREDICTED: autophagy-related protein 101 [Drosophila miranda
>gi|1036711013|ref|XP_017139265.1| PREDICTED: uncharacterized protein LOC108153661 [Drosophila
>gi|1036711065|ref|XP_017139300.1| PREDICTED: coactosin-like protein [Drosophila miranda]
>gi|1036756147|ref|XP_017156652.1| PREDICTED: gastrin/cholecystokinin type B receptor [Drosophi
>gi|1036705770|ref|XP_017135820.1| PREDICTED: cofilin/actin-depolymerizing factor homolog [Dros
>gi|1036700156|ref|XP_017147331.1| PREDICTED: gastrin/cholecystokinin type B receptor [Drosophi
>gi|1036754941|ref|XP_017156588.1| PREDICTED: probable cytochrome P450 308a1 isoform X1 [Droso
>gi|1036754959|ref|XP_017156589.1| PREDICTED: probable cytochrome P450 308a1 isoform X2 [Droso
>gi|1036754978|ref|XP_017156590.1| PREDICTED: probable cytochrome P450 308a1 isoform X3 [Droso
>gi|1036744903|ref|XP_017156015.1| PREDICTED: cytochrome P450 18a1 [Drosophila miranda]
>gi|1036744886|ref|XP_017156013.1| PREDICTED: cytochrome P450 306a1 [Drosophila miranda]
>gi|1036700171|ref|XP_017147467.1| PREDICTED: zinc metalloproteinase nas-7 [Drosophila miranda]
>gi|1036727073|ref|XP_017148167.1| PREDICTED: serine/threonine-protein kinase fused [Drosophila
>gi|1036727039|ref|XP_017148142.1| PREDICTED: C-mannosyltransferase dpy-19 homolog [Drosophila
>gi|1036727056|ref|XP_017148152.1| PREDICTED: C-mannosyltransferase dpy-19 homolog [Drosophila
>gi|1036746173|ref|XP_017156090.1| PREDICTED: inhibitor of growth protein 3 [Drosophila miranda
>gi|1036728732|ref|XP_017149133.1| PREDICTED: glucose-induced degradation protein 8 homolog [Dr
>gi|1036728713|ref|XP_017149120.1| PREDICTED: nucleoporin NUP53 [Drosophila miranda]
>gi|1036728639|ref|XP_017149086.1| PREDICTED: rab11 family-interacting protein 1 isoform X1 [Dr
>gi|1036728675|ref|XP_017149098.1| PREDICTED: rab11 family-interacting protein 1 isoform X2 [Dr
>gi|1036728694|ref|XP_017149108.1| PREDICTED: uncharacterized protein LOC108159941 isoform X3
>gi|1036746156|ref|XP_017156089.1| PREDICTED: protein valois-like [Drosophila miranda]
>gi|1036742880|ref|XP_017155911.1| PREDICTED: tumor necrosis factor receptor superfamily member
>gi|1036742899|ref|XP_017155912.1| PREDICTED: tumor necrosis factor receptor superfamily member
>gi|1036742917|ref|XP_017155913.1| PREDICTED: tumor necrosis factor receptor superfamily member
>gi|1036753121|ref|XP_017156479.1| PREDICTED: uncharacterized protein LOC108164975 isoform X1
>gi|1036753136|ref|XP_017156480.1| PREDICTED: uncharacterized protein LOC108164975 isoform X1
>gi|1036753153|ref|XP_017156481.1| PREDICTED: uncharacterized protein LOC108164975 isoform X1
>gi|1036753170|ref|XP_017156482.1| PREDICTED: uncharacterized protein LOC108164975 isoform X2
>gi|1036728284|ref|XP_017148876.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036728303|ref|XP_017148899.1| PREDICTED: uncharacterized protein LOC108159829 [Drosophila
>gi|1036728247|ref|XP_017148856.1| PREDICTED: histidine--tRNA ligase, cytoplasmic isoform X2 [
>gi|1036728265|ref|XP_017148864.1| PREDICTED: histidine--tRNA ligase, cytoplasmic isoform X3 [
>gi|1036728225|ref|XP_017148847.1| PREDICTED: histidine--tRNA ligase, cytoplasmic isoform X1 [
>gi|1036752173|ref|XP_017156420.1| PREDICTED: gamma-glutamyltranspeptidase 1 [Drosophila miran
>gi|1036738919|ref|XP_017154680.1| PREDICTED: N-acetylgalactosaminyltransferase 7 [Drosophila m
>gi|1036756273|ref|XP_017156660.1| PREDICTED: uncharacterized protein LOC108165112 [Drosophila
>gi|1036756290|ref|XP_017156661.1| PREDICTED: uncharacterized protein LOC108165113 [Drosophila
>gi|1036746772|ref|XP_017156129.1| PREDICTED: uncharacterized protein LOC108164744 isoform X1
>gi|1036746791|ref|XP_017156130.1| PREDICTED: uncharacterized protein LOC108164744 isoform X2
>gi|1036746756|ref|XP_017156128.1| PREDICTED: uncharacterized protein LOC108164743 [Drosophila
>gi|1036746722|ref|XP_017156126.1| PREDICTED: serine protease persephone-like isoform X1 [Dros
>gi|1036746739|ref|XP_017156127.1| PREDICTED: serine protease persephone-like isoform X2 [Dros
>gi|1036746685|ref|XP_017156123.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036746706|ref|XP_017156124.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036700188|ref|XP_017147595.1| PREDICTED: uncharacterized protein LOC108159096 [Drosophila
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>gi|1036757062|ref|XP_017156703.1| PREDICTED: uncharacterized protein LOC108165146 [Drosophila
>gi|1036757043|ref|XP_017156702.1| PREDICTED: transcription factor kayak [Drosophila miranda]
>gi|1036703565|ref|XP_017157020.1| PREDICTED: cuticle protein 21.3 isoform X2 [Drosophila miran
>gi|1036703548|ref|XP_017157013.1| PREDICTED: cuticle protein 16.5 isoform X1 [Drosophila miran
>gi|1036733150|ref|XP_017151479.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X1 [Dros
>gi|1036733169|ref|XP_017151487.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X1 [Dros
>gi|1036733204|ref|XP_017151500.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X3 [Dros
>gi|1036733221|ref|XP_017151508.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X4 [Dros
>gi|1036733188|ref|XP_017151494.1| PREDICTED: cytosolic purine 5'-nucleotidase isoform X2 [Dros
>gi|1036753777|ref|XP_017156518.1| PREDICTED: transferrin [Drosophila miranda]
>gi|1036739630|ref|XP_017155095.1| PREDICTED: coatomer subunit beta [Drosophila miranda]
>gi|1036736349|ref|XP_017153284.1| PREDICTED: nitric oxide synthase-interacting protein homolog
>gi|1036736333|ref|XP_017153272.1| PREDICTED: protein-serine O-palmitoleoyltransferase porcupin
>gi|1036731200|ref|XP_017150599.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731088|ref|XP_017150547.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731107|ref|XP_017150555.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731127|ref|XP_017150565.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731145|ref|XP_017150573.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731163|ref|XP_017150583.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731181|ref|XP_017150591.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731218|ref|XP_017150608.1| PREDICTED: cyclic AMP response element-binding protein B iso
>gi|1036731072|ref|XP_017150535.1| PREDICTED: allantoinase [Drosophila miranda]
>gi|1036731015|ref|XP_017150515.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036703264|ref|XP_017156851.1| PREDICTED: uncharacterized protein LOC108165325 [Drosophila
>gi|1036746880|ref|XP_017156135.1| PREDICTED: uncharacterized protein LOC108164749 isoform X2
>gi|1036746863|ref|XP_017156134.1| PREDICTED: uncharacterized protein LOC108164749 isoform X1
>gi|1036705738|ref|XP_017135798.1| PREDICTED: uncharacterized protein LOC108151612 [Drosophila
>gi|1036705969|ref|XP_017135969.1| PREDICTED: uncharacterized protein LOC108151706 [Drosophila
>gi|1036755886|ref|XP_017156638.1| PREDICTED: uncharacterized protein LOC108165097 isoform X1
>gi|1036755905|ref|XP_017156639.1| PREDICTED: uncharacterized protein LOC108165097 isoform X2
>gi|1036745905|ref|XP_017156073.1| PREDICTED: protein Exd1 homolog isoform X1 [Drosophila miran
>gi|1036745923|ref|XP_017156074.1| PREDICTED: protein Exd1 homolog isoform X2 [Drosophila miran
>gi|1036724408|ref|XP_017146537.1| PREDICTED: SWI5-dependent HO expression protein 3-like [Dros
>gi|1036701212|ref|XP_017154222.1| PREDICTED: uncharacterized protein LOC108163424 [Drosophila
>gi|1036701229|ref|XP_017154347.1| PREDICTED: TPR-containing protein DDB_G0280363 [Drosophila m
>gi|1036758205|ref|XP_017156763.1| PREDICTED: cytochrome c oxidase subunit 4 isoform 2, mitoch
>gi|1036816707|ref|XP_017137410.1| PREDICTED: protein yippee-like CG15309 [Drosophila miranda]
>gi|1036816726|ref|XP_017137411.1| PREDICTED: protein yippee-like CG15309 [Drosophila miranda]
>gi|1036816649|ref|XP_017137406.1| PREDICTED: kelch domain-containing protein 3 isoform X1 [Dro
>gi|1036816667|ref|XP_017137408.1| PREDICTED: kelch domain-containing protein 3 isoform X2 [Dro
>gi|1036816688|ref|XP_017137409.1| PREDICTED: kelch domain-containing protein 3 isoform X2 [Dro
>gi|1036816745|ref|XP_017137412.1| PREDICTED: INO80 complex subunit C [Drosophila miranda]
>gi|1036830533|ref|XP_017138182.1| PREDICTED: mannosyl-oligosaccharide glucosidase [Drosophila
>gi|1036799288|ref|XP_017136446.1| PREDICTED: transmembrane protein 68 [Drosophila miranda]
>gi|1036799253|ref|XP_017136444.1| PREDICTED: transport and Golgi organization protein 2 isofor
>gi|1036799271|ref|XP_017136445.1| PREDICTED: transport and Golgi organization protein 2 isofor
>gi|1036799234|ref|XP_017136443.1| PREDICTED: RRP12-like protein [Drosophila miranda]
>gi|1036784211|ref|XP_017135592.1| PREDICTED: uncharacterized protein LOC108151473 isoform X3
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>gi|1036784247|ref|XP_017135594.1| PREDICTED: uncharacterized protein LOC108151473 isoform X5
>gi|1036784192|ref|XP_017135591.1| PREDICTED: uncharacterized protein LOC108151473 isoform X2
>gi|1036784173|ref|XP_017135590.1| PREDICTED: uncharacterized protein LOC108151473 isoform X1
>gi|1036784230|ref|XP_017135593.1| PREDICTED: uncharacterized protein LOC108151473 isoform X4
>gi|1036832323|ref|XP_017138279.1| PREDICTED: DNA ligase 4 [Drosophila miranda]
>gi|1036838628|ref|XP_017138644.1| PREDICTED: ribonuclease H2 subunit B [Drosophila miranda]
>gi|1036832475|ref|XP_017138289.1| PREDICTED: uncharacterized protein LOC108153041 isoform X1
>gi|1036832495|ref|XP_017138290.1| PREDICTED: uncharacterized protein LOC108153041 isoform X2
>gi|1036827231|ref|XP_017137990.1| PREDICTED: fatty acid hydroxylase domain-containing protein
>gi|1036827212|ref|XP_017137989.1| PREDICTED: RNA-binding protein FUS [Drosophila miranda]
>gi|1036827248|ref|XP_017137991.1| PREDICTED: selenoprotein BthD [Drosophila miranda]
>gi|1036795987|ref|XP_017136259.1| PREDICTED: actin cytoskeleton-regulatory complex protein pa
>gi|1036796006|ref|XP_017136260.1| PREDICTED: actin cytoskeleton-regulatory complex protein pa
>gi|1036795931|ref|XP_017136257.1| PREDICTED: neuroglian isoform X1 [Drosophila miranda]
>gi|1036795968|ref|XP_017136258.1| PREDICTED: neuroglian isoform X2 [Drosophila miranda]
>gi|1036826162|ref|XP_017137928.1| PREDICTED: integrator complex subunit 4 [Drosophila miranda]
>gi|1036775700|ref|XP_017135090.1| PREDICTED: uncharacterized protein LOC108151171 isoform X2
>gi|1036775683|ref|XP_017135089.1| PREDICTED: uncharacterized protein LOC108151171 isoform X1
>gi|1036775719|ref|XP_017135091.1| PREDICTED: uncharacterized protein LOC108151171 isoform X3
>gi|1036775740|ref|XP_017135092.1| PREDICTED: uncharacterized protein LOC108151171 isoform X4
>gi|1036775664|ref|XP_017135088.1| PREDICTED: LOW QUALITY PROTEIN: E3 ubiquitin-protein ligase
>gi|1036758220|ref|XP_017156764.1| PREDICTED: uncharacterized protein LOC108165236 [Drosophila
>gi|1036832871|ref|XP_017138312.1| PREDICTED: uncharacterized protein LOC108153050 [Drosophila
>gi|1036832836|ref|XP_017138310.1| PREDICTED: uncharacterized protein LOC108153049 [Drosophila
>gi|1036832854|ref|XP_017138311.1| PREDICTED: uncharacterized protein LOC108153049 [Drosophila
>gi|1036832817|ref|XP_017138309.1| PREDICTED: zinc finger C4H2 domain-containing protein [Droso
>gi|1036812587|ref|XP_017137180.1| PREDICTED: dnaJ homolog subfamily C member 22 [Drosophila mi
>gi|1036812606|ref|XP_017137181.1| PREDICTED: dnaJ homolog subfamily C member 22 [Drosophila mi
>gi|1036812549|ref|XP_017137178.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036812568|ref|XP_017137179.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036812625|ref|XP_017137183.1| PREDICTED: homeotic protein female sterile-like [Drosophila
>gi|1036812644|ref|XP_017137184.1| PREDICTED: homeotic protein female sterile-like [Drosophila
>gi|1036812662|ref|XP_017137185.1| PREDICTED: uncharacterized protein LOC108152395 isoform X1
>gi|1036812683|ref|XP_017137186.1| PREDICTED: uncharacterized protein LOC108152395 isoform X2
>gi|1036838429|ref|XP_017138629.1| PREDICTED: uncharacterized protein LOC108153271 [Drosophila
>gi|1036789975|ref|XP_017135921.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC9 [Dro
>gi|1036789919|ref|XP_017135918.1| PREDICTED: density-regulated protein homolog [Drosophila mi
>gi|1036789883|ref|XP_017135916.1| PREDICTED: nitrogen permease regulator 2-like protein [Droso
>gi|1036789901|ref|XP_017135917.1| PREDICTED: ketimine reductase mu-crystallin [Drosophila mira
>gi|1036789938|ref|XP_017135919.1| PREDICTED: uncharacterized protein LOC108151686 isoform X1
>gi|1036789956|ref|XP_017135920.1| PREDICTED: uncharacterized protein LOC108151686 isoform X2
>gi|1036789831|ref|XP_017135913.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036789850|ref|XP_017135914.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036789867|ref|XP_017135915.1| PREDICTED: regulator of G-protein signaling 7 [Drosophila mi
>gi|1036839647|ref|XP_017138697.1| PREDICTED: uncharacterized protein LOC108153304 isoform X1
>gi|1036839662|ref|XP_017138698.1| PREDICTED: uncharacterized protein LOC108153304 isoform X2
>gi|1036839680|ref|XP_017138699.1| PREDICTED: uncharacterized protein LOC108153304 isoform X2
>gi|1036778909|ref|XP_017135282.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
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>gi|1036778776|ref|XP_017135273.1| PREDICTED: uncharacterized protein LOC108151262 isoform X5
>gi|1036778758|ref|XP_017135272.1| PREDICTED: uncharacterized protein LOC108151262 isoform X4
>gi|1036778704|ref|XP_017135269.1| PREDICTED: uncharacterized protein LOC108151262 isoform X1
>gi|1036778740|ref|XP_017135271.1| PREDICTED: uncharacterized protein LOC108151262 isoform X3
>gi|1036778722|ref|XP_017135270.1| PREDICTED: uncharacterized protein LOC108151262 isoform X2
>gi|1036778855|ref|XP_017135278.1| PREDICTED: LOW QUALITY PROTEIN: proteasome inhibitor PI31 su
>gi|1036778796|ref|XP_017135274.1| PREDICTED: uncharacterized protein LOC108151264 [Drosophila
>gi|1036778815|ref|XP_017135275.1| PREDICTED: uncharacterized protein LOC108151265 [Drosophila
>gi|1036778834|ref|XP_017135277.1| PREDICTED: uncharacterized protein LOC108151266 [Drosophila
>gi|1036778874|ref|XP_017135279.1| PREDICTED: 40S ribosomal protein S19a-like [Drosophila miran
>gi|1036778892|ref|XP_017135280.1| PREDICTED: uncharacterized protein LOC108151269 [Drosophila
>gi|1036767279|ref|XP_017134623.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036851147|ref|XP_017139330.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036767393|ref|XP_017134627.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036845649|ref|XP_017139012.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036845667|ref|XP_017139014.1| PREDICTED: 60 kDa heat shock protein, mitochondrial-like [Di
>gi|1036758238|ref|XP_017156765.1| PREDICTED: 60 kDa heat shock protein, mitochondrial-like [Di
>gi|1036845632|ref|XP_017139011.1| PREDICTED: calcitonin gene-related peptide type 1 receptor
>gi|1036843301|ref|XP_017138895.1| PREDICTED: ELAV-like protein 2 isoform X2 [Drosophila miran
>gi|1036843283|ref|XP_017138894.1| PREDICTED: ELAV-like protein 1 isoform X1 [Drosophila miran
>gi|1036838410|ref|XP_017138627.1| PREDICTED: protein YIPF1 [Drosophila miranda]
>gi|1036835152|ref|XP_017138433.1| PREDICTED: breast cancer metastasis-suppressor 1 homolog [Di
>gi|1036835186|ref|XP_017138435.1| PREDICTED: probable 39S ribosomal protein L49, mitochondrial
>gi|1036835167|ref|XP_017138434.1| PREDICTED: probable 39S ribosomal protein L49, mitochondrial
>gi|1036849963|ref|XP_017139262.1| PREDICTED: uncharacterized protein LOC108153662 isoform X1
>gi|1036849983|ref|XP_017139263.1| PREDICTED: uncharacterized protein LOC108153662 isoform X2
>gi|1036827395|ref|XP_017138001.1| PREDICTED: FAD synthase [Drosophila miranda]
>gi|1036827413|ref|XP_017138002.1| PREDICTED: FAD synthase [Drosophila miranda]
>gi|1036827378|ref|XP_017137999.1| PREDICTED: aarF domain-containing protein kinase 4 [Drosophi
>gi|1036811059|ref|XP_017137099.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036811019|ref|XP_017137097.1| PREDICTED: uncharacterized protein LOC108152344 [Drosophila
>gi|1036811038|ref|XP_017137098.1| PREDICTED: neprilysin-11 [Drosophila miranda]
>gi|1036839828|ref|XP_017138707.1| PREDICTED: uncharacterized protein LOC108153308 [Drosophila
>gi|1036839846|ref|XP_017138708.1| PREDICTED: uncharacterized protein LOC108153308 [Drosophila
>gi|1036842877|ref|XP_017138871.1| PREDICTED: polyhomeotic-proximal chromatin protein-like [Dro
>gi|1036758253|ref|XP_017156766.1| PREDICTED: bromodomain-containing factor 1-like [Drosophila
>gi|1036790512|ref|XP_017135950.1| PREDICTED: LOW QUALITY PROTEIN: polyhomeotic-proximal chroma
>gi|1036790419|ref|XP_017135945.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790435|ref|XP_017135946.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790455|ref|XP_017135947.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790474|ref|XP_017135948.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036790493|ref|XP_017135949.1| PREDICTED: D-2-hydroxyglutarate dehydrogenase, mitochondrial
>gi|1036812222|ref|XP_017137160.1| PREDICTED: 6-phosphogluconate dehydrogenase, decarboxylating
>gi|1036812297|ref|XP_017137164.1| PREDICTED: protein bcn92 [Drosophila miranda]
>gi|1036812239|ref|XP_017137161.1| PREDICTED: MAP kinase-activated protein kinase 2 [Drosophila
>gi|1036812260|ref|XP_017137162.1| PREDICTED: MAP kinase-activated protein kinase 2 [Drosophila
>gi|1036812279|ref|XP_017137163.1| PREDICTED: MAP kinase-activated protein kinase 2 [Drosophila
>gi|1036800012|ref|XP_017136486.1| PREDICTED: N-acylneuraminate-9-phosphatase [Drosophila miran
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>gi|1036800050|ref|XP_017136488.1| PREDICTED: protein lin-52 homolog [Drosophila miranda]
>gi|1036800032|ref|XP_017136487.1| PREDICTED: alpha-protein kinase 1 [Drosophila miranda]
>gi|1036799994|ref|XP_017136485.1| PREDICTED: regulator of telomere elongation helicase 1 homol
>gi|1036805859|ref|XP_017136816.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805878|ref|XP_017136818.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805897|ref|XP_017136819.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805916|ref|XP_017136820.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805934|ref|XP_017136821.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036805951|ref|XP_017136822.1| PREDICTED: integrator complex subunit 6 [Drosophila miranda]
>gi|1036784600|ref|XP_017135614.1| PREDICTED: sorting nexin-27 isoform X1 [Drosophila miranda]
>gi|1036784638|ref|XP_017135616.1| PREDICTED: uncharacterized protein LOC108151488 [Drosophila
>gi|1036784657|ref|XP_017135618.1| PREDICTED: 60S ribosomal protein L28 [Drosophila miranda]
>gi|1036784621|ref|XP_017135615.1| PREDICTED: sorting nexin-27 isoform X2 [Drosophila miranda]
>gi|1036784485|ref|XP_017135608.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784506|ref|XP_017135609.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784525|ref|XP_017135610.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784544|ref|XP_017135611.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784563|ref|XP_017135612.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 is
>gi|1036784449|ref|XP_017135606.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784412|ref|XP_017135604.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784430|ref|XP_017135605.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784466|ref|XP_017135607.1| PREDICTED: A disintegrin and metalloproteinase with thrombos
>gi|1036784582|ref|XP_017135613.1| PREDICTED: NADH-ubiquinone oxidoreductase 75 kDa subunit, m
>gi|1036758272|ref|XP_017156767.1| PREDICTED: protein trapped in endoderm-1-like, partial [Dros
>gi|1036819313|ref|XP_017137554.1| PREDICTED: thioredoxin reductase 1, mitochondrial isoform X2
>gi|1036819294|ref|XP_017137553.1| PREDICTED: thioredoxin reductase 1, mitochondrial isoform X1
>gi|1036819332|ref|XP_017137555.1| PREDICTED: C-factor isoform X1 [Drosophila miranda]
>gi|1036819351|ref|XP_017137556.1| PREDICTED: C-factor isoform X2 [Drosophila miranda]
>gi|1036819374|ref|XP_017137557.1| PREDICTED: uncharacterized protein LOC108152606 [Drosophila
>gi|1036837757|ref|XP_017138589.1| PREDICTED: troponin I isoform X1 [Drosophila miranda]
>gi|1036837813|ref|XP_017138592.1| PREDICTED: troponin I isoform X4 [Drosophila miranda]
>gi|1036837795|ref|XP_017138591.1| PREDICTED: troponin I isoform X3 [Drosophila miranda]
>gi|1036837776|ref|XP_017138590.1| PREDICTED: troponin I isoform X2 [Drosophila miranda]
>gi|1036837868|ref|XP_017138595.1| PREDICTED: troponin I isoform X7 [Drosophila miranda]
>gi|1036837922|ref|XP_017138599.1| PREDICTED: troponin I isoform X10 [Drosophila miranda]
>gi|1036837903|ref|XP_017138598.1| PREDICTED: troponin I isoform X9 [Drosophila miranda]
>gi|1036837885|ref|XP_017138597.1| PREDICTED: troponin I isoform X8 [Drosophila miranda]
>gi|1036837831|ref|XP_017138593.1| PREDICTED: troponin I isoform X5 [Drosophila miranda]
>gi|1036837850|ref|XP_017138594.1| PREDICTED: troponin I isoform X6 [Drosophila miranda]
>gi|1036837941|ref|XP_017138600.1| PREDICTED: troponin I isoform X11 [Drosophila miranda]
>gi|1036837960|ref|XP_017138601.1| PREDICTED: troponin I isoform X12 [Drosophila miranda]
>gi|1036848353|ref|XP_017139165.1| PREDICTED: casein kinase II subunit beta'-like [Drosophila m
>gi|1036808771|ref|XP_017136975.1| PREDICTED: interference hedgehog isoform X2 [Drosophila mira
>gi|1036808655|ref|XP_017136969.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808674|ref|XP_017136970.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808694|ref|XP_017136971.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808713|ref|XP_017136972.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808732|ref|XP_017136973.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
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>gi|1036808753|ref|XP_017136974.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808788|ref|XP_017136976.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808806|ref|XP_017136977.1| PREDICTED: interference hedgehog isoform X1 [Drosophila mira
>gi|1036808827|ref|XP_017136978.1| PREDICTED: regulatory protein zeste [Drosophila miranda]
>gi|1036803920|ref|XP_017136705.1| PREDICTED: basement membrane-specific heparan sulfate protec
>gi|1036803937|ref|XP_017136706.1| PREDICTED: basement membrane-specific heparan sulfate protec
>gi|1036846112|ref|XP_017139040.1| PREDICTED: PDF receptor [Drosophila miranda]
>gi|1036850930|ref|XP_017139318.1| PREDICTED: probable V-type proton ATPase subunit D 2 [Drosop
>gi|1036828261|ref|XP_017138050.1| PREDICTED: dopamine N-acetyltransferase isoform X1 [Drosophi
>gi|1036828280|ref|XP_017138052.1| PREDICTED: dopamine N-acetyltransferase isoform X2 [Drosophi
>gi|1036828299|ref|XP_017138053.1| PREDICTED: dopamine N-acetyltransferase isoform X2 [Drosophi
>gi|1036828338|ref|XP_017138055.1| PREDICTED: protein GUCD1 isoform X2 [Drosophila miranda]
>gi|1036828319|ref|XP_017138054.1| PREDICTED: protein GUCD1 isoform X1 [Drosophila miranda]
>gi|1036828243|ref|XP_017138049.1| PREDICTED: zinc finger and BTB domain-containing protein 49
>gi|1036833065|ref|XP_017138323.1| PREDICTED: PHD finger protein 12 [Drosophila miranda]
>gi|1036815826|ref|XP_017137357.1| PREDICTED: uncharacterized protein LOC108152492 [Drosophila
>gi|1036815807|ref|XP_017137356.1| PREDICTED: uncharacterized protein LOC108152491 [Drosophila
>gi|1036815784|ref|XP_017137354.1| PREDICTED: mitochondrial ribonuclease P protein 3 [Drosophi
>gi|1036815763|ref|XP_017137353.1| PREDICTED: DNA repair endonuclease XPF [Drosophila miranda]
>gi|1036851487|ref|XP_017139351.1| PREDICTED: uncharacterized protein LOC108153733 [Drosophila
>gi|1036767666|ref|XP_017134640.1| PREDICTED: uncharacterized protein LOC108150845 [Drosophila
>gi|1036758291|ref|XP_017156768.1| PREDICTED: uncharacterized protein LOC108165242 [Drosophila
>gi|1036765327|ref|XP_017157164.1| PREDICTED: uncharacterized protein LOC108165601 [Drosophila
>gi|1036769627|ref|XP_017134750.1| PREDICTED: uncharacterized protein LOC108150931 [Drosophila
>gi|1036766542|ref|XP_017134580.1| PREDICTED: uncharacterized protein LOC108150789 isoform X2
>gi|1036766523|ref|XP_017134579.1| PREDICTED: uncharacterized protein LOC108150789 isoform X1
>gi|1036852340|ref|XP_017139405.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036763406|ref|XP_017157064.1| PREDICTED: translation initiation factor IF-2 [Drosophila m
>gi|1036851345|ref|XP_017139342.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1036766958|ref|XP_017134604.1| PREDICTED: uncharacterized protein LOC108150810 [Drosophila
>gi|1036797845|ref|XP_017136363.1| PREDICTED: fasciclin-2 isoform X1 [Drosophila miranda]
>gi|1036797879|ref|XP_017136365.1| PREDICTED: fasciclin-2 isoform X3 [Drosophila miranda]
>gi|1036797862|ref|XP_017136364.1| PREDICTED: fasciclin-2 isoform X2 [Drosophila miranda]
>gi|1036797898|ref|XP_017136366.1| PREDICTED: fasciclin-2 isoform X4 [Drosophila miranda]
>gi|1036797917|ref|XP_017136367.1| PREDICTED: fasciclin-2 isoform X5 [Drosophila miranda]
>gi|1036797936|ref|XP_017136369.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucuro
>gi|1036846146|ref|XP_017139041.1| PREDICTED: translation machinery-associated protein 7 homolo
>gi|1036827266|ref|XP_017137993.1| PREDICTED: mitochondrial tRNA-specific 2-thiouridylase 1 [Dr
>gi|1036827284|ref|XP_017137994.1| PREDICTED: monocyte to macrophage differentiation factor [Dr
>gi|1036827303|ref|XP_017137995.1| PREDICTED: monocyte to macrophage differentiation factor [Dr
>gi|1036801742|ref|XP_017136587.1| PREDICTED: HEAT repeat-containing protein 6 [Drosophila mira
>gi|1036801760|ref|XP_017136588.1| PREDICTED: retinoblastoma family protein [Drosophila miranda
>gi|1036852545|ref|XP_017139417.1| PREDICTED: uncharacterized protein LOC108153784 [Drosophila
>gi|1036758307|ref|XP_017156769.1| PREDICTED: ESX-1 secretion-associated protein EspI [Drosophi
>gi|1036791737|ref|XP_017136020.1| PREDICTED: vesicle-trafficking protein SEC22b-B [Drosophila
>gi|1036791635|ref|XP_017136013.1| PREDICTED: uncharacterized protein CG43867 isoform X7 [Droso
>gi|1036791652|ref|XP_017136014.1| PREDICTED: uncharacterized protein CG43867 isoform X8 [Droso
>gi|1036791616|ref|XP_017136012.1| PREDICTED: uncharacterized protein CG43867 isoform X6 [Droso
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>gi|1036791579|ref|XP_017136010.1| PREDICTED: uncharacterized protein CG43867 isoform X4 [Dros
>gi|1036791539|ref|XP_017136008.1| PREDICTED: uncharacterized protein CG43867 isoform X2 [Dros
>gi|1036791558|ref|XP_017136009.1| PREDICTED: uncharacterized protein CG43867 isoform X3 [Dros
>gi|1036791520|ref|XP_017136007.1| PREDICTED: uncharacterized protein CG43867 isoform X1 [Dros
>gi|1036791598|ref|XP_017136011.1| PREDICTED: uncharacterized protein CG43867 isoform X5 [Dros
>gi|1036791671|ref|XP_017136015.1| PREDICTED: uncharacterized protein CG43867 isoform X9 [Dros
>gi|1036791703|ref|XP_017136016.1| PREDICTED: uncharacterized protein CG43867 isoform X10 [Dros
>gi|1036791720|ref|XP_017136017.1| PREDICTED: uncharacterized protein CG43867 isoform X11 [Dros
>gi|1036791774|ref|XP_017136022.1| PREDICTED: uncharacterized protein LOC108151733 [Drosophila
>gi|1036791755|ref|XP_017136021.1| PREDICTED: uncharacterized protein LOC108151732 [Drosophila
>gi|1036846346|ref|XP_017139052.1| PREDICTED: protein lunapark-B-like isoform X2 [Drosophila m
>gi|1036846363|ref|XP_017139053.1| PREDICTED: protein lunapark-B-like isoform X3 [Drosophila m
>gi|1036846327|ref|XP_017139051.1| PREDICTED: protein lunapark-B-like isoform X1 [Drosophila m
>gi|1036827152|ref|XP_017137986.1| PREDICTED: leucine-zipper-like transcriptional regulator 1
>gi|1036843707|ref|XP_017138920.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036843688|ref|XP_017138919.1| PREDICTED: nucleosome assembly protein 1-like 1-A [Drosophil
>gi|1036824926|ref|XP_017137860.1| PREDICTED: uncharacterized protein LOC108152787 [Drosophila
>gi|1036824944|ref|XP_017137861.1| PREDICTED: cell wall integrity and stress response component
>gi|1036824963|ref|XP_017137862.1| PREDICTED: translocating chain-associated membrane protein
>gi|1036842197|ref|XP_017138834.1| PREDICTED: crossover junction endonuclease MUS81 [Drosophila
>gi|1036842181|ref|XP_017138833.1| PREDICTED: LOW QUALITY PROTEIN: RUN domain-containing prote
>gi|1036846681|ref|XP_017139072.1| PREDICTED: 3-oxoacyl-[acyl-carrier-protein] reductase FabG
>gi|1036846658|ref|XP_017139071.1| PREDICTED: LOW QUALITY PROTEIN: synaptic vesicle glycoprote
>gi|1036758325|ref|XP_017156770.1| PREDICTED: NADH-quinone oxidoreductase subunit F 1-like [Dr
>gi|1036758343|ref|XP_017156771.1| PREDICTED: NADH-quinone oxidoreductase subunit G-like [Dros
>gi|1036846381|ref|XP_017139054.1| PREDICTED: uncharacterized protein LOC108153509 [Drosophila
>gi|1036833321|ref|XP_017138338.1| PREDICTED: 60S ribosomal protein L13a-like [Drosophila mira
>gi|1036758359|ref|XP_017156774.1| PREDICTED: ERI1 exoribonuclease 2-like [Drosophila miranda]
>gi|1036765347|ref|XP_017157165.1| PREDICTED: uncharacterized protein DDB_G0283357 [Drosophila
>gi|1036821039|ref|XP_017137644.1| PREDICTED: probable calcium-binding protein CML12 [Drosophil
>gi|1036821057|ref|XP_017137645.1| PREDICTED: probable calcium-binding protein CML12 [Drosophil
>gi|1036821000|ref|XP_017137641.1| PREDICTED: adenylate cyclase, terminal-differentiation speci
>gi|1036821113|ref|XP_017137647.1| PREDICTED: uncharacterized protein LOC108152662 [Drosophila
>gi|1036820981|ref|XP_017137640.1| PREDICTED: cell division control protein 45 homolog [Drosoph
>gi|1036821094|ref|XP_017137646.1| PREDICTED: ribonuclease P protein subunit p21 [Drosophila m
>gi|1036821018|ref|XP_017137643.1| PREDICTED: gamma-butyrobetaine dioxygenase [Drosophila mira
>gi|1036824082|ref|XP_017137814.1| PREDICTED: protein suppressor of white apricot isoform X1 [
>gi|1036824101|ref|XP_017137815.1| PREDICTED: protein suppressor of white apricot isoform X2 [
>gi|1036824136|ref|XP_017137817.1| PREDICTED: protein suppressor of white apricot isoform X3 [
>gi|1036824155|ref|XP_017137818.1| PREDICTED: protein suppressor of white apricot isoform X4 [
>gi|1036767334|ref|XP_017134624.1| PREDICTED: chymotrypsin-elastase inhibitor ixodidin [Drosop
>gi|1036848879|ref|XP_017139196.1| PREDICTED: 27 kDa hemolymph protein [Drosophila miranda]
>gi|1036765926|ref|XP_017134548.1| PREDICTED: uncharacterized protein LOC108150765 [Drosophila
>gi|1036852071|ref|XP_017139388.1| PREDICTED: succinyl-CoA ligase subunit alpha, mitochondrial
>gi|1036812434|ref|XP_017137172.1| PREDICTED: uncharacterized protein LOC108152386 [Drosophila
>gi|1036812491|ref|XP_017137175.1| PREDICTED: uncharacterized protein LOC108152388 [Drosophila
>gi|1036812452|ref|XP_017137173.1| PREDICTED: uncharacterized protein LOC108152387 isoform X1
>gi|1036812470|ref|XP_017137174.1| PREDICTED: uncharacterized protein LOC108152387 isoform X2
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>gi|1036765745|ref|XP_017134537.1| PREDICTED: uncharacterized protein LOC108150754 [Drosophila
>gi|1036758375|ref|XP_017156775.1| PREDICTED: pollen-specific leucine-rich repeat extensin-like
>gi|1036764635|ref|XP_017157127.1| PREDICTED: uncharacterized protein LOC108165564 [Drosophila
>gi|1036849174|ref|XP_017139213.1| PREDICTED: 27 kDa hemolymph protein [Drosophila miranda]
>gi|1036765996|ref|XP_017134551.1| PREDICTED: uncharacterized protein LOC108150768 [Drosophila
>gi|1036765975|ref|XP_017134550.1| PREDICTED: uncharacterized protein LOC108150767 [Drosophila
>gi|1036770401|ref|XP_017134793.1| PREDICTED: probable basic-leucine zipper transcription facto
>gi|1036758394|ref|XP_017156776.1| PREDICTED: uncharacterized protein LOC108165249 [Drosophila
>gi|1036852162|ref|XP_017139394.1| PREDICTED: alpha-protein kinase 1 [Drosophila miranda]
>gi|1036758413|ref|XP_017156777.1| PREDICTED: uncharacterized protein LOC108165250 [Drosophila
>gi|1036852090|ref|XP_017139390.1| PREDICTED: GATA zinc finger domain-containing protein 10 [D
>gi|1036758432|ref|XP_017156778.1| PREDICTED: uncharacterized protein LOC108165251 [Drosophila
>gi|1036764243|ref|XP_017157110.1| PREDICTED: hornerin [Drosophila miranda]
>gi|1036846417|ref|XP_017139056.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1036801064|ref|XP_017136546.1| PREDICTED: LOW QUALITY PROTEIN: polyhomeotic-proximal chroma
>gi|1036835060|ref|XP_017138428.1| PREDICTED: probable ATP-dependent DNA helicase DDX11 [Drosop
>gi|1036832108|ref|XP_017138270.1| PREDICTED: activating transcription factor 3 [Drosophila mi
>gi|1036764738|ref|XP_017157133.1| PREDICTED: calcineurin B homologous protein 1 [Drosophila m
>gi|1036758451|ref|XP_017156779.1| PREDICTED: uncharacterized protein LOC108165252 [Drosophila
>gi|1036850799|ref|XP_017139310.1| PREDICTED: uncharacterized protein LOC108153696 [Drosophila
>gi|1036796526|ref|XP_017136288.1| PREDICTED: disheveled-associated activator of morphogenesis
>gi|1036796507|ref|XP_017136287.1| PREDICTED: disheveled-associated activator of morphogenesis
>gi|1036796545|ref|XP_017136289.1| PREDICTED: golgin subfamily A member 6C-like [Drosophila mi
>gi|1036849821|ref|XP_017139253.1| PREDICTED: cell wall protein DAN4 [Drosophila miranda]
>gi|1036831922|ref|XP_017138259.1| PREDICTED: N-alpha-acetyltransferase 30 isoform X1 [Drosoph
>gi|1036831941|ref|XP_017138260.1| PREDICTED: N-alpha-acetyltransferase 30 isoform X2 [Drosoph
>gi|1036837347|ref|XP_017138563.1| PREDICTED: poly(A) RNA polymerase, mitochondrial [Drosophila
>gi|1036837366|ref|XP_017138565.1| PREDICTED: CD9 antigen [Drosophila miranda]
>gi|1036807353|ref|XP_017136897.1| PREDICTED: solute carrier family 12 member 8 [Drosophila mi
>gi|1036807335|ref|XP_017136896.1| PREDICTED: ESF1 homolog [Drosophila miranda]
>gi|1036807389|ref|XP_017136898.1| PREDICTED: serine/threonine-protein kinase Nek8 [Drosophila
>gi|1036767411|ref|XP_017134628.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036811422|ref|XP_017137116.1| PREDICTED: sex-lethal homolog isoform X2 [Drosophila miranda
>gi|1036811403|ref|XP_017137115.1| PREDICTED: sex-lethal homolog isoform X1 [Drosophila miranda
>gi|1036811441|ref|XP_017137117.1| PREDICTED: RNA-binding protein FUS-like [Drosophila miranda]
>gi|1036801929|ref|XP_017136598.1| PREDICTED: 5'-AMP-activated protein kinase catalytic subunit
>gi|1036801949|ref|XP_017136599.1| PREDICTED: thioredoxin, mitochondrial [Drosophila miranda]
>gi|1036801781|ref|XP_017136589.1| PREDICTED: rho GTPase-activating protein gacZ isoform X1 [D
>gi|1036801800|ref|XP_017136590.1| PREDICTED: rho GTPase-activating protein gacZ isoform X2 [D
>gi|1036801817|ref|XP_017136591.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [D
>gi|1036801836|ref|XP_017136592.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [D
>gi|1036801855|ref|XP_017136593.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [D
>gi|1036801874|ref|XP_017136594.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [D
>gi|1036801893|ref|XP_017136595.1| PREDICTED: rho GTPase-activating protein gacZ isoform X3 [D
>gi|1036801911|ref|XP_017136597.1| PREDICTED: rho GTPase-activating protein gacZ isoform X4 [D
>gi|1036830077|ref|XP_017138153.1| PREDICTED: RILP-like protein homolog isoform X2 [Drosophila
>gi|1036830058|ref|XP_017138152.1| PREDICTED: RILP-like protein homolog isoform X1 [Drosophila
>gi|1036825259|ref|XP_017137880.1| PREDICTED: LOW QUALITY PROTEIN: microtubule-associated prote
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>gi|1036758470|ref|XP_017156780.1| PREDICTED: putative gustatory receptor 2a [Drosophila miran
>gi|1036849335|ref|XP_017139222.1| PREDICTED: uncharacterized protein LOC108153629 [Drosophila
>gi|1036849349|ref|XP_017139223.1| PREDICTED: uncharacterized protein LOC108153630 [Drosophila
>gi|1036826589|ref|XP_017137953.1| PREDICTED: leucine-rich PPR motif-containing protein, mitoch
>gi|1036834104|ref|XP_017138374.1| PREDICTED: enoyl-CoA delta isomerase 2, mitochondrial [Droso
>gi|1036834085|ref|XP_017138373.1| PREDICTED: large subunit GTPase 1 homolog [Drosophila miran
>gi|1036783797|ref|XP_017135569.1| PREDICTED: GDP-fucose protein O-fucosyltransferase 2 [Drosop
>gi|1036784078|ref|XP_017135585.1| PREDICTED: PXMP2/4 family protein 4 [Drosophila miranda]
>gi|1036784097|ref|XP_017135586.1| PREDICTED: PXMP2/4 family protein 4 [Drosophila miranda]
>gi|1036783816|ref|XP_017135570.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036784021|ref|XP_017135582.1| PREDICTED: protein sym1 isoform X1 [Drosophila miranda]
>gi|1036784040|ref|XP_017135583.1| PREDICTED: protein sym1 isoform X1 [Drosophila miranda]
>gi|1036784059|ref|XP_017135584.1| PREDICTED: protein sym1 isoform X2 [Drosophila miranda]
>gi|1036784002|ref|XP_017135581.1| PREDICTED: uncharacterized protein LOC108151467 [Drosophila
>gi|1036783889|ref|XP_017135574.1| PREDICTED: trypsin eta [Drosophila miranda]
>gi|1036783927|ref|XP_017135577.1| PREDICTED: ras-related protein Rab-27A isoform X1 [Drosophil
>gi|1036783946|ref|XP_017135578.1| PREDICTED: ras-related protein Rab-27A isoform X2 [Drosophil
>gi|1036783965|ref|XP_017135579.1| PREDICTED: ras-related protein Rab-27A isoform X2 [Drosophil
>gi|1036783983|ref|XP_017135580.1| PREDICTED: ras-related protein Rab-27A isoform X3 [Drosophil
>gi|1036783834|ref|XP_017135571.1| PREDICTED: uncharacterized protein LOC108151462 [Drosophila
>gi|1036783853|ref|XP_017135572.1| PREDICTED: pleckstrin homology domain-containing family F me
>gi|1036783872|ref|XP_017135573.1| PREDICTED: pleckstrin homology domain-containing family F me
>gi|1036783908|ref|XP_017135576.1| PREDICTED: 40S ribosomal protein SA [Drosophila miranda]
>gi|1036849305|ref|XP_017139220.1| PREDICTED: glutamate receptor ionotropic, NMDA 2B isoform X
>gi|1036849270|ref|XP_017139218.1| PREDICTED: glutamate receptor ionotropic, NMDA 2B isoform X
>gi|1036849289|ref|XP_017139219.1| PREDICTED: glutamate receptor ionotropic, NMDA 2B isoform X
>gi|1036764581|ref|XP_017157124.1| PREDICTED: uncharacterized protein LOC108165560 [Drosophila
>gi|1036820681|ref|XP_017137623.1| PREDICTED: BTB/POZ domain-containing protein KCTD5 [Drosoph
>gi|1036758489|ref|XP_017156781.1| PREDICTED: LOW QUALITY PROTEIN: mucin-5AC [Drosophila miran
>gi|1036770545|ref|XP_017134801.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770578|ref|XP_017134803.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770512|ref|XP_017134799.1| PREDICTED: broad-complex core protein isoform X1 [Drosophila
>gi|1036770529|ref|XP_017134800.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770561|ref|XP_017134802.1| PREDICTED: broad-complex core protein isoforms 1/2/3/4/5 iso
>gi|1036770494|ref|XP_017134798.1| PREDICTED: vacuolar protein sorting-associated protein 18 ho
>gi|1036770656|ref|XP_017134808.1| PREDICTED: protein C19orf12 homolog [Drosophila miranda]
>gi|1036770597|ref|XP_017134804.1| PREDICTED: protein halfway isoform X1 [Drosophila miranda]
>gi|1036770616|ref|XP_017134805.1| PREDICTED: protein halfway isoform X2 [Drosophila miranda]
>gi|1036770438|ref|XP_017134795.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770420|ref|XP_017134794.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770475|ref|XP_017134797.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770456|ref|XP_017134796.1| PREDICTED: coiled-coil domain-containing protein CG32809 iso
>gi|1036770676|ref|XP_017134809.1| PREDICTED: uncharacterized protein LOC108150978 [Drosophila
>gi|1036770637|ref|XP_017134807.1| PREDICTED: uncharacterized protein LOC108150976 [Drosophila
>gi|1036834752|ref|XP_017138409.1| PREDICTED: protein a6 [Drosophila miranda]
>gi|1036758508|ref|XP_017156782.1| PREDICTED: trypsin-3 [Drosophila miranda]
>gi|1036850967|ref|XP_017139319.1| PREDICTED: delta-sarcoglycan [Drosophila miranda]
>gi|1036766628|ref|XP_017134584.1| PREDICTED: uncharacterized protein LOC108150793 [Drosophila
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>gi|1036765469|ref|XP_017157173.1| PREDICTED: uncharacterized protein LOC108165607 [Drosophila
>gi|1036803038|ref|XP_017136656.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802982|ref|XP_017136653.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802961|ref|XP_017136652.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802942|ref|XP_017136651.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036802924|ref|XP_017136650.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036803000|ref|XP_017136654.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036803019|ref|XP_017136655.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036803059|ref|XP_017136657.1| PREDICTED: double-stranded RNA-specific editase Adar isofor
>gi|1036758526|ref|XP_017156783.1| PREDICTED: LOW QUALITY PROTEIN: TRAF3-interacting protein 1-
>gi|1036776046|ref|XP_017135108.1| PREDICTED: neurobeachin isoform X5 [Drosophila miranda]
>gi|1036776063|ref|XP_017135109.1| PREDICTED: neurobeachin isoform X6 [Drosophila miranda]
>gi|1036776084|ref|XP_017135110.1| PREDICTED: neurobeachin isoform X7 [Drosophila miranda]
>gi|1036776196|ref|XP_017135116.1| PREDICTED: neurobeachin isoform X13 [Drosophila miranda]
>gi|1036776141|ref|XP_017135113.1| PREDICTED: neurobeachin isoform X10 [Drosophila miranda]
>gi|1036776234|ref|XP_017135118.1| PREDICTED: neurobeachin isoform X15 [Drosophila miranda]
>gi|1036776308|ref|XP_017135122.1| PREDICTED: proteasomal ubiquitin receptor ADRM1 homolog [Dro
>gi|1036776327|ref|XP_017135123.1| PREDICTED: proteasome subunit alpha type-3-like [Drosophila
>gi|1036776346|ref|XP_017135124.1| PREDICTED: uncharacterized protein LOC108151189 [Drosophila
>gi|1036776270|ref|XP_017135120.1| PREDICTED: uncharacterized protein LOC108151185 [Drosophila
>gi|1036776289|ref|XP_017135121.1| PREDICTED: uncharacterized protein LOC108151185 [Drosophila
>gi|1036776251|ref|XP_017135119.1| PREDICTED: uncharacterized abhydrolase domain-containing pro
>gi|1036775966|ref|XP_017135104.1| PREDICTED: neurobeachin isoform X1 [Drosophila miranda]
>gi|1036776160|ref|XP_017135114.1| PREDICTED: neurobeachin isoform X11 [Drosophila miranda]
>gi|1036775985|ref|XP_017135105.1| PREDICTED: neurobeachin isoform X2 [Drosophila miranda]
>gi|1036776004|ref|XP_017135106.1| PREDICTED: neurobeachin isoform X3 [Drosophila miranda]
>gi|1036776023|ref|XP_017135107.1| PREDICTED: neurobeachin isoform X4 [Drosophila miranda]
>gi|1036776179|ref|XP_017135115.1| PREDICTED: neurobeachin isoform X12 [Drosophila miranda]
>gi|1036776103|ref|XP_017135111.1| PREDICTED: neurobeachin isoform X8 [Drosophila miranda]
>gi|1036776120|ref|XP_017135112.1| PREDICTED: neurobeachin isoform X9 [Drosophila miranda]
>gi|1036776215|ref|XP_017135117.1| PREDICTED: neurobeachin isoform X14 [Drosophila miranda]
>gi|1036776383|ref|XP_017135127.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1036776365|ref|XP_017135125.1| PREDICTED: 60S ribosomal protein L30-like [Drosophila miran
>gi|1036767718|ref|XP_017134642.1| PREDICTED: uncharacterized protein LOC108150849 [Drosophila
>gi|1036852578|ref|XP_017139419.1| PREDICTED: dnaJ homolog subfamily C member 7-like [Drosophil
>gi|1036764462|ref|XP_017157119.1| PREDICTED: uncharacterized protein LOC108165557 [Drosophila
>gi|1036773580|ref|XP_017134973.1| PREDICTED: uncharacterized protein LOC108151094 isoform X1
>gi|1036773597|ref|XP_017134974.1| PREDICTED: uncharacterized protein LOC108151094 isoform X2
>gi|1036773394|ref|XP_017134962.1| PREDICTED: uncharacterized protein LOC108151087 [Drosophila
>gi|1036773671|ref|XP_017134979.1| PREDICTED: uncharacterized protein LOC108151096 [Drosophila
>gi|1036773564|ref|XP_017134972.1| PREDICTED: cyclin-dependent kinase 7 [Drosophila miranda]
>gi|1036773690|ref|XP_017134980.1| PREDICTED: U1 small nuclear ribonucleoprotein A [Drosophila
>gi|1036773709|ref|XP_017134981.1| PREDICTED: thioredoxin-2 [Drosophila miranda]
>gi|1036773728|ref|XP_017134982.1| PREDICTED: thioredoxin-1 [Drosophila miranda]
>gi|1036773431|ref|XP_017134964.1| PREDICTED: uncharacterized protein LOC108151090 [Drosophila
>gi|1036773413|ref|XP_017134963.1| PREDICTED: uncharacterized protein LOC108151089 [Drosophila
>gi|1036773542|ref|XP_017134971.1| PREDICTED: something about silencing protein 10 [Drosophila
>gi|1036773375|ref|XP_017134961.1| PREDICTED: RNA-binding protein 4F [Drosophila miranda]
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>gi|1036773469|ref|XP_017134966.1| PREDICTED: acid phosphatase type 7 isoform X2 [Drosophila m:
>gi|1036773523|ref|XP_017134970.1| PREDICTED: acid phosphatase type 7 isoform X5 [Drosophila m:
>gi|1036773486|ref|XP_017134968.1| PREDICTED: acid phosphatase type 7 isoform X3 [Drosophila m:
>gi|1036773505|ref|XP_017134969.1| PREDICTED: acid phosphatase type 7 isoform X4 [Drosophila m:
>gi|1036773450|ref|XP_017134965.1| PREDICTED: acid phosphatase type 7 isoform X1 [Drosophila m:
>gi|1036773652|ref|XP_017134978.1| PREDICTED: shematrin-like protein 2 isoform X3 [Drosophila m
>gi|1036773634|ref|XP_017134977.1| PREDICTED: glycine-rich cell wall structural protein 2 isofo
>gi|1036773616|ref|XP_017134975.1| PREDICTED: glycine-rich RNA-binding protein 3, mitochondrial
>gi|1036824695|ref|XP_017137850.1| PREDICTED: RNA-binding protein 33 [Drosophila miranda]
>gi|1036763208|ref|XP_017157053.1| PREDICTED: uncharacterized protein LOC108165508 [Drosophila
>gi|1036790788|ref|XP_017135965.1| PREDICTED: zinc finger protein 12 [Drosophila miranda]
>gi|1036790846|ref|XP_017135968.1| PREDICTED: probable 6-phosphogluconolactonase [Drosophila m:
>gi|1036790807|ref|XP_017135966.1| PREDICTED: nuclear RNA export factor 1 [Drosophila miranda]
>gi|1036790825|ref|XP_017135967.1| PREDICTED: putative sodium-dependent multivitamin transporte
>gi|1036790917|ref|XP_017135972.1| PREDICTED: uncharacterized protein LOC108151711 [Drosophila
>gi|1036790934|ref|XP_017135973.1| PREDICTED: uncharacterized protein LOC108151712 [Drosophila
>gi|1036790863|ref|XP_017135970.1| PREDICTED: uncharacterized protein LOC108151710 isoform X1
>gi|1036790882|ref|XP_017135971.1| PREDICTED: uncharacterized protein LOC108151710 isoform X2
>gi|1036812105|ref|XP_017137153.1| PREDICTED: uncharacterized protein LOC108152374 [Drosophila
>gi|1036812026|ref|XP_017137149.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036812047|ref|XP_017137150.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036812066|ref|XP_017137151.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036812087|ref|XP_017137152.1| PREDICTED: 1-phosphatidylinositol 4,5-bisphosphate phosphodi
>gi|1036832144|ref|XP_017138273.1| PREDICTED: pyruvate dehydrogenase phosphatase regulatory sul
>gi|1036762241|ref|XP_017157000.1| PREDICTED: optomotor-blind protein [Drosophila miranda]
>gi|1036762365|ref|XP_017157008.1| PREDICTED: uncharacterized protein CG3556 [Drosophila mirand
>gi|1036762403|ref|XP_017157010.1| PREDICTED: protein TonB [Drosophila miranda]
>gi|1036762223|ref|XP_017156999.1| PREDICTED: AP-3 complex subunit beta-2 [Drosophila miranda]
>gi|1036762437|ref|XP_017157012.1| PREDICTED: protein SREK1IP1 [Drosophila miranda]
>gi|1036762384|ref|XP_017157009.1| PREDICTED: uncharacterized protein LOC108165469 [Drosophila
>gi|1036762418|ref|XP_017157011.1| PREDICTED: serine--pyruvate aminotransferase, mitochondrial
>gi|1036762455|ref|XP_017157014.1| PREDICTED: 60S ribosomal protein L7a [Drosophila miranda]
>gi|1036762259|ref|XP_017157002.1| PREDICTED: protein deltex [Drosophila miranda]
>gi|1036762014|ref|XP_017156992.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762026|ref|XP_017156993.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762131|ref|XP_017156994.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762148|ref|XP_017156995.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762166|ref|XP_017156996.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036761940|ref|XP_017156988.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036761959|ref|XP_017156989.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dro
>gi|1036761998|ref|XP_017156991.1| PREDICTED: uncharacterized protein LOC108165461 isoform X3
>gi|1036762204|ref|XP_017156998.1| PREDICTED: uncharacterized protein LOC108165461 isoform X10
>gi|1036762186|ref|XP_017156997.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036762560|ref|XP_017157019.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC:
>gi|1036761977|ref|XP_017156990.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dro
>gi|1036762493|ref|XP_017157016.1| PREDICTED: uncharacterized protein LOC108165476 [Drosophila
>gi|1036762511|ref|XP_017157017.1| PREDICTED: 40S ribosomal protein S12, mitochondrial [Drosoph
>gi|1036758542|ref|XP_017156784.1| PREDICTED: protein giant [Drosophila miranda]
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>gi|1036762524|ref|XP_017157018.1| PREDICTED: uncharacterized protein LOC108165478 [Drosophila
>gi|1036758559|ref|XP_017156785.1| PREDICTED: LOW QUALITY PROTEIN: protein regulator of cytoki
>gi|1036758578|ref|XP_017156786.1| PREDICTED: protein argonaute-2-like, partial [Drosophila mi
>gi|1036762474|ref|XP_017157015.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1036762347|ref|XP_017157007.1| PREDICTED: uncharacterized protein LOC108165467 isoform X4
>gi|1036762314|ref|XP_017157005.1| PREDICTED: uncharacterized protein LOC108165467 isoform X2
>gi|1036762295|ref|XP_017157004.1| PREDICTED: uncharacterized protein LOC108165467 isoform X1
>gi|1036762332|ref|XP_017157006.1| PREDICTED: uncharacterized protein LOC108165467 isoform X3
>gi|1036758596|ref|XP_017156787.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036762277|ref|XP_017157003.1| PREDICTED: zinc finger and BTB domain-containing protein 48
>gi|1036817445|ref|XP_017137449.1| PREDICTED: casein kinase I [Drosophila miranda]
>gi|1036811137|ref|XP_017137103.1| PREDICTED: syntaxin-binding protein 5 isoform X3 [Drosophila
>gi|1036811117|ref|XP_017137102.1| PREDICTED: syntaxin-binding protein 5 isoform X2 [Drosophila
>gi|1036811155|ref|XP_017137104.1| PREDICTED: syntaxin-binding protein 5 isoform X4 [Drosophila
>gi|1036811078|ref|XP_017137100.1| PREDICTED: syntaxin-binding protein 5 isoform X1 [Drosophila
>gi|1036811097|ref|XP_017137101.1| PREDICTED: syntaxin-binding protein 5 isoform X1 [Drosophila
>gi|1036811213|ref|XP_017137107.1| PREDICTED: syntaxin-binding protein 5 isoform X7 [Drosophila
>gi|1036811194|ref|XP_017137106.1| PREDICTED: syntaxin-binding protein 5 isoform X6 [Drosophila
>gi|1036811173|ref|XP_017137105.1| PREDICTED: syntaxin-binding protein 5 isoform X5 [Drosophila
>gi|1036811232|ref|XP_017137108.1| PREDICTED: putative glutathione-specific gamma-glutamylcyclo
>gi|1036845029|ref|XP_017138984.1| PREDICTED: uncharacterized protein LOC108153465 [Drosophila
>gi|1036765005|ref|XP_017157146.1| PREDICTED: uncharacterized protein LOC108165585 [Drosophila
>gi|1036766684|ref|XP_017134587.1| PREDICTED: sperm flagellar protein 1 [Drosophila miranda]
>gi|1036825751|ref|XP_017137904.1| PREDICTED: NK1 transcription factor-related protein 2 [Droso
>gi|1036825694|ref|XP_017137900.1| PREDICTED: folylpolyglutamate synthase, mitochondrial [Droso
>gi|1036825713|ref|XP_017137901.1| PREDICTED: folylpolyglutamate synthase, mitochondrial [Droso
>gi|1036825732|ref|XP_017137903.1| PREDICTED: box C/D snoRNA protein 1 [Drosophila miranda]
>gi|1036800658|ref|XP_017136523.1| PREDICTED: uncharacterized protein LOC108152037 isoform X3
>gi|1036800675|ref|XP_017136524.1| PREDICTED: uncharacterized protein LOC108152037 isoform X4
>gi|1036800584|ref|XP_017136519.1| PREDICTED: uncharacterized protein LOC108152037 isoform X1
>gi|1036800603|ref|XP_017136520.1| PREDICTED: uncharacterized protein LOC108152037 isoform X2
>gi|1036800621|ref|XP_017136521.1| PREDICTED: uncharacterized protein LOC108152037 isoform X2
>gi|1036800639|ref|XP_017136522.1| PREDICTED: uncharacterized protein LOC108152037 isoform X2
>gi|1036837275|ref|XP_017138559.1| PREDICTED: ATP-binding cassette sub-family B member 8, mito
>gi|1036823056|ref|XP_017137756.1| PREDICTED: LIM domain kinase 1 isoform X1 [Drosophila miran
>gi|1036823075|ref|XP_017137757.1| PREDICTED: LIM domain kinase 1 isoform X1 [Drosophila miran
>gi|1036823094|ref|XP_017137758.1| PREDICTED: LIM domain kinase 1 isoform X2 [Drosophila miran
>gi|1036819003|ref|XP_017137539.1| PREDICTED: interferon regulatory factor 2-binding protein-l
>gi|1036819022|ref|XP_017137540.1| PREDICTED: interferon regulatory factor 2-binding protein-l
>gi|1036833557|ref|XP_017138351.1| PREDICTED: uncharacterized protein LOC108153079 [Drosophila
>gi|1036833576|ref|XP_017138353.1| PREDICTED: larval cuticle protein LCP-17 [Drosophila miranda
>gi|1036837219|ref|XP_017138555.1| PREDICTED: rho GTPase-activating protein gacZ [Drosophila mi
>gi|1036849748|ref|XP_017139248.1| PREDICTED: Meckel syndrome type 1 protein [Drosophila miran
>gi|1036765417|ref|XP_017157169.1| PREDICTED: pupal cuticle protein 36 [Drosophila miranda]
>gi|1036831303|ref|XP_017138226.1| PREDICTED: uncharacterized protein LOC108152999 isoform X1
>gi|1036831322|ref|XP_017138227.1| PREDICTED: uncharacterized protein LOC108152999 isoform X2
>gi|1036831338|ref|XP_017138228.1| PREDICTED: uncharacterized protein LOC108152999 isoform X2
>gi|1036832992|ref|XP_017138319.1| PREDICTED: ras-related protein Rab-40C [Drosophila miranda]
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>gi|1036758615|ref|XP_017156788.1| PREDICTED: glycine-rich cell wall structural protein 1.8 [D:
>gi|1036758634|ref|XP_017156789.1| PREDICTED: uncharacterized protein LOC108165264 [Drosophila
>gi|1036837488|ref|XP_017138572.1| PREDICTED: putative protein kinase C delta type homolog iso:
>gi|1036837506|ref|XP_017138573.1| PREDICTED: putative protein kinase C delta type homolog iso:
>gi|1036837522|ref|XP_017138574.1| PREDICTED: putative protein kinase C delta type homolog iso:
>gi|1036837472|ref|XP_017138571.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1036837454|ref|XP_017138570.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1036764720|ref|XP_017157132.1| PREDICTED: sodium/potassium-transporting ATPase subunit beta
>gi|1036851255|ref|XP_017139337.1| PREDICTED: actin-57B-like [Drosophila miranda]
>gi|1036765890|ref|XP_017134545.1| PREDICTED: ubiquitin-conjugating enzyme E2 D2 [Drosophila m:
>gi|1036766155|ref|XP_017134561.1| PREDICTED: protein arginine N-methyltransferase 1-B-like [D:
>gi|1036851579|ref|XP_017139357.1| PREDICTED: 14-3-3 protein epsilon-like [Drosophila miranda]
>gi|1036848183|ref|XP_017139155.1| PREDICTED: midasin-like [Drosophila miranda]
>gi|1036814388|ref|XP_017137278.1| PREDICTED: uncharacterized protein LOC108152443 [Drosophila
>gi|1036764670|ref|XP_017157129.1| PREDICTED: casein kinase I [Drosophila miranda]
>gi|1036758650|ref|XP_017156790.1| PREDICTED: uncharacterized protein LOC108165266 [Drosophila
>gi|1036850217|ref|XP_017139278.1| PREDICTED: uncharacterized protein LOC108153673 isoform X2
>gi|1036850198|ref|XP_017139277.1| PREDICTED: uncharacterized protein LOC108153673 isoform X1
>gi|1036758668|ref|XP_017156791.1| PREDICTED: LOW QUALITY PROTEIN: chromosomal serine/threonine
>gi|1036784116|ref|XP_017135587.1| PREDICTED: teneurin-a [Drosophila miranda]
>gi|1036784135|ref|XP_017135588.1| PREDICTED: uncharacterized protein LOC108151472 isoform X1
>gi|1036784154|ref|XP_017135589.1| PREDICTED: uncharacterized protein LOC108151472 isoform X2
>gi|1036839595|ref|XP_017138694.1| PREDICTED: cell surface glycoprotein 1 isoform X5 [Drosophil
>gi|1036839577|ref|XP_017138693.1| PREDICTED: cell surface glycoprotein 1 isoform X4 [Drosophil
>gi|1036839558|ref|XP_017138691.1| PREDICTED: cell surface glycoprotein 1 isoform X3 [Drosophil
>gi|1036839611|ref|XP_017138695.1| PREDICTED: cell surface glycoprotein 1 isoform X6 [Drosophil
>gi|1036839540|ref|XP_017138690.1| PREDICTED: cell surface glycoprotein 1 isoform X2 [Drosophil
>gi|1036839521|ref|XP_017138689.1| PREDICTED: cell surface glycoprotein 1 isoform X1 [Drosophil
>gi|1036758687|ref|XP_017156792.1| PREDICTED: ADP-ribosylation factor-like protein 3 [Drosophil
>gi|1036764599|ref|XP_017157125.1| PREDICTED: uncharacterized protein LOC108165561 [Drosophila
>gi|1036842018|ref|XP_017138823.1| PREDICTED: sporozoite surface protein 2 isoform X2 [Drosophi
>gi|1036841804|ref|XP_017138822.1| PREDICTED: proteoglycan 4 isoform X1 [Drosophila miranda]
>gi|1036758706|ref|XP_017156794.1| PREDICTED: protein cutoff-like [Drosophila miranda]
>gi|1036842214|ref|XP_017138835.1| PREDICTED: flightin [Drosophila miranda]
>gi|1036837630|ref|XP_017138581.1| PREDICTED: protein commissureless [Drosophila miranda]
>gi|1036848387|ref|XP_017139167.1| PREDICTED: uncharacterized protein LOC108153597 [Drosophila
>gi|1036848420|ref|XP_017139169.1| PREDICTED: uncharacterized protein LOC108153598 [Drosophila
>gi|1036848434|ref|XP_017139170.1| PREDICTED: uncharacterized protein LOC108153598 [Drosophila
>gi|1036840797|ref|XP_017138761.1| PREDICTED: uncharacterized protein LOC108153336 [Drosophila
>gi|1036767145|ref|XP_017134615.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036765814|ref|XP_017134541.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036768422|ref|XP_017134681.1| PREDICTED: uncharacterized protein LOC108150883 [Drosophila
>gi|1036768703|ref|XP_017134696.1| PREDICTED: uncharacterized protein LOC108150897 [Drosophila
>gi|1036758723|ref|XP_017156795.1| PREDICTED: uncharacterized protein LOC108165271 [Drosophila
>gi|1036758741|ref|XP_017156796.1| PREDICTED: gamete and mating-type specific protein A-like [D
>gi|1036805803|ref|XP_017136813.1| PREDICTED: uncharacterized protein LOC108152183 isoform X2
>gi|1036805765|ref|XP_017136811.1| PREDICTED: uncharacterized protein LOC108152183 isoform X1
>gi|1036805784|ref|XP_017136812.1| PREDICTED: uncharacterized protein LOC108152183 isoform X1
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>gi|1036805822|ref|XP_017136814.1| PREDICTED: uncharacterized protein LOC108152183 isoform X3
>gi|1036805746|ref|XP_017136810.1| PREDICTED: zinc finger and BTB domain-containing protein 49
>gi|1036770123|ref|XP_017134776.1| PREDICTED: uncharacterized protein LOC108150953 [Drosophila
>gi|1036768344|ref|XP_017134678.1| PREDICTED: phosducin-like protein [Drosophila miranda]
>gi|1036768153|ref|XP_017134666.1| PREDICTED: uncharacterized protein LOC108150872 isoform X3
>gi|1036768117|ref|XP_017134664.1| PREDICTED: uncharacterized protein LOC108150872 isoform X1
>gi|1036768134|ref|XP_017134665.1| PREDICTED: uncharacterized protein LOC108150872 isoform X2
>gi|1036768172|ref|XP_017134667.1| PREDICTED: rho GTPase-activating protein 20 isoform X4 [Dros
>gi|1036768327|ref|XP_017134676.1| PREDICTED: ubiquitin-conjugating enzyme E2 R2 [Drosophila mi
>gi|1036768267|ref|XP_017134673.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768246|ref|XP_017134672.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768306|ref|XP_017134675.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768286|ref|XP_017134674.1| PREDICTED: probable aminoacyl tRNA synthase complex-interact
>gi|1036768189|ref|XP_017134669.1| PREDICTED: U3 small nucleolar RNA-associated protein 14 homo
>gi|1036768227|ref|XP_017134671.1| PREDICTED: general transcription factor IIH subunit 4 [Dros
>gi|1036768208|ref|XP_017134670.1| PREDICTED: probable Ufm1-specific protease 2 [Drosophila mi
>gi|1036768098|ref|XP_017134663.1| PREDICTED: microtubule-associated serine/threonine-protein
>gi|1036768078|ref|XP_017134662.1| PREDICTED: microtubule-associated serine/threonine-protein
>gi|1036772210|ref|XP_017134895.1| PREDICTED: DCN1-like protein [Drosophila miranda]
>gi|1036771984|ref|XP_017134882.1| PREDICTED: T-cell immunomodulatory protein [Drosophila miran
>gi|1036772002|ref|XP_017134883.1| PREDICTED: protein argonaute-2 [Drosophila miranda]
>gi|1036772367|ref|XP_017134903.1| PREDICTED: LOW QUALITY PROTEIN: protein disulfide-isomerase
>gi|1036772406|ref|XP_017134905.1| PREDICTED: LOW QUALITY PROTEIN: protein cutoff-like [Drosoph
>gi|1036772040|ref|XP_017134885.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772021|ref|XP_017134884.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772057|ref|XP_017134886.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772075|ref|XP_017134887.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772092|ref|XP_017134888.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772115|ref|XP_017134889.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772134|ref|XP_017134890.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772153|ref|XP_017134891.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036772172|ref|XP_017134892.1| PREDICTED: cyclic AMP response element-binding protein A iso
>gi|1036758761|ref|XP_017156797.1| PREDICTED: uncharacterized protein LOC108165274 [Drosophila
>gi|1036772327|ref|XP_017134901.1| PREDICTED: uncharacterized protein LOC108151041 [Drosophila
>gi|1036772306|ref|XP_017134900.1| PREDICTED: uncharacterized protein LOC108151040 [Drosophila
>gi|1036758779|ref|XP_017156799.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036772287|ref|XP_017134899.1| PREDICTED: BAG family molecular chaperone regulator 2 isofor
>gi|1036772230|ref|XP_017134896.1| PREDICTED: BAG family molecular chaperone regulator 2 isofor
>gi|1036772248|ref|XP_017134897.1| PREDICTED: BAG family molecular chaperone regulator 2 isofor
>gi|1036772268|ref|XP_017134898.1| PREDICTED: BAG family molecular chaperone regulator 2 isofor
>gi|1036772193|ref|XP_017134893.1| PREDICTED: uncharacterized protein LOC108151037 [Drosophila
>gi|1036772385|ref|XP_017134904.1| PREDICTED: protein midgut expression 1 [Drosophila miranda]
>gi|1036772341|ref|XP_017134902.1| PREDICTED: uncharacterized protein LOC108151042 [Drosophila
>gi|1036834954|ref|XP_017138421.1| PREDICTED: peptide methionine sulfoxide reductase isoform X2
>gi|1036834935|ref|XP_017138420.1| PREDICTED: peptide methionine sulfoxide reductase isoform X1
>gi|1036834970|ref|XP_017138422.1| PREDICTED: peptide methionine sulfoxide reductase isoform X3
>gi|1036834989|ref|XP_017138423.1| PREDICTED: gonadal protein gdl isoform X1 [Drosophila mirand
>gi|1036835004|ref|XP_017138424.1| PREDICTED: gonadal protein gdl isoform X2 [Drosophila mirand
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>gi|1036763476|ref|XP_017157068.1| PREDICTED: protein Z600 [Drosophila miranda]
>gi|1036841769|ref|XP_017138820.1| PREDICTED: uncharacterized protein LOC108153364 [Drosophila
>gi|1036822908|ref|XP_017137747.1| PREDICTED: OTU domain-containing protein 6B [Drosophila mira
>gi|1036822925|ref|XP_017137748.1| PREDICTED: sugar transporter SWEET1 [Drosophila miranda]
>gi|1036822889|ref|XP_017137746.1| PREDICTED: DDB1- and CUL4-associated factor 13 [Drosophila m
>gi|1036835719|ref|XP_017138466.1| PREDICTED: uncharacterized protein LOC108153172 [Drosophila
>gi|1036835738|ref|XP_017138468.1| PREDICTED: mpv17-like protein isoform X1 [Drosophila miranda
>gi|1036835756|ref|XP_017138469.1| PREDICTED: mpv17-like protein isoform X2 [Drosophila miranda
>gi|1036835776|ref|XP_017138470.1| PREDICTED: mpv17-like protein isoform X2 [Drosophila miranda
>gi|1036829575|ref|XP_017138124.1| PREDICTED: protein toll [Drosophila miranda]
>gi|1036823723|ref|XP_017137793.1| PREDICTED: cationic amino acid transporter 2 isoform X2 [Dro
>gi|1036823665|ref|XP_017137790.1| PREDICTED: high affinity cationic amino acid transporter 1 i
>gi|1036823683|ref|XP_017137791.1| PREDICTED: high affinity cationic amino acid transporter 1 i
>gi|1036823704|ref|XP_017137792.1| PREDICTED: high affinity cationic amino acid transporter 1 i
>gi|1036823740|ref|XP_017137794.1| PREDICTED: bestrophin-2 [Drosophila miranda]
>gi|1036852454|ref|XP_017139412.1| PREDICTED: bestrophin-4 [Drosophila miranda]
>gi|1036813967|ref|XP_017137253.1| PREDICTED: protein toll [Drosophila miranda]
>gi|1036782493|ref|XP_017135490.1| PREDICTED: endoplasmic reticulum-Golgi intermediate compartm
>gi|1036782507|ref|XP_017135491.1| PREDICTED: reactive oxygen species modulator 1 [Drosophila m
>gi|1036782455|ref|XP_017135488.1| PREDICTED: U4/U6 small nuclear ribonucleoprotein Prp31 [Dros
>gi|1036782379|ref|XP_017135484.1| PREDICTED: probable DNA mismatch repair protein Msh6 [Drosop
>gi|1036782397|ref|XP_017135485.1| PREDICTED: glycoprotein 3-alpha-L-fucosyltransferase A [Dros
>gi|1036782419|ref|XP_017135486.1| PREDICTED: glycoprotein 3-alpha-L-fucosyltransferase A [Dros
>gi|1036782437|ref|XP_017135487.1| PREDICTED: glycoprotein 3-alpha-L-fucosyltransferase A [Dros
>gi|1036782474|ref|XP_017135489.1| PREDICTED: peroxisomal biogenesis factor 3 [Drosophila mira
>gi|1036815904|ref|XP_017137361.1| PREDICTED: pyroglutamyl-peptidase 1 [Drosophila miranda]
>gi|1036815923|ref|XP_017137362.1| PREDICTED: uncharacterized protein LOC108152498 [Drosophila
>gi|1036815883|ref|XP_017137360.1| PREDICTED: protein disulfide-isomerase [Drosophila miranda]
>gi|1036790178|ref|XP_017135932.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X1
>gi|1036790197|ref|XP_017135933.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X2
>gi|1036790215|ref|XP_017135934.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X2
>gi|1036790236|ref|XP_017135935.1| PREDICTED: uncharacterized protein DDB_G0284459 isoform X2
>gi|1036790159|ref|XP_017135931.1| PREDICTED: centrosomal protein of 135 kDa isoform X4 [Drosop
>gi|1036790064|ref|XP_017135926.1| PREDICTED: uncharacterized protein LOC108151691 isoform X1
>gi|1036790083|ref|XP_017135927.1| PREDICTED: uncharacterized protein LOC108151691 isoform X2
>gi|1036790140|ref|XP_017135930.1| PREDICTED: centrosomal protein of 135 kDa isoform X3 [Drosop
>gi|1036790121|ref|XP_017135929.1| PREDICTED: centrosomal protein of 135 kDa isoform X2 [Drosop
>gi|1036790102|ref|XP_017135928.1| PREDICTED: centrosomal protein of 135 kDa isoform X1 [Drosop
>gi|1036790252|ref|XP_017135936.1| PREDICTED: uncharacterized protein LOC108151695 isoform X1
>gi|1036790269|ref|XP_017135937.1| PREDICTED: uncharacterized protein LOC108151695 isoform X2
>gi|1036852144|ref|XP_017139393.1| PREDICTED: uncharacterized protein LOC108153765 [Drosophila
>gi|1036813503|ref|XP_017137226.1| PREDICTED: CTP synthase isoform X1 [Drosophila miranda]
>gi|1036813557|ref|XP_017137229.1| PREDICTED: CTP synthase isoform X3 [Drosophila miranda]
>gi|1036813522|ref|XP_017137227.1| PREDICTED: uncharacterized protein LOC108152422 isoform X2
>gi|1036813537|ref|XP_017137228.1| PREDICTED: uncharacterized protein LOC108152422 isoform X2
>gi|1036813597|ref|XP_017137231.1| PREDICTED: uncharacterized protein LOC108152423 isoform X2
>gi|1036813578|ref|XP_017137230.1| PREDICTED: uncharacterized protein LOC108152423 isoform X1
>gi|1036833393|ref|XP_017138343.1| PREDICTED: glycine--tRNA ligase [Drosophila miranda]
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>gi|1036777693|ref|XP_017135206.1| PREDICTED: putative uncharacterized protein DDB_G0271606 [Di
>gi|1036777453|ref|XP_017135192.1| PREDICTED: A-kinase anchor protein 9 isoform X2 [Drosophila
>gi|1036777509|ref|XP_017135196.1| PREDICTED: golgin subfamily A member 4 isoform X5 [Drosophil
>gi|1036777528|ref|XP_017135197.1| PREDICTED: golgin subfamily A member 4 isoform X6 [Drosophil
>gi|1036777566|ref|XP_017135199.1| PREDICTED: pericentrin isoform X8 [Drosophila miranda]
>gi|1036777472|ref|XP_017135193.1| PREDICTED: A-kinase anchor protein 9 isoform X3 [Drosophila
>gi|1036777547|ref|XP_017135198.1| PREDICTED: golgin subfamily A member 4 isoform X7 [Drosophil
>gi|1036777582|ref|XP_017135200.1| PREDICTED: pericentrin isoform X9 [Drosophila miranda]
>gi|1036777601|ref|XP_017135201.1| PREDICTED: pericentrin isoform X10 [Drosophila miranda]
>gi|1036777620|ref|XP_017135202.1| PREDICTED: golgin subfamily A member 4 isoform X11 [Drosophi
>gi|1036777434|ref|XP_017135191.1| PREDICTED: A-kinase anchor protein 9 isoform X1 [Drosophila
>gi|1036777490|ref|XP_017135195.1| PREDICTED: A-kinase anchor protein 9 isoform X4 [Drosophila
>gi|1036777637|ref|XP_017135203.1| PREDICTED: A-kinase anchor protein 9 isoform X12 [Drosophila
>gi|1036777786|ref|XP_017135212.1| PREDICTED: putative odorant receptor 71a [Drosophila miranda
>gi|1036777656|ref|XP_017135204.1| PREDICTED: pericentrin isoform X13 [Drosophila miranda]
>gi|1036777675|ref|XP_017135205.1| PREDICTED: trichohyalin isoform X14 [Drosophila miranda]
>gi|1036777824|ref|XP_017135214.1| PREDICTED: uncharacterized protein LOC108151228 [Drosophila
>gi|1036777842|ref|XP_017135215.1| PREDICTED: uncharacterized protein LOC108151229 [Drosophila
>gi|1036777905|ref|XP_017135219.1| PREDICTED: prostatic spermine-binding protein [Drosophila m
>gi|1036777870|ref|XP_017135218.1| PREDICTED: probable DNA-directed RNA polymerase subunit del
>gi|1036777857|ref|XP_017135216.1| PREDICTED: uncharacterized protein LOC108151230 [Drosophila
>gi|1036777767|ref|XP_017135211.1| PREDICTED: synaptotagmin-5 isoform X2 [Drosophila miranda]
>gi|1036777805|ref|XP_017135213.1| PREDICTED: GS homeobox 1 [Drosophila miranda]
>gi|1036777710|ref|XP_017135207.1| PREDICTED: synaptotagmin-3 isoform X1 [Drosophila miranda]
>gi|1036777729|ref|XP_017135208.1| PREDICTED: synaptotagmin-3 isoform X1 [Drosophila miranda]
>gi|1036777748|ref|XP_017135210.1| PREDICTED: synaptotagmin-3 isoform X1 [Drosophila miranda]
>gi|1036765796|ref|XP_017134540.1| PREDICTED: uncharacterized protein LOC108150757 [Drosophila
>gi|1036829999|ref|XP_017138148.1| PREDICTED: reversion-inducing cysteine-rich protein with Ka
>gi|1036830018|ref|XP_017138149.1| PREDICTED: reversion-inducing cysteine-rich protein with Ka
>gi|1036830037|ref|XP_017138150.1| PREDICTED: proteasome subunit beta type-7 [Drosophila miran
>gi|1036833027|ref|XP_017138321.1| PREDICTED: 39S ribosomal protein L39, mitochondrial [Drosoph
>gi|1036833010|ref|XP_017138320.1| PREDICTED: zinc transporter ZIP10 [Drosophila miranda]
>gi|1036821311|ref|XP_017137658.1| PREDICTED: Y+L amino acid transporter 2 [Drosophila miranda]
>gi|1036821330|ref|XP_017137659.1| PREDICTED: Y+L amino acid transporter 2 [Drosophila miranda]
>gi|1036821349|ref|XP_017137660.1| PREDICTED: angio-associated migratory cell protein [Drosophi
>gi|1036848860|ref|XP_017139195.1| PREDICTED: uncharacterized protein LOC108153608 isoform X2
>gi|1036848841|ref|XP_017139194.1| PREDICTED: uncharacterized protein LOC108153608 isoform X1
>gi|1036763155|ref|XP_017157052.1| PREDICTED: enhancer of split malpha protein [Drosophila mira
>gi|1036763331|ref|XP_017157059.1| PREDICTED: uncharacterized protein LOC108165514 [Drosophila
>gi|1036846698|ref|XP_017139073.1| PREDICTED: enhancer of split malpha protein [Drosophila mira
>gi|1036850162|ref|XP_017139274.1| PREDICTED: uridine phosphorylase 1-like [Drosophila miranda]
>gi|1036763315|ref|XP_017157058.1| PREDICTED: uncharacterized protein LOC108165513 [Drosophila
>gi|1036758797|ref|XP_017156800.1| PREDICTED: uncharacterized protein LOC108165276 [Drosophila
>gi|1036849104|ref|XP_017139209.1| PREDICTED: echinoderm microtubule-associated protein-like CO
>gi|1036767087|ref|XP_017134612.1| PREDICTED: uncharacterized protein LOC108150817 isoform X1
>gi|1036767108|ref|XP_017134613.1| PREDICTED: uncharacterized protein LOC108150817 isoform X2
>gi|1036830096|ref|XP_017138154.1| PREDICTED: Ig-like and fibronectin type-III domain-containin
>gi|1036812414|ref|XP_017137171.1| PREDICTED: uncharacterized protein LOC108152385 [Drosophila
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>gi|1036812336|ref|XP_017137166.1| PREDICTED: probable helicase with zinc finger domain [Droso]
>gi|1036812359|ref|XP_017137167.1| PREDICTED: probable helicase with zinc finger domain [Droso]
>gi|1036812376|ref|XP_017137168.1| PREDICTED: probable helicase with zinc finger domain [Droso]
>gi|1036812395|ref|XP_017137169.1| PREDICTED: probable helicase with zinc finger domain [Droso]
>gi|1036838573|ref|XP_017138637.1| PREDICTED: H/ACA ribonucleoprotein complex subunit 2-like p]
>gi|1036833046|ref|XP_017138322.1| PREDICTED: tudor domain-containing protein 3 [Drosophila mi]
>gi|1036827923|ref|XP_017138033.1| PREDICTED: patatin-like phospholipase domain-containing pro]
>gi|1036811742|ref|XP_017137134.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 2]
>gi|1036825053|ref|XP_017137867.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylg]
>gi|1036825071|ref|XP_017137868.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylg]
>gi|1036825088|ref|XP_017137870.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylg]
>gi|1036825107|ref|XP_017137871.1| PREDICTED: alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylg]
>gi|1036774850|ref|XP_017135049.1| PREDICTED: uncharacterized protein LOC108151135 [Drosophila]
>gi|1036774781|ref|XP_017135044.1| PREDICTED: transcription factor GAGA isoform X4 [Drosophila]
>gi|1036774727|ref|XP_017135042.1| PREDICTED: transcription factor GAGA isoform X2 [Drosophila]
>gi|1036774762|ref|XP_017135043.1| PREDICTED: transcription factor GAGA isoform X3 [Drosophila]
>gi|1036774708|ref|XP_017135041.1| PREDICTED: transcription factor GAGA isoform X1 [Drosophila]
>gi|1036774816|ref|XP_017135046.1| PREDICTED: protein roadkill [Drosophila miranda]
>gi|1036774632|ref|XP_017135037.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X1 [Drosoph]
>gi|1036774651|ref|XP_017135038.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X2 [Drosoph]
>gi|1036774670|ref|XP_017135039.1| PREDICTED: venom dipeptidyl peptidase 4 isoform X3 [Drosoph]
>gi|1036774903|ref|XP_017135052.1| PREDICTED: thioredoxin-2 [Drosophila miranda]
>gi|1036774885|ref|XP_017135051.1| PREDICTED: uncharacterized protein LOC108151137 [Drosophila]
>gi|1036774689|ref|XP_017135040.1| PREDICTED: myosin heavy chain, striated muscle [Drosophila ]
>gi|1036774922|ref|XP_017135053.1| PREDICTED: uncharacterized protein LOC108151139 [Drosophila]
>gi|1036774868|ref|XP_017135050.1| PREDICTED: uncharacterized protein LOC108151136 [Drosophila]
>gi|1036774943|ref|XP_017135054.1| PREDICTED: uncharacterized protein LOC108151140 [Drosophila]
>gi|1036774834|ref|XP_017135048.1| PREDICTED: protein PIH1D3 [Drosophila miranda]
>gi|1036774799|ref|XP_017135045.1| PREDICTED: venom protease [Drosophila miranda]
>gi|1036815942|ref|XP_017137363.1| PREDICTED: glypican-4 [Drosophila miranda]
>gi|1036834338|ref|XP_017138384.1| PREDICTED: polycystic kidney disease 2-like 2 protein [Dros]
>gi|1036834375|ref|XP_017138386.1| PREDICTED: nucleoside-triphosphatase THEP1 [Drosophila mira]
>gi|1036834356|ref|XP_017138385.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC5 [Dr]
>gi|1036835252|ref|XP_017138439.1| PREDICTED: methyltransferase-like protein 22 [Drosophila mi]
>gi|1036835236|ref|XP_017138437.1| PREDICTED: decaprenyl-diphosphate synthase subunit 2 [Droso]
>gi|1036801320|ref|XP_017136561.1| PREDICTED: mediator of RNA polymerase II transcription subu]
>gi|1036801339|ref|XP_017136562.1| PREDICTED: mediator of RNA polymerase II transcription subu]
>gi|1036801358|ref|XP_017136563.1| PREDICTED: mediator of RNA polymerase II transcription subu]
>gi|1036801377|ref|XP_017136565.1| PREDICTED: mediator of RNA polymerase II transcription subu]
>gi|1036764686|ref|XP_017157130.1| PREDICTED: uncharacterized protein LOC108165567 [Drosophila]
>gi|1036851440|ref|XP_017139348.1| PREDICTED: uncharacterized protein LOC108153730 [Drosophila]
>gi|1036843488|ref|XP_017138907.1| PREDICTED: uncharacterized protein LOC108153417 isoform X1 ]
>gi|1036843507|ref|XP_017138908.1| PREDICTED: uncharacterized protein LOC108153417 isoform X2 ]
>gi|1036843526|ref|XP_017138909.1| PREDICTED: uncharacterized protein LOC108153417 isoform X3 ]
>gi|1036843471|ref|XP_017138906.1| PREDICTED: uncharacterized protein LOC108153416 [Drosophila]
>gi|1036843545|ref|XP_017138910.1| PREDICTED: trypsin alpha-3 [Drosophila miranda]
>gi|1036840814|ref|XP_017138762.1| PREDICTED: fringe glycosyltransferase [Drosophila miranda]
>gi|1036840831|ref|XP_017138763.1| PREDICTED: uncharacterized protein LOC108153339 [Drosophila]
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>gi|1036758816|ref|XP_017156801.1| PREDICTED: uncharacterized protein LOC108165277 [Drosophila
>gi|1036830224|ref|XP_017138163.1| PREDICTED: uncharacterized protein LOC108152966 isoform X1
>gi|1036830241|ref|XP_017138164.1| PREDICTED: uncharacterized protein LOC108152966 isoform X2
>gi|1036830316|ref|XP_017138168.1| PREDICTED: proteasome subunit alpha type-1-like [Drosophila
>gi|1036830260|ref|XP_017138165.1| PREDICTED: zinc finger protein 48 isoform X1 [Drosophila mi
>gi|1036830279|ref|XP_017138166.1| PREDICTED: zinc finger protein 48 isoform X1 [Drosophila mi
>gi|1036830298|ref|XP_017138167.1| PREDICTED: zinc finger protein 48 isoform X2 [Drosophila mi
>gi|1036840407|ref|XP_017138739.1| PREDICTED: transcriptional protein SWT1 isoform X1 [Drosophi
>gi|1036840423|ref|XP_017138740.1| PREDICTED: transcriptional protein SWT1 isoform X2 [Drosophi
>gi|1036798087|ref|XP_017136377.1| PREDICTED: ADP-ribosylation factor 1-like isoform X2 [Droso
>gi|1036798068|ref|XP_017136376.1| PREDICTED: ADP-ribosylation factor 1-like isoform X1 [Droso
>gi|1036798105|ref|XP_017136378.1| PREDICTED: ADP-ribosylation factor 1-like isoform X1 [Droso
>gi|1036797978|ref|XP_017136371.1| PREDICTED: COP9 signalosome complex subunit 3 isoform X1 [D
>gi|1036797997|ref|XP_017136372.1| PREDICTED: COP9 signalosome complex subunit 3 isoform X2 [D
>gi|1036798161|ref|XP_017136381.1| PREDICTED: uncharacterized protein LOC108151954 [Drosophila
>gi|1036798052|ref|XP_017136375.1| PREDICTED: uncharacterized protein LOC108151949 [Drosophila
>gi|1036798014|ref|XP_017136373.1| PREDICTED: cAMP-dependent protein kinase type I regulatory s
>gi|1036798033|ref|XP_017136374.1| PREDICTED: cAMP-dependent protein kinase type I regulatory s
>gi|1036798143|ref|XP_017136380.1| PREDICTED: uncharacterized protein LOC108151953 [Drosophila
>gi|1036798124|ref|XP_017136379.1| PREDICTED: uncharacterized protein LOC108151952 [Drosophila
>gi|1036797960|ref|XP_017136370.1| PREDICTED: uncharacterized protein LOC108151946 [Drosophila
>gi|1036800529|ref|XP_017136514.1| PREDICTED: neurofilament heavy polypeptide isoform X2 [Droso
>gi|1036800510|ref|XP_017136513.1| PREDICTED: uncharacterized protein LOC108152034 isoform X1
>gi|1036800565|ref|XP_017136518.1| PREDICTED: protein ST7 homolog [Drosophila miranda]
>gi|1036800548|ref|XP_017136515.1| PREDICTED: uncharacterized protein LOC108152035 [Drosophila
>gi|1036806686|ref|XP_017136858.1| PREDICTED: translocation protein SEC63 homolog [Drosophila r
>gi|1036806723|ref|XP_017136860.1| PREDICTED: neurofilament medium polypeptide isoform X2 [Dros
>gi|1036806704|ref|XP_017136859.1| PREDICTED: methyl-CpG-binding domain-containing protein 10 i
>gi|1036811459|ref|XP_017137118.1| PREDICTED: akirin [Drosophila miranda]
>gi|1036811478|ref|XP_017137120.1| PREDICTED: sphingomyelin synthase-related 1 [Drosophila mira
>gi|1036822326|ref|XP_017137714.1| PREDICTED: nuclear valosin-containing protein-like isoform X
>gi|1036822345|ref|XP_017137715.1| PREDICTED: nuclear valosin-containing protein-like isoform X
>gi|1036822364|ref|XP_017137716.1| PREDICTED: nuclear valosin-containing protein-like isoform X
>gi|1036822383|ref|XP_017137717.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036851560|ref|XP_017139356.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036764838|ref|XP_017157139.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036764003|ref|XP_017157099.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036851726|ref|XP_017139366.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036850303|ref|XP_017139281.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036831754|ref|XP_017138250.1| PREDICTED: ras-related protein Rac2 [Drosophila miranda]
>gi|1036831716|ref|XP_017138248.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036831735|ref|XP_017138249.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036766702|ref|XP_017134589.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1036766270|ref|XP_017134563.1| PREDICTED: protein late bloomer isoform X1 [Drosophila mira
>gi|1036766289|ref|XP_017134564.1| PREDICTED: protein late bloomer isoform X2 [Drosophila mira
>gi|1036766308|ref|XP_017134565.1| PREDICTED: protein late bloomer isoform X3 [Drosophila mira
>gi|1036769646|ref|XP_017134752.1| PREDICTED: cuticle protein 76 [Drosophila miranda]
>gi|1036768775|ref|XP_017134700.1| PREDICTED: cuticle protein 16.5 [Drosophila miranda]
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>gi|1036768517|ref|XP_017134686.1| PREDICTED: uncharacterized protein LOC108150888 isoform X1
>gi|1036768536|ref|XP_017134687.1| PREDICTED: uncharacterized protein LOC108150888 isoform X2
>gi|1036768558|ref|XP_017134688.1| PREDICTED: uncharacterized protein LOC108150888 isoform X3
>gi|1036787543|ref|XP_017135780.1| PREDICTED: probable cytochrome P450 316a1 [Drosophila mirand
>gi|1036787505|ref|XP_017135778.1| PREDICTED: cytochrome P450 4d8 isoform X1 [Drosophila mirand
>gi|1036787524|ref|XP_017135779.1| PREDICTED: cytochrome P450 4d8 isoform X2 [Drosophila mirand
>gi|1036787559|ref|XP_017135781.1| PREDICTED: carboxypeptidase B [Drosophila miranda]
>gi|1036758835|ref|XP_017156802.1| PREDICTED: zinc carboxypeptidase-like [Drosophila miranda]
>gi|1036787577|ref|XP_017135782.1| PREDICTED: cuticle protein LPCP-23 [Drosophila miranda]
>gi|1036787372|ref|XP_017135770.1| PREDICTED: large proline-rich protein BAG6 isoform X1 [Dros
>gi|1036787391|ref|XP_017135772.1| PREDICTED: large proline-rich protein BAG6 isoform X1 [Dros
>gi|1036787410|ref|XP_017135773.1| PREDICTED: large proline-rich protein BAG6 isoform X1 [Dros
>gi|1036787446|ref|XP_017135775.1| PREDICTED: large proline-rich protein BAG6 isoform X3 [Dros
>gi|1036787429|ref|XP_017135774.1| PREDICTED: large proline-rich protein BAG6 isoform X2 [Dros
>gi|1036787467|ref|XP_017135776.1| PREDICTED: large proline-rich protein BAG6 isoform X4 [Dros
>gi|1036787486|ref|XP_017135777.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036814640|ref|XP_017137292.1| PREDICTED: epsin-1 isoform X1 [Drosophila miranda]
>gi|1036814661|ref|XP_017137293.1| PREDICTED: epsin-1 isoform X1 [Drosophila miranda]
>gi|1036814698|ref|XP_017137295.1| PREDICTED: epsin-1 isoform X3 [Drosophila miranda]
>gi|1036814717|ref|XP_017137296.1| PREDICTED: epsin-1 isoform X4 [Drosophila miranda]
>gi|1036814736|ref|XP_017137297.1| PREDICTED: epsin-1 isoform X5 [Drosophila miranda]
>gi|1036814680|ref|XP_017137294.1| PREDICTED: epsin-1 isoform X2 [Drosophila miranda]
>gi|1036764616|ref|XP_017157126.1| PREDICTED: trypsin [Drosophila miranda]
>gi|1036758851|ref|XP_017156803.1| PREDICTED: trypsin beta-like [Drosophila miranda]
>gi|1036758870|ref|XP_017156804.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036846846|ref|XP_017139081.1| PREDICTED: WD repeat-containing protein 63 [Drosophila miran
>gi|1036850818|ref|XP_017139311.1| PREDICTED: gamma-tubulin complex component 2 homolog [Drosop
>gi|1036814926|ref|XP_017137308.1| PREDICTED: uncharacterized protein LOC108152455 [Drosophila
>gi|1036814850|ref|XP_017137304.1| PREDICTED: uncharacterized protein LOC108152453 isoform X1
>gi|1036814888|ref|XP_017137306.1| PREDICTED: protein late bloomer-like isoform X1 [Drosophila
>gi|1036814907|ref|XP_017137307.1| PREDICTED: uncharacterized protein LOC108152454 isoform X2
>gi|1036814869|ref|XP_017137305.1| PREDICTED: pleckstrin homology domain-containing family G me
>gi|1036814966|ref|XP_017137309.1| PREDICTED: ubiquitin-related modifier 1 homolog [Drosophila
>gi|1036836279|ref|XP_017138500.1| PREDICTED: uncharacterized protein LOC108153193 [Drosophila
>gi|1036836298|ref|XP_017138501.1| PREDICTED: transmembrane protein 70 homolog, mitochondrial
>gi|1036758889|ref|XP_017156805.1| PREDICTED: peptidoglycan-recognition protein SD [Drosophila
>gi|1036836598|ref|XP_017138520.1| PREDICTED: uncharacterized protein LOC108153209 [Drosophila
>gi|1036798880|ref|XP_017136422.1| PREDICTED: ankyrin-3 isoform X5 [Drosophila miranda]
>gi|1036798842|ref|XP_017136420.1| PREDICTED: ankyrin-2 isoform X3 [Drosophila miranda]
>gi|1036798823|ref|XP_017136419.1| PREDICTED: ankyrin-2 isoform X2 [Drosophila miranda]
>gi|1036798787|ref|XP_017136417.1| PREDICTED: ankyrin-2 isoform X1 [Drosophila miranda]
>gi|1036798804|ref|XP_017136418.1| PREDICTED: ankyrin-2 isoform X1 [Drosophila miranda]
>gi|1036798961|ref|XP_017136427.1| PREDICTED: ankyrin-2 isoform X9 [Drosophila miranda]
>gi|1036798941|ref|XP_017136426.1| PREDICTED: ankyrin-2 isoform X8 [Drosophila miranda]
>gi|1036798998|ref|XP_017136429.1| PREDICTED: ankyrin-2 isoform X11 [Drosophila miranda]
>gi|1036798979|ref|XP_017136428.1| PREDICTED: ankyrin-2 isoform X10 [Drosophila miranda]
>gi|1036798922|ref|XP_017136425.1| PREDICTED: ankyrin-2 isoform X7 [Drosophila miranda]
>gi|1036798861|ref|XP_017136421.1| PREDICTED: ankyrin-2 isoform X4 [Drosophila miranda]
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>gi|1036799016|ref|XP_017136430.1| PREDICTED: uncharacterized protein LOC108151976 [Drosophila
>gi|1036798899|ref|XP_017136423.1| PREDICTED: ankyrin-3 isoform X6 [Drosophila miranda]
>gi|1036794192|ref|XP_017136157.1| PREDICTED: probable medium-chain specific acyl-CoA dehydroge
>gi|1036794140|ref|XP_017136154.1| PREDICTED: tonsoku-like protein [Drosophila miranda]
>gi|1036794156|ref|XP_017136155.1| PREDICTED: zinc finger RNA-binding protein isoform X1 [Droso
>gi|1036794174|ref|XP_017136156.1| PREDICTED: zinc finger RNA-binding protein isoform X2 [Droso
>gi|1036819780|ref|XP_017137580.1| PREDICTED: integrator complex subunit 9 [Drosophila miranda]
>gi|1036819761|ref|XP_017137579.1| PREDICTED: isoleucine--tRNA ligase, mitochondrial [Drosophil
>gi|1036849676|ref|XP_017139244.1| PREDICTED: ATP synthase subunit beta, mitochondrial [Drosoph
>gi|1036778461|ref|XP_017135254.1| PREDICTED: uncharacterized protein LOC108151250 [Drosophila
>gi|1036778443|ref|XP_017135253.1| PREDICTED: H(+)/Cl(-) exchange transporter 3 [Drosophila mir
>gi|1036778479|ref|XP_017135255.1| PREDICTED: threonine aspartase 1 [Drosophila miranda]
>gi|1036778497|ref|XP_017135257.1| PREDICTED: uncharacterized protein LOC108151252 [Drosophila
>gi|1036778423|ref|XP_017135252.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778388|ref|XP_017135250.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778367|ref|XP_017135249.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A iso
>gi|1036778405|ref|XP_017135251.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778348|ref|XP_017135248.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778183|ref|XP_017135238.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A iso
>gi|1036778166|ref|XP_017135237.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778311|ref|XP_017135246.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778255|ref|XP_017135243.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778236|ref|XP_017135241.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778217|ref|XP_017135240.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A iso
>gi|1036778200|ref|XP_017135239.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778274|ref|XP_017135244.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778149|ref|XP_017135236.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778293|ref|XP_017135245.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036778329|ref|XP_017135247.1| PREDICTED: protein phosphatase 1 regulatory subunit 12B iso
>gi|1036813691|ref|XP_017137237.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813616|ref|XP_017137232.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813635|ref|XP_017137233.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813653|ref|XP_017137234.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036813672|ref|XP_017137236.1| PREDICTED: death-associated inhibitor of apoptosis 1 isofor
>gi|1036830115|ref|XP_017138155.1| PREDICTED: uncharacterized protein LOC108152963 [Drosophila
>gi|1036830170|ref|XP_017138159.1| PREDICTED: uncharacterized protein LOC108152965 isoform X1
>gi|1036830205|ref|XP_017138162.1| PREDICTED: uncharacterized protein LOC108152965 isoform X3
>gi|1036830187|ref|XP_017138160.1| PREDICTED: uncharacterized protein LOC108152965 isoform X2
>gi|1036830133|ref|XP_017138157.1| PREDICTED: uncharacterized protein LOC108152964 [Drosophila
>gi|1036830152|ref|XP_017138158.1| PREDICTED: uncharacterized protein LOC108152964 [Drosophila
>gi|1036824294|ref|XP_017137825.1| PREDICTED: uncharacterized protein LOC108152764 isoform X1
>gi|1036824313|ref|XP_017137827.1| PREDICTED: putative dipeptidase CPC735_014430 isoform X2 [Dr
>gi|1036824452|ref|XP_017137836.1| PREDICTED: mpv17-like protein 2 isoform X1 [Drosophila miran
>gi|1036824469|ref|XP_017137837.1| PREDICTED: mpv17-like protein 2 isoform X2 [Drosophila miran
>gi|1036824415|ref|XP_017137833.1| PREDICTED: uncharacterized protein LOC108152769 isoform X1
>gi|1036824434|ref|XP_017137835.1| PREDICTED: uncharacterized protein LOC108152769 isoform X2
>gi|1036824379|ref|XP_017137831.1| PREDICTED: uncharacterized protein LOC108152767 [Drosophila
>gi|1036824396|ref|XP_017137832.1| PREDICTED: peritrophin-44 [Drosophila miranda]
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>gi|1036824362|ref|XP_017137830.1| PREDICTED: succinate dehydrogenase [ubiquinone] flavoprotei
>gi|1036824329|ref|XP_017137828.1| PREDICTED: T-related protein isoform X1 [Drosophila miranda]
>gi|1036824346|ref|XP_017137829.1| PREDICTED: T-related protein isoform X2 [Drosophila miranda]
>gi|1036851218|ref|XP_017139335.1| PREDICTED: endonuclease G, mitochondrial [Drosophila miranda
>gi|1036849544|ref|XP_017139236.1| PREDICTED: 26S protease regulatory subunit 10B [Drosophila m
>gi|1036849561|ref|XP_017139237.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036771620|ref|XP_017134861.1| PREDICTED: chondroitin proteoglycan 1 [Drosophila miranda]
>gi|1036771637|ref|XP_017134862.1| PREDICTED: chondroitin proteoglycan 2 [Drosophila miranda]
>gi|1036771748|ref|XP_017134868.1| PREDICTED: chondroitin proteoglycan 2 [Drosophila miranda]
>gi|1036771565|ref|XP_017134857.1| PREDICTED: uncharacterized protein LOC108151010 isoform X1
>gi|1036771601|ref|XP_017134860.1| PREDICTED: uncharacterized protein LOC108151010 isoform X3
>gi|1036771584|ref|XP_017134858.1| PREDICTED: uncharacterized protein LOC108151010 isoform X2
>gi|1036771767|ref|XP_017134869.1| PREDICTED: uncharacterized protein LOC108151021 [Drosophila
>gi|1036758905|ref|XP_017156806.1| PREDICTED: uncharacterized protein LOC108165282 [Drosophila
>gi|1036771471|ref|XP_017134852.1| PREDICTED: glutenin, high molecular weight subunit DX5 isof
>gi|1036771452|ref|XP_017134851.1| PREDICTED: uncharacterized PE-PGRS family protein PE_PGRS54
>gi|1036771545|ref|XP_017134856.1| PREDICTED: flocculation protein FLO11 isoform X3 [Drosophila
>gi|1036771508|ref|XP_017134854.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036771526|ref|XP_017134855.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036771821|ref|XP_017134873.1| PREDICTED: peritrophin-1-like [Drosophila miranda]
>gi|1036771802|ref|XP_017134872.1| PREDICTED: integumentary mucin B.1-like [Drosophila miranda]
>gi|1036771269|ref|XP_017134840.1| PREDICTED: uncharacterized protein LOC108151001 [Drosophila
>gi|1036771489|ref|XP_017134853.1| PREDICTED: anoctamin-4 [Drosophila miranda]
>gi|1036771729|ref|XP_017134867.1| PREDICTED: tubulin polyglutamylase complex subunit 2 [Droso
>gi|1036771786|ref|XP_017134871.1| PREDICTED: viral IAP-associated factor homolog [Drosophila m
>gi|1036771435|ref|XP_017134850.1| PREDICTED: uncharacterized protein LOC108151005 [Drosophila
>gi|1036771872|ref|XP_017134875.1| PREDICTED: cAMP-regulated phosphoprotein 19 [Drosophila mir
>gi|1036771287|ref|XP_017134841.1| PREDICTED: tuberin [Drosophila miranda]
>gi|1036771710|ref|XP_017134866.1| PREDICTED: uncharacterized protein LOC108151016 [Drosophila
>gi|1036771656|ref|XP_017134863.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036771692|ref|XP_017134865.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036771400|ref|XP_017134848.1| PREDICTED: uncharacterized protein LOC108151004 isoform X4
>gi|1036771342|ref|XP_017134844.1| PREDICTED: uncharacterized protein LOC108151004 isoform X3
>gi|1036771361|ref|XP_017134845.1| PREDICTED: uncharacterized protein LOC108151004 isoform X3
>gi|1036771380|ref|XP_017134847.1| PREDICTED: uncharacterized protein LOC108151004 isoform X3
>gi|1036771323|ref|XP_017134843.1| PREDICTED: uncharacterized protein LOC108151004 isoform X2
>gi|1036771304|ref|XP_017134842.1| PREDICTED: uncharacterized protein LOC108151004 isoform X1
>gi|1036771418|ref|XP_017134849.1| PREDICTED: uncharacterized protein LOC108151004 isoform X5
>gi|1036771840|ref|XP_017134874.1| PREDICTED: uncharacterized protein LOC108151025 [Drosophila
>gi|1036771891|ref|XP_017134876.1| PREDICTED: uncharacterized protein LOC108151027 [Drosophila
>gi|1036771674|ref|XP_017134864.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036764876|ref|XP_017157141.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036849232|ref|XP_017139216.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036850651|ref|XP_017139302.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036765507|ref|XP_017157175.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036764281|ref|XP_017157112.1| PREDICTED: peritrophin-44 isoform X2 [Drosophila miranda]
>gi|1036764261|ref|XP_017157111.1| PREDICTED: peritrophin-44 isoform X1 [Drosophila miranda]
>gi|1036764965|ref|XP_017157144.1| PREDICTED: peritrophin-44 [Drosophila miranda]
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>gi|1036824848|ref|XP_017137857.1| PREDICTED: peritrophin-44 [Drosophila miranda]
>gi|1036824808|ref|XP_017137855.1| PREDICTED: possible lysine-specific histone demethylase 1 [
>gi|1036824829|ref|XP_017137856.1| PREDICTED: possible lysine-specific histone demethylase 1 [
>gi|1036800896|ref|XP_017136536.1| PREDICTED: uncharacterized protein LOC108152044 [Drosophila
>gi|1036800861|ref|XP_017136534.1| PREDICTED: neurogenic protein mastermind isoform X3 [Drosop
>gi|1036800842|ref|XP_017136533.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosop
>gi|1036800804|ref|XP_017136531.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosop
>gi|1036800823|ref|XP_017136532.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosop
>gi|1036800880|ref|XP_017136535.1| PREDICTED: myb-like protein Q isoform X4 [Drosophila miranda
>gi|1036800936|ref|XP_017136538.1| PREDICTED: probable deoxycytidylate deaminase [Drosophila m
>gi|1036800917|ref|XP_017136537.1| PREDICTED: clathrin light chain [Drosophila miranda]
>gi|1036758923|ref|XP_017156807.1| PREDICTED: spore coat protein SP96-like [Drosophila miranda
>gi|1036845837|ref|XP_017139023.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036845800|ref|XP_017139021.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036845818|ref|XP_017139022.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036845856|ref|XP_017139024.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036845875|ref|XP_017139025.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036845892|ref|XP_017139027.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036845911|ref|XP_017139028.1| PREDICTED: serine/threonine-protein phosphatase rdgC isofor
>gi|1036786846|ref|XP_017135742.1| PREDICTED: uncharacterized protein LOC108151577 [Drosophila
>gi|1036786864|ref|XP_017135743.1| PREDICTED: uncharacterized protein C9orf117 homolog [Drosop
>gi|1036786917|ref|XP_017135746.1| PREDICTED: protein FRG1 homolog [Drosophila miranda]
>gi|1036786827|ref|XP_017135741.1| PREDICTED: tectonin beta-propeller repeat-containing protei
>gi|1036786932|ref|XP_017135747.1| PREDICTED: DCN1-like protein 5 isoform X1 [Drosophila miran
>gi|1036786968|ref|XP_017135749.1| PREDICTED: DCN1-like protein 5 isoform X1 [Drosophila miran
>gi|1036786901|ref|XP_017135745.1| PREDICTED: lysosome-associated membrane glycoprotein 5 isof
>gi|1036786883|ref|XP_017135744.1| PREDICTED: lysosome-associated membrane glycoprotein 5 isof
>gi|1036786987|ref|XP_017135750.1| PREDICTED: DCN1-like protein 4 isoform X2 [Drosophila miran
>gi|1036788166|ref|XP_017135816.1| PREDICTED: alpha-soluble NSF attachment protein [Drosophila
>gi|1036788111|ref|XP_017135813.1| PREDICTED: leukocyte elastase inhibitor [Drosophila miranda
>gi|1036788094|ref|XP_017135812.1| PREDICTED: retinoblastoma-binding protein 5 homolog [Drosop
>gi|1036788130|ref|XP_017135814.1| PREDICTED: eukaryotic peptide chain release factor subunit
>gi|1036788148|ref|XP_017135815.1| PREDICTED: eukaryotic peptide chain release factor subunit
>gi|1036788075|ref|XP_017135811.1| PREDICTED: cysteine sulfinic acid decarboxylase [Drosophila
>gi|1036822962|ref|XP_017137750.1| PREDICTED: DNA/RNA-binding protein KIN17 [Drosophila miranda
>gi|1036822981|ref|XP_017137751.1| PREDICTED: LOW QUALITY PROTEIN: vitellogenin-3 [Drosophila m
>gi|1036822944|ref|XP_017137749.1| PREDICTED: DNA primase large subunit [Drosophila miranda]
>gi|1036832401|ref|XP_017138284.1| PREDICTED: toll-like receptor 1 isoform X4 [Drosophila mira
>gi|1036832361|ref|XP_017138282.1| PREDICTED: toll-like receptor 1 isoform X2 [Drosophila mira
>gi|1036832382|ref|XP_017138283.1| PREDICTED: toll-like receptor 1 isoform X3 [Drosophila mira
>gi|1036832342|ref|XP_017138280.1| PREDICTED: toll-like receptor 1 isoform X1 [Drosophila mira
>gi|1036832419|ref|XP_017138285.1| PREDICTED: protein inturned [Drosophila miranda]
>gi|1036832456|ref|XP_017138288.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036832438|ref|XP_017138287.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036846436|ref|XP_017139057.1| PREDICTED: serine protease 3 [Drosophila miranda]
>gi|1036810377|ref|XP_017137060.1| PREDICTED: cyclin-T [Drosophila miranda]
>gi|1036828764|ref|XP_017138079.1| PREDICTED: ecdysone-induced protein 74EF isoform X2 [Drosop
>gi|1036828745|ref|XP_017138078.1| PREDICTED: ecdysone-induced protein 74EF isoform X1 [Drosop
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>gi|1036828781|ref|XP_017138080.1| PREDICTED: rho GTPase-activating protein gacII-like [Drosop
>gi|1036845048|ref|XP_017138985.1| PREDICTED: charged multivesicular body protein 5 [Drosophil
>gi|1036802166|ref|XP_017136610.1| PREDICTED: uncharacterized protein LOC108152084 isoform X1
>gi|1036802185|ref|XP_017136611.1| PREDICTED: AF4/FMR2 family member 4 isoform X2 [Drosophila m
>gi|1036802242|ref|XP_017136614.1| PREDICTED: protein melted isoform X2 [Drosophila miranda]
>gi|1036802204|ref|XP_017136612.1| PREDICTED: protein melted isoform X1 [Drosophila miranda]
>gi|1036802223|ref|XP_017136613.1| PREDICTED: protein melted isoform X1 [Drosophila miranda]
>gi|1036802263|ref|XP_017136616.1| PREDICTED: acyl-CoA-binding protein homolog [Drosophila mira
>gi|1036765957|ref|XP_017134549.1| PREDICTED: acyl-CoA-binding protein-like [Drosophila miranda
>gi|1036849714|ref|XP_017139246.1| PREDICTED: acyl-CoA-binding protein [Drosophila miranda]
>gi|1036849731|ref|XP_017139247.1| PREDICTED: acyl-CoA-binding protein homolog [Drosophila mira
>gi|1036849695|ref|XP_017139245.1| PREDICTED: Bardet-Biedl syndrome 1 protein [Drosophila miran
>gi|1036825035|ref|XP_017137866.1| PREDICTED: protein male-specific lethal-3 [Drosophila mirand
>gi|1036825020|ref|XP_017137865.1| PREDICTED: FH2 domain-containing protein 1 [Drosophila miran
>gi|1036758939|ref|XP_017156808.1| PREDICTED: uncharacterized protein LOC108165284 [Drosophila
>gi|1036768403|ref|XP_017134679.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036767849|ref|XP_017134649.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036767870|ref|XP_017134650.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036848986|ref|XP_017139202.1| PREDICTED: uncharacterized protein LOC108153615 isoform X1
>gi|1036849005|ref|XP_017139203.1| PREDICTED: uncharacterized protein LOC108153615 isoform X2
>gi|1036849045|ref|XP_017139205.1| PREDICTED: uncharacterized protein LOC108153616 isoform X2
>gi|1036849024|ref|XP_017139204.1| PREDICTED: uncharacterized protein LOC108153616 isoform X1
>gi|1036848967|ref|XP_017139201.1| PREDICTED: GTP-binding protein Rhes [Drosophila miranda]
>gi|1036838898|ref|XP_017138661.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X9 [Drosop
>gi|1036838861|ref|XP_017138658.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X7 [Drosop
>gi|1036838880|ref|XP_017138659.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X8 [Drosop
>gi|1036838842|ref|XP_017138657.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X6 [Drosop
>gi|1036838823|ref|XP_017138656.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X5 [Drosop
>gi|1036838806|ref|XP_017138655.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X4 [Drosop
>gi|1036838787|ref|XP_017138654.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X3 [Drosop
>gi|1036838768|ref|XP_017138653.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X2 [Drosop
>gi|1036838730|ref|XP_017138651.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X1 [Drosop
>gi|1036838749|ref|XP_017138652.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X1 [Drosop
>gi|1036838971|ref|XP_017138665.1| PREDICTED: collagen alpha-1(XV) chain isoform X13 [Drosophil
>gi|1036838953|ref|XP_017138664.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X12 [Drosop
>gi|1036838934|ref|XP_017138663.1| PREDICTED: collagen alpha-1(XV) chain isoform X11 [Drosophil
>gi|1036838915|ref|XP_017138662.1| PREDICTED: collagen alpha-1(XVIII) chain isoform X10 [Drosop
>gi|1036838990|ref|XP_017138666.1| PREDICTED: uncharacterized protein LOC108153289 [Drosophila
>gi|1036812163|ref|XP_017137156.1| PREDICTED: differentially expressed in FDCP 8 homolog [Drosc
>gi|1036812123|ref|XP_017137154.1| PREDICTED: protein KRI1 homolog [Drosophila miranda]
>gi|1036812142|ref|XP_017137155.1| PREDICTED: protein KRI1 homolog [Drosophila miranda]
>gi|1036812201|ref|XP_017137159.1| PREDICTED: autophagy protein 12-like [Drosophila miranda]
>gi|1036812182|ref|XP_017137158.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1036774472|ref|XP_017135027.1| PREDICTED: rho GTPase-activating protein 68F isoform X1 [Dro
>gi|1036774491|ref|XP_017135028.1| PREDICTED: rho GTPase-activating protein 68F isoform X1 [Dro
>gi|1036774510|ref|XP_017135029.1| PREDICTED: rho GTPase-activating protein 68F isoform X2 [Dro
>gi|1036774432|ref|XP_017135025.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1036774340|ref|XP_017135019.1| PREDICTED: neurexin-4 isoform X3 [Drosophila miranda]
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>gi|1036774359|ref|XP_017135020.1| PREDICTED: neurexin-4 isoform X4 [Drosophila miranda]
>gi|1036774302|ref|XP_017135017.1| PREDICTED: neurexin-4 isoform X1 [Drosophila miranda]
>gi|1036774321|ref|XP_017135018.1| PREDICTED: neurexin-4 isoform X2 [Drosophila miranda]
>gi|1036774565|ref|XP_017135033.1| PREDICTED: L-aminoadipate-semialdehyde dehydrogenase-phospho
>gi|1036774597|ref|XP_017135036.1| PREDICTED: uncharacterized protein LOC108151126 [Drosophila
>gi|1036774451|ref|XP_017135026.1| PREDICTED: opsin, ultraviolet-sensitive [Drosophila miranda]
>gi|1036774547|ref|XP_017135032.1| PREDICTED: carbonic anhydrase 2 [Drosophila miranda]
>gi|1036774583|ref|XP_017135034.1| PREDICTED: inositol oxygenase [Drosophila miranda]
>gi|1036774414|ref|XP_017135024.1| PREDICTED: esterase-5C [Drosophila miranda]
>gi|1036774397|ref|XP_017135022.1| PREDICTED: esterase-5B [Drosophila miranda]
>gi|1036774378|ref|XP_017135021.1| PREDICTED: esterase-5A isoform X5 [Drosophila miranda]
>gi|1036774529|ref|XP_017135030.1| PREDICTED: THO complex subunit 6 [Drosophila miranda]
>gi|1036851872|ref|XP_017139375.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036827881|ref|XP_017138030.1| PREDICTED: adenylate kinase isoenzyme 1-like isoform X2 [Dro
>gi|1036827863|ref|XP_017138029.1| PREDICTED: adenylate kinase isoenzyme 1-like isoform X1 [Dro
>gi|1036827904|ref|XP_017138032.1| PREDICTED: adenylate kinase isoenzyme 1-like [Drosophila mi
>gi|1036827768|ref|XP_017138023.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827787|ref|XP_017138024.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827806|ref|XP_017138025.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827825|ref|XP_017138027.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036827844|ref|XP_017138028.1| PREDICTED: uncharacterized protein LOC108152894 [Drosophila
>gi|1036804750|ref|XP_017136753.1| PREDICTED: probable protein S-acyltransferase 4 [Drosophila
>gi|1036804768|ref|XP_017136754.1| PREDICTED: dnaJ protein ERDJ2A-like [Drosophila miranda]
>gi|1036830663|ref|XP_017138189.1| PREDICTED: uncharacterized protein LOC108152974 [Drosophila
>gi|1036830684|ref|XP_017138190.1| PREDICTED: delta-aminolevulinic acid dehydratase [Drosophila
>gi|1036824752|ref|XP_017137853.1| PREDICTED: spermine synthase isoform X1 [Drosophila miranda]
>gi|1036824771|ref|XP_017137854.1| PREDICTED: spermine synthase isoform X2 [Drosophila miranda]
>gi|1036840223|ref|XP_017138729.1| PREDICTED: 72 kDa inositol polyphosphate 5-phosphatase [Dros
>gi|1036764987|ref|XP_017157145.1| PREDICTED: LIM homeobox transcription factor 1-beta [Drosoph
>gi|1036849511|ref|XP_017139233.1| PREDICTED: LIM homeobox transcription factor 1-beta [Drosoph
>gi|1036842423|ref|XP_017138844.1| PREDICTED: uncharacterized protein LOC108153383 [Drosophila
>gi|1036842441|ref|XP_017138845.1| PREDICTED: uncharacterized protein LOC108153383 [Drosophila
>gi|1036819818|ref|XP_017137582.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso
>gi|1036819892|ref|XP_017137585.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso
>gi|1036819873|ref|XP_017137584.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso
>gi|1036819911|ref|XP_017137586.1| PREDICTED: calcium-transporting ATPase type 2C member 1 iso
>gi|1036827671|ref|XP_017138017.1| PREDICTED: ras-GEF domain-containing family member 1B-B [Dro
>gi|1036841699|ref|XP_017138815.1| PREDICTED: growth arrest and DNA damage-inducible proteins-i
>gi|1036841685|ref|XP_017138814.1| PREDICTED: uncharacterized protein LOC108153359 [Drosophila
>gi|1036842894|ref|XP_017138872.1| PREDICTED: neuropeptides capa receptor isoform X1 [Drosophil
>gi|1036842913|ref|XP_017138873.1| PREDICTED: neuropeptides capa receptor isoform X1 [Drosophil
>gi|1036842950|ref|XP_017138876.1| PREDICTED: neuropeptides capa receptor isoform X3 [Drosophil
>gi|1036842931|ref|XP_017138875.1| PREDICTED: neuropeptides capa receptor isoform X2 [Drosophil
>gi|1036851128|ref|XP_017139329.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036817942|ref|XP_017137479.1| PREDICTED: hemicentin-1 isoform X7 [Drosophila miranda]
>gi|1036817965|ref|XP_017137480.1| PREDICTED: uncharacterized protein LOC108152559 isoform X8
>gi|1036817923|ref|XP_017137478.1| PREDICTED: hemicentin-1 isoform X6 [Drosophila miranda]
>gi|1036817906|ref|XP_017137477.1| PREDICTED: uncharacterized protein LOC108152559 isoform X5
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>gi|1036817871|ref|XP_017137475.1| PREDICTED: uncharacterized protein LOC108152559 isoform X3
>gi|1036817855|ref|XP_017137473.1| PREDICTED: uncharacterized protein LOC108152559 isoform X2
>gi|1036817840|ref|XP_017137472.1| PREDICTED: uncharacterized protein LOC108152559 isoform X1
>gi|1036817888|ref|XP_017137476.1| PREDICTED: uncharacterized protein LOC108152559 isoform X4
>gi|1036758957|ref|XP_017156809.1| PREDICTED: muscle M-line assembly protein unc-89-like [Droso
>gi|1036817984|ref|XP_017137481.1| PREDICTED: uncharacterized protein LOC108152560 [Drosophila
>gi|1036827432|ref|XP_017138003.1| PREDICTED: uncharacterized protein LOC108152881 [Drosophila
>gi|1036851309|ref|XP_017139340.1| PREDICTED: cathepsin L1 [Drosophila miranda]
>gi|1036846093|ref|XP_017139038.1| PREDICTED: uncharacterized protein LOC108153498 [Drosophila
>gi|1036767596|ref|XP_017134638.1| PREDICTED: uncharacterized protein LOC108150840 [Drosophila
>gi|1036782830|ref|XP_017135511.1| PREDICTED: cationic amino acid transporter 3 isoform X1 [Dro
>gi|1036782849|ref|XP_017135512.1| PREDICTED: cationic amino acid transporter 3 isoform X2 [Dro
>gi|1036782867|ref|XP_017135513.1| PREDICTED: cationic amino acid transporter 3 isoform X2 [Dro
>gi|1036782884|ref|XP_017135514.1| PREDICTED: cationic amino acid transporter 3 isoform X2 [Dro
>gi|1036782902|ref|XP_017135515.1| PREDICTED: general transcription factor IIH subunit 2 [Droso
>gi|1036782795|ref|XP_017135509.1| PREDICTED: uncharacterized protein LOC108151428 [Drosophila
>gi|1036782813|ref|XP_017135510.1| PREDICTED: uncharacterized protein LOC108151428 [Drosophila
>gi|1036782921|ref|XP_017135516.1| PREDICTED: putative methyltransferase NSUN6 [Drosophila mira
>gi|1036782938|ref|XP_017135517.1| PREDICTED: ADP-ribosylation factor 1 [Drosophila miranda]
>gi|1036782957|ref|XP_017135518.1| PREDICTED: ADP-ribosylation factor 1 [Drosophila miranda]
>gi|1036831611|ref|XP_017138243.1| PREDICTED: ER membrane protein complex subunit 4 [Drosophila
>gi|1036831648|ref|XP_017138246.1| PREDICTED: uncharacterized protein LOC108153010 isoform X2
>gi|1036831629|ref|XP_017138245.1| PREDICTED: uncharacterized protein LOC108153010 isoform X1
>gi|1036831667|ref|XP_017138247.1| PREDICTED: single-pass membrane and coiled-coil domain-conta
>gi|1036766868|ref|XP_017134598.1| PREDICTED: uncharacterized protein LOC108150806 [Drosophila
>gi|1036758975|ref|XP_017156810.1| PREDICTED: oleosin-B6 [Drosophila miranda]
>gi|1036850074|ref|XP_017139269.1| PREDICTED: aspartic and glutamic acid-rich protein [Drosophi
>gi|1036850093|ref|XP_017139270.1| PREDICTED: uncharacterized protein LOC108153667 [Drosophila
>gi|1036824174|ref|XP_017137819.1| PREDICTED: myeloid zinc finger 1 isoform X1 [Drosophila mira
>gi|1036824195|ref|XP_017137820.1| PREDICTED: myeloid zinc finger 1 isoform X2 [Drosophila mira
>gi|1036824214|ref|XP_017137821.1| PREDICTED: myeloid zinc finger 1 isoform X3 [Drosophila mira
>gi|1036824237|ref|XP_017137822.1| PREDICTED: myeloid zinc finger 1 isoform X4 [Drosophila mira
>gi|1036824256|ref|XP_017137823.1| PREDICTED: zinc finger protein 768 isoform X5 [Drosophila mi
>gi|1036824275|ref|XP_017137824.1| PREDICTED: uncharacterized protein LOC108152763 [Drosophila
>gi|1036763048|ref|XP_017157049.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm2 [Drosoph
>gi|1036846165|ref|XP_017139042.1| PREDICTED: gonadotropin-releasing hormone receptor [Drosophi
>gi|1036831227|ref|XP_017138221.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1036831208|ref|XP_017138220.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1036831172|ref|XP_017138218.1| PREDICTED: solute carrier family 12 member 2 isoform X1 [Dro
>gi|1036831191|ref|XP_017138219.1| PREDICTED: solute carrier family 12 member 2 isoform X1 [Dro
>gi|1036817463|ref|XP_017137450.1| PREDICTED: RNA polymerase II-associated protein 1 [Drosophil
>gi|1036817519|ref|XP_017137454.1| PREDICTED: general odorant-binding protein 69a [Drosophila r
>gi|1036817481|ref|XP_017137452.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosoph
>gi|1036817501|ref|XP_017137453.1| PREDICTED: SET and MYND domain-containing protein 4 [Drosoph
>gi|1036826387|ref|XP_017137941.1| PREDICTED: calcium-binding mitochondrial carrier protein SCa
>gi|1036826406|ref|XP_017137942.1| PREDICTED: calcium-binding mitochondrial carrier protein SCa
>gi|1036826424|ref|XP_017137943.1| PREDICTED: probable calcium-binding mitochondrial carrier Cl
>gi|1036826442|ref|XP_017137944.1| PREDICTED: probable calcium-binding mitochondrial carrier Cl
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>gi|1036836023|ref|XP_017138485.1| PREDICTED: ADP-ribosylation factor GTPase-activating protei
>gi|1036836004|ref|XP_017138484.1| PREDICTED: ADP-ribosylation factor GTPase-activating protei
>gi|1036847661|ref|XP_017139127.1| PREDICTED: paired box protein Pax-6 isoform X1 [Drosophila m
>gi|1036847680|ref|XP_017139128.1| PREDICTED: paired box protein Pax-1 isoform X2 [Drosophila m
>gi|1036846398|ref|XP_017139055.1| PREDICTED: paired box protein Pax-6 [Drosophila miranda]
>gi|1036768499|ref|XP_017134685.1| PREDICTED: cuticle protein 38-like [Drosophila miranda]
>gi|1036768575|ref|XP_017134689.1| PREDICTED: uncharacterized protein LOC108150890 [Drosophila
>gi|1036766481|ref|XP_017134577.1| PREDICTED: uncharacterized protein LOC108150787 [Drosophila
>gi|1036828127|ref|XP_017138043.1| PREDICTED: protein odr-4 homolog [Drosophila miranda]
>gi|1036828091|ref|XP_017138040.1| PREDICTED: uncharacterized protein LOC108152905 [Drosophila
>gi|1036828110|ref|XP_017138041.1| PREDICTED: uncharacterized protein LOC108152905 [Drosophila
>gi|1036847085|ref|XP_017139095.1| PREDICTED: uncharacterized protein LOC108153540 isoform X1
>gi|1036847104|ref|XP_017139096.1| PREDICTED: uncharacterized protein LOC108153540 isoform X2
>gi|1036848931|ref|XP_017139199.1| PREDICTED: clavesin-1 [Drosophila miranda]
>gi|1036780551|ref|XP_017135379.1| PREDICTED: 1,5-anhydro-D-fructose reductase isoform X2 [Dros
>gi|1036780532|ref|XP_017135378.1| PREDICTED: 1,5-anhydro-D-fructose reductase isoform X1 [Dros
>gi|1036780570|ref|XP_017135380.1| PREDICTED: aldose reductase isoform X3 [Drosophila miranda]
>gi|1036780589|ref|XP_017135381.1| PREDICTED: kxDL motif-containing protein CG10681 isoform X1
>gi|1036780607|ref|XP_017135383.1| PREDICTED: kxDL motif-containing protein CG10681 isoform X2
>gi|1036780514|ref|XP_017135377.1| PREDICTED: protein TSSC1 [Drosophila miranda]
>gi|1036780625|ref|XP_017135384.1| PREDICTED: ubiquitin-conjugating enzyme E2 C [Drosophila mi
>gi|1036780498|ref|XP_017135376.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1036780479|ref|XP_017135375.1| PREDICTED: citron Rho-interacting kinase isoform X2 [Drosop
>gi|1036780461|ref|XP_017135374.1| PREDICTED: citron Rho-interacting kinase isoform X1 [Drosop
>gi|1036781203|ref|XP_017135419.1| PREDICTED: protein LSM14 homolog B-A isoform X1 [Drosophila
>gi|1036781222|ref|XP_017135420.1| PREDICTED: protein LSM14 homolog B-A isoform X2 [Drosophila
>gi|1036781184|ref|XP_017135418.1| PREDICTED: histone acetyltransferase KAT2A [Drosophila mira
>gi|1036781278|ref|XP_017135423.1| PREDICTED: uncharacterized protein LOC108151372 [Drosophila
>gi|1036781259|ref|XP_017135422.1| PREDICTED: eukaryotic translation initiation factor 2 subun
>gi|1036781240|ref|XP_017135421.1| PREDICTED: phosphoacetylglucosamine mutase [Drosophila mira
>gi|1036785067|ref|XP_017135642.1| PREDICTED: melanotransferrin [Drosophila miranda]
>gi|1036785159|ref|XP_017135647.1| PREDICTED: uncharacterized protein LOC108151514 [Drosophila
>gi|1036785141|ref|XP_017135646.1| PREDICTED: probable phosphomannomutase [Drosophila miranda]
>gi|1036785084|ref|XP_017135643.1| PREDICTED: kelch domain-containing protein 4 [Drosophila mi
>gi|1036785103|ref|XP_017135644.1| PREDICTED: uncharacterized protein LOC108151510 [Drosophila
>gi|1036785122|ref|XP_017135645.1| PREDICTED: ankyrin repeat domain-containing protein SOWAHA-
>gi|1036843319|ref|XP_017138896.1| PREDICTED: homeobox protein araucan [Drosophila miranda]
>gi|1036837311|ref|XP_017138561.1| PREDICTED: homeobox protein caupolican [Drosophila miranda]
>gi|1036832127|ref|XP_017138271.1| PREDICTED: homeobox protein caupolican [Drosophila miranda]
>gi|1036834640|ref|XP_017138403.1| PREDICTED: protein ABHD18 [Drosophila miranda]
>gi|1036812315|ref|XP_017137165.1| PREDICTED: tyrosine-protein phosphatase 69D [Drosophila mira
>gi|1036796922|ref|XP_017136311.1| PREDICTED: uncharacterized protein LOC108151911 [Drosophila
>gi|1036796937|ref|XP_017136312.1| PREDICTED: probable small nuclear ribonucleoprotein Sm D1 [
>gi|1036796871|ref|XP_017136308.1| PREDICTED: kinesin light chain [Drosophila miranda]
>gi|1036796888|ref|XP_017136309.1| PREDICTED: kinesin light chain [Drosophila miranda]
>gi|1036796855|ref|XP_017136307.1| PREDICTED: ankyrin repeat domain-containing protein 11 [Dros
>gi|1036796905|ref|XP_017136310.1| PREDICTED: hsp70 nucleotide exchange factor fes1 [Drosophila
>gi|1036781387|ref|XP_017135430.1| PREDICTED: vacuolar protein sorting-associated protein 13D
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>gi|1036781425|ref|XP_017135432.1| PREDICTED: huntingtin-interacting protein 1 isoform X2 [Dros
>gi|1036781406|ref|XP_017135431.1| PREDICTED: huntingtin-interacting protein 1 isoform X1 [Dros
>gi|1036758994|ref|XP_017156811.1| PREDICTED: ATP-binding cassette sub-family A member 17-like
>gi|1036759010|ref|XP_017156812.1| PREDICTED: uncharacterized protein LOC108165288 [Drosophila
>gi|1036759029|ref|XP_017156813.1| PREDICTED: protein SDA1 homolog [Drosophila miranda]
>gi|1036828835|ref|XP_017138082.1| PREDICTED: calcium channel flower isoform X1 [Drosophila mir
>gi|1036828853|ref|XP_017138083.1| PREDICTED: calcium channel flower isoform X2 [Drosophila mir
>gi|1036799054|ref|XP_017136432.1| PREDICTED: zinc finger protein 140 [Drosophila miranda]
>gi|1036799035|ref|XP_017136431.1| PREDICTED: mRNA-decapping enzyme subunit 2 [Drosophila mirar
>gi|1036825673|ref|XP_017137899.1| PREDICTED: kelch-like protein diablo [Drosophila miranda]
>gi|1036834122|ref|XP_017138375.1| PREDICTED: uncharacterized protein C15orf41 homolog [Drosopl
>gi|1036834138|ref|XP_017138376.1| PREDICTED: zinc finger protein 706-like [Drosophila miranda]
>gi|1036834157|ref|XP_017138377.1| PREDICTED: zinc finger protein 706-like [Drosophila miranda]
>gi|1036847586|ref|XP_017139122.1| PREDICTED: uncharacterized protein LOC108153561 isoform X1
>gi|1036847605|ref|XP_017139123.1| PREDICTED: uncharacterized protein LOC108153561 isoform X1
>gi|1036847622|ref|XP_017139124.1| PREDICTED: uncharacterized protein LOC108153561 isoform X2
>gi|1036840742|ref|XP_017138758.1| PREDICTED: uncharacterized protein LOC108153334 isoform X11
>gi|1036840704|ref|XP_017138756.1| PREDICTED: uncharacterized protein LOC108153334 isoform X9
>gi|1036840614|ref|XP_017138750.1| PREDICTED: uncharacterized protein LOC108153334 isoform X4
>gi|1036840668|ref|XP_017138754.1| PREDICTED: uncharacterized protein LOC108153334 isoform X7
>gi|1036840723|ref|XP_017138757.1| PREDICTED: uncharacterized protein LOC108153334 isoform X10
>gi|1036840649|ref|XP_017138753.1| PREDICTED: uncharacterized protein LOC108153334 isoform X6
>gi|1036840481|ref|XP_017138743.1| PREDICTED: uncharacterized protein LOC108153334 isoform X2
>gi|1036840779|ref|XP_017138760.1| PREDICTED: gamma-aminobutyric acid receptor subunit alpha-3
>gi|1036840687|ref|XP_017138755.1| PREDICTED: uncharacterized protein LOC108153334 isoform X8
>gi|1036840760|ref|XP_017138759.1| PREDICTED: gamma-aminobutyric acid receptor subunit alpha-3
>gi|1036840633|ref|XP_017138752.1| PREDICTED: uncharacterized protein LOC108153334 isoform X5
>gi|1036840595|ref|XP_017138749.1| PREDICTED: uncharacterized protein LOC108153334 isoform X3
>gi|1036840444|ref|XP_017138741.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840462|ref|XP_017138742.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840500|ref|XP_017138744.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840517|ref|XP_017138745.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840536|ref|XP_017138746.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840557|ref|XP_017138747.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036840576|ref|XP_017138748.1| PREDICTED: uncharacterized protein LOC108153334 isoform X1
>gi|1036844916|ref|XP_017138978.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X6
>gi|1036844879|ref|XP_017138976.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X4
>gi|1036844897|ref|XP_017138977.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X5
>gi|1036844858|ref|XP_017138975.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X3
>gi|1036844837|ref|XP_017138974.1| PREDICTED: serine/threonine-protein kinase BRSK2 isoform X2
>gi|1036844784|ref|XP_017138971.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X1
>gi|1036844803|ref|XP_017138972.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X1
>gi|1036844819|ref|XP_017138973.1| PREDICTED: serine/threonine-protein kinase BRSK1 isoform X1
>gi|1036845067|ref|XP_017138986.1| PREDICTED: protein kinase DC2 isoform X1 [Drosophila miranda
>gi|1036845082|ref|XP_017138987.1| PREDICTED: protein kinase DC2 isoform X1 [Drosophila miranda
>gi|1036845099|ref|XP_017138988.1| PREDICTED: protein kinase DC2 isoform X2 [Drosophila miranda
>gi|1036845390|ref|XP_017139006.1| PREDICTED: group XIIA secretory phospholipase A2 [Drosophila
>gi|1036835833|ref|XP_017138473.1| PREDICTED: E3 ubiquitin-protein ligase NRDP1 [Drosophila mir
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>gi|1036835852|ref|XP_017138474.1| PREDICTED: uncharacterized protein LOC108153177 [Drosophila
>gi|1036799217|ref|XP_017136442.1| PREDICTED: putative U5 small nuclear ribonucleoprotein 200 k
>gi|1036797300|ref|XP_017136333.1| PREDICTED: chromosomal serine/threonine-protein kinase JIL-1
>gi|1036797319|ref|XP_017136334.1| PREDICTED: chromosomal serine/threonine-protein kinase JIL-1
>gi|1036834679|ref|XP_017138405.1| PREDICTED: iodotyrosine deiodinase 1-like [Drosophila miran
>gi|1036834661|ref|XP_017138404.1| PREDICTED: uncharacterized protein LOC108153126 [Drosophila
>gi|1036834714|ref|XP_017138407.1| PREDICTED: uncharacterized protein LOC108153129 [Drosophila
>gi|1036834696|ref|XP_017138406.1| PREDICTED: CCAAT/enhancer-binding protein gamma [Drosophila
>gi|1036841595|ref|XP_017138809.1| PREDICTED: nedd8-activating enzyme E1 regulatory subunit [Dr
>gi|1036759048|ref|XP_017156814.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1036813031|ref|XP_017137198.1| PREDICTED: uncharacterized protein LOC108152406 [Drosophila
>gi|1036813052|ref|XP_017137199.1| PREDICTED: uncharacterized protein LOC108152406 [Drosophila
>gi|1036813072|ref|XP_017137200.1| PREDICTED: uncharacterized protein LOC108152406 [Drosophila
>gi|1036812791|ref|XP_017137187.1| PREDICTED: adenosine deaminase CECR1 [Drosophila miranda]
>gi|1036812810|ref|XP_017137188.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812829|ref|XP_017137189.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812848|ref|XP_017137190.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812867|ref|XP_017137191.1| PREDICTED: adenosine deaminase CECR1-A [Drosophila miranda]
>gi|1036812884|ref|XP_017137193.1| PREDICTED: adenosine deaminase CECR1 [Drosophila miranda]
>gi|1036812901|ref|XP_017137194.1| PREDICTED: basic-leucine zipper transcription factor A [Dros
>gi|1036812918|ref|XP_017137195.1| PREDICTED: basic-leucine zipper transcription factor A [Dros
>gi|1036813012|ref|XP_017137197.1| PREDICTED: spermine oxidase-like [Drosophila miranda]
>gi|1036812937|ref|XP_017137196.1| PREDICTED: peroxisomal N(1)-acetyl-spermine/spermidine oxida
>gi|1036850379|ref|XP_017139285.1| PREDICTED: uncharacterized protein LOC108153680 [Drosophila
>gi|1036823821|ref|XP_017137799.1| PREDICTED: nuclear transcription factor Y subunit beta [Dros
>gi|1036847217|ref|XP_017139102.1| PREDICTED: 15 kDa selenoprotein [Drosophila miranda]
>gi|1036783551|ref|XP_017135555.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036783570|ref|XP_017135556.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036783591|ref|XP_017135557.1| PREDICTED: protein spitz [Drosophila miranda]
>gi|1036783513|ref|XP_017135553.1| PREDICTED: NEDD4 family-interacting protein 1 [Drosophila mi
>gi|1036783532|ref|XP_017135554.1| PREDICTED: NEDD4 family-interacting protein 1 [Drosophila mi
>gi|1036783435|ref|XP_017135548.1| PREDICTED: zinc finger protein on ecdysone puffs isoform X1
>gi|1036783454|ref|XP_017135549.1| PREDICTED: zinc finger protein on ecdysone puffs isoform X1
>gi|1036783473|ref|XP_017135551.1| PREDICTED: zinc finger protein on ecdysone puffs isoform X2
>gi|1036783494|ref|XP_017135552.1| PREDICTED: uncharacterized protein LOC108151448 [Drosophila
>gi|1036849083|ref|XP_017139208.1| PREDICTED: prostaglandin D2 receptor [Drosophila miranda]
>gi|1036820663|ref|XP_017137622.1| PREDICTED: integral membrane protein GPR155 [Drosophila mira
>gi|1036766046|ref|XP_017134554.1| PREDICTED: RNA-binding motif protein, X-linked 2 isoform X1
>gi|1036766064|ref|XP_017134556.1| PREDICTED: RNA-binding motif protein, X-linked 2 isoform X2
>gi|1036841110|ref|XP_017138781.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841091|ref|XP_017138780.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840981|ref|XP_017138773.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841055|ref|XP_017138778.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036840943|ref|XP_017138771.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841072|ref|XP_017138779.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
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>gi|1036840909|ref|XP_017138768.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036841017|ref|XP_017138775.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
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>gi|1036840926|ref|XP_017138769.1| PREDICTED: Down syndrome cell adhesion molecule-like protein
>gi|1036840962|ref|XP_017138772.1| PREDICTED: Down syndrome cell adhesion molecule-like protein
>gi|1036840998|ref|XP_017138774.1| PREDICTED: Down syndrome cell adhesion molecule-like protein
>gi|1036840888|ref|XP_017138767.1| PREDICTED: Down syndrome cell adhesion molecule-like protein
>gi|1036841128|ref|XP_017138782.1| PREDICTED: Down syndrome cell adhesion molecule-like protein
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>gi|1036841147|ref|XP_017138783.1| PREDICTED: Down syndrome cell adhesion molecule-like protein
>gi|1036813396|ref|XP_017137219.1| PREDICTED: uncharacterized protein LOC108152417 [Drosophila
>gi|1036813356|ref|XP_017137217.1| PREDICTED: uncharacterized protein LOC108152415 isoform X1
>gi|1036813375|ref|XP_017137218.1| PREDICTED: uncharacterized protein LOC108152415 isoform X2
>gi|1036813314|ref|XP_017137215.1| PREDICTED: folliculin [Drosophila miranda]
>gi|1036813430|ref|XP_017137221.1| PREDICTED: 60S ribosomal protein L18 [Drosophila miranda]
>gi|1036813413|ref|XP_017137220.1| PREDICTED: 39S ribosomal protein L50, mitochondrial [Drosoph
>gi|1036813337|ref|XP_017137216.1| PREDICTED: nuclear receptor coactivator 5 [Drosophila miran
>gi|1036770311|ref|XP_017134787.1| PREDICTED: uncharacterized protein LOC108150964 [Drosophila
>gi|1036770197|ref|XP_017134780.1| PREDICTED: cell division cycle protein 27 homolog [Drosophi
>gi|1036770273|ref|XP_017134785.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036770159|ref|XP_017134778.1| PREDICTED: protein KIAA0100 [Drosophila miranda]
>gi|1036770178|ref|XP_017134779.1| PREDICTED: trafficking protein particle complex subunit 11
>gi|1036770237|ref|XP_017134782.1| PREDICTED: trypsin alpha [Drosophila miranda]
>gi|1036770216|ref|XP_017134781.1| PREDICTED: trypsin beta-like [Drosophila miranda]
>gi|1036758083|ref|XP_017156756.1| PREDICTED: transmembrane protease serine 9 [Drosophila mira
>gi|1036770255|ref|XP_017134783.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1036770292|ref|XP_017134786.1| PREDICTED: stress-associated endoplasmic reticulum protein 2
>gi|1036770349|ref|XP_017134790.1| PREDICTED: UPF0235 protein C15orf40 homolog [Drosophila mir
>gi|1036770330|ref|XP_017134788.1| PREDICTED: heat shock protein 83 [Drosophila miranda]
>gi|1036794352|ref|XP_017136167.1| PREDICTED: uncharacterized protein LOC108151819 [Drosophila
>gi|1036759067|ref|XP_017156815.1| PREDICTED: uncharacterized protein LOC108165292 [Drosophila
>gi|1036794387|ref|XP_017136169.1| PREDICTED: cell wall protein IFF6 [Drosophila miranda]
>gi|1036794371|ref|XP_017136168.1| PREDICTED: uncharacterized protein LOC108151820 [Drosophila
>gi|1036794317|ref|XP_017136165.1| PREDICTED: daple-like protein isoform X2 [Drosophila miranda
>gi|1036794302|ref|XP_017136164.1| PREDICTED: girdin isoform X1 [Drosophila miranda]
>gi|1036794333|ref|XP_017136166.1| PREDICTED: girdin isoform X3 [Drosophila miranda]
>gi|1036829922|ref|XP_017138144.1| PREDICTED: titin homolog [Drosophila miranda]
>gi|1036829960|ref|XP_017138146.1| PREDICTED: uncharacterized protein LOC108152957 [Drosophila
>gi|1036829978|ref|XP_017138147.1| PREDICTED: uncharacterized protein LOC108152958 [Drosophila
>gi|1036829941|ref|XP_017138145.1| PREDICTED: tRNA (guanine(37)-N1)-methyltransferase [Drosophi
>gi|1036843652|ref|XP_017138917.1| PREDICTED: short/branched chain specific acyl-CoA dehydroge
>gi|1036843670|ref|XP_017138918.1| PREDICTED: ninjurin-1 [Drosophila miranda]
>gi|1036782545|ref|XP_017135493.1| PREDICTED: long-chain-fatty-acid--CoA ligase 5 isoform X1 [
>gi|1036782564|ref|XP_017135494.1| PREDICTED: long-chain-fatty-acid--CoA ligase 5 isoform X1 [
>gi|1036782582|ref|XP_017135495.1| PREDICTED: long-chain-fatty-acid--CoA ligase 5 isoform X2 [
>gi|1036782526|ref|XP_017135492.1| PREDICTED: UDP-glucose:glycoprotein glucosyltransferase [Dro
>gi|1036782600|ref|XP_017135497.1| PREDICTED: cell cycle checkpoint control protein RAD9A [Dros
>gi|1036782657|ref|XP_017135500.1| PREDICTED: biogenesis of lysosome-related organelles comple
>gi|1036782674|ref|XP_017135501.1| PREDICTED: glutaredoxin-C4 [Drosophila miranda]
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>gi|1036782638|ref|XP_017135499.1| PREDICTED: dysbindin protein homolog [Drosophila miranda]
>gi|1036782619|ref|XP_017135498.1| PREDICTED: ADP-ribosylation factor-like protein 6-interacti
>gi|1036759086|ref|XP_017156816.1| PREDICTED: transmembrane protease serine 9 [Drosophila mira
>gi|1036820924|ref|XP_017137637.1| PREDICTED: protein I'm not dead yet isoform X1 [Drosophila
>gi|1036820943|ref|XP_017137638.1| PREDICTED: protein I'm not dead yet isoform X2 [Drosophila
>gi|1036847497|ref|XP_017139119.1| PREDICTED: uncharacterized protein LOC108153557 isoform X1
>gi|1036847514|ref|XP_017139120.1| PREDICTED: uncharacterized protein LOC108153557 isoform X2
>gi|1036832533|ref|XP_017138292.1| PREDICTED: catalase isoform X1 [Drosophila miranda]
>gi|1036832552|ref|XP_017138293.1| PREDICTED: catalase isoform X2 [Drosophila miranda]
>gi|1036844356|ref|XP_017138953.1| PREDICTED: uncharacterized protein LOC108153447 [Drosophila
>gi|1036844374|ref|XP_017138954.1| PREDICTED: uncharacterized protein LOC108153447 [Drosophila
>gi|1036808939|ref|XP_017136984.1| PREDICTED: uncharacterized protein LOC108152274 [Drosophila
>gi|1036808919|ref|XP_017136983.1| PREDICTED: uncharacterized protein LOC108152273 [Drosophila
>gi|1036826370|ref|XP_017137940.1| PREDICTED: nuclear hormone receptor FTZ-F1 [Drosophila mira
>gi|1036820721|ref|XP_017137625.1| PREDICTED: ubiquitin-like-conjugating enzyme ATG3 [Drosophi
>gi|1036820700|ref|XP_017137624.1| PREDICTED: nuclear fragile X mental retardation-interacting
>gi|1036820741|ref|XP_017137626.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036763115|ref|XP_017157050.1| PREDICTED: protein terminus [Drosophila miranda]
>gi|1036764894|ref|XP_017157142.1| PREDICTED: M protein, serotype 2.1-like [Drosophila miranda
>gi|1036847760|ref|XP_017139132.1| PREDICTED: uncharacterized protein LOC108153567 [Drosophila
>gi|1036849122|ref|XP_017139210.1| PREDICTED: uncharacterized protein LOC108153620 isoform X1
>gi|1036849139|ref|XP_017139211.1| PREDICTED: uncharacterized protein LOC108153620 isoform X1
>gi|1036849155|ref|XP_017139212.1| PREDICTED: uncharacterized protein LOC108153620 isoform X2
>gi|1036851689|ref|XP_017139363.1| PREDICTED: uncharacterized protein LOC108153745 [Drosophila
>gi|1036847869|ref|XP_017139138.1| PREDICTED: uncharacterized protein LOC108153571 [Drosophila
>gi|1036759105|ref|XP_017156817.1| PREDICTED: putative odorant receptor 69a [Drosophila miranda
>gi|1036759126|ref|XP_017156819.1| PREDICTED: putative odorant receptor 69a [Drosophila miranda
>gi|1036765869|ref|XP_017134544.1| PREDICTED: malate dehydrogenase, mitochondrial [Drosophila
>gi|1036764947|ref|XP_017157143.1| PREDICTED: malate dehydrogenase, mitochondrial [Drosophila
>gi|1036841227|ref|XP_017138787.1| PREDICTED: cullin homolog 1 [Drosophila miranda]
>gi|1036818364|ref|XP_017137502.1| PREDICTED: 10 kDa heat shock protein, mitochondrial [Drosop
>gi|1036818346|ref|XP_017137501.1| PREDICTED: 39S ribosomal protein L20, mitochondrial [Drosop
>gi|1036818328|ref|XP_017137500.1| PREDICTED: protein-S-isoprenylcysteine O-methyltransferase
>gi|1036818309|ref|XP_017137499.1| PREDICTED: adenosine kinase isoform X2 [Drosophila miranda]
>gi|1036818290|ref|XP_017137498.1| PREDICTED: adenosine kinase isoform X1 [Drosophila miranda]
>gi|1036829655|ref|XP_017138128.1| PREDICTED: 40S ribosomal protein S12 [Drosophila miranda]
>gi|1036829594|ref|XP_017138125.1| PREDICTED: zinc finger MYND domain-containing protein 10 hor
>gi|1036829636|ref|XP_017138127.1| PREDICTED: matrix-remodeling-associated protein 7 [Drosophi
>gi|1036829615|ref|XP_017138126.1| PREDICTED: uncharacterized protein LOC108152942 [Drosophila
>gi|1036843889|ref|XP_017138928.1| PREDICTED: cyclin-H [Drosophila miranda]
>gi|1036764131|ref|XP_017157102.1| PREDICTED: protein embryonic gonad [Drosophila miranda]
>gi|1036825202|ref|XP_017137876.1| PREDICTED: ras-interacting protein RIP3-like [Drosophila mi
>gi|1036825183|ref|XP_017137875.1| PREDICTED: ras-interacting protein RIP3-like [Drosophila mi
>gi|1036759145|ref|XP_017156820.1| PREDICTED: catalase C-like [Drosophila miranda]
>gi|1036844555|ref|XP_017138965.1| PREDICTED: beta-1-syntrophin isoform X1 [Drosophila miranda
>gi|1036844573|ref|XP_017138966.1| PREDICTED: beta-1-syntrophin isoform X1 [Drosophila miranda
>gi|1036844727|ref|XP_017138967.1| PREDICTED: beta-1-syntrophin isoform X1 [Drosophila miranda
>gi|1036844746|ref|XP_017138969.1| PREDICTED: beta-1-syntrophin isoform X2 [Drosophila miranda
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>gi|1036844765|ref|XP_017138970.1| PREDICTED: brain acid soluble protein 1 [Drosophila miranda]
>gi|1036770366|ref|XP_017134791.1| PREDICTED: uncharacterized protein LOC108150967 [Drosophila
>gi|1036767446|ref|XP_017134630.1| PREDICTED: uncharacterized protein LOC108150834 [Drosophila
>gi|1036849381|ref|XP_017139225.1| PREDICTED: uncharacterized protein LOC108153633 [Drosophila
>gi|1036849365|ref|XP_017139224.1| PREDICTED: translation initiation factor IF-2 [Drosophila m:
>gi|1036767741|ref|XP_017134643.1| PREDICTED: uncharacterized protein LOC108150851 [Drosophila
>gi|1036768687|ref|XP_017134695.1| PREDICTED: mucin-7 [Drosophila miranda]
>gi|1036823459|ref|XP_017137780.1| PREDICTED: uncharacterized protein LOC108152743 [Drosophila
>gi|1036823436|ref|XP_017137779.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036823478|ref|XP_017137781.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu:
>gi|1036823417|ref|XP_017137777.1| PREDICTED: zinc finger protein OZF [Drosophila miranda]
>gi|1036836973|ref|XP_017138540.1| PREDICTED: uncharacterized protein LOC108153223 isoform X1
>gi|1036836992|ref|XP_017138542.1| PREDICTED: uncharacterized protein LOC108153223 isoform X2
>gi|1036832570|ref|XP_017138294.1| PREDICTED: ninein [Drosophila miranda]
>gi|1036759164|ref|XP_017156821.1| PREDICTED: acanthoscurrin-2 [Drosophila miranda]
>gi|1036759183|ref|XP_017156822.1| PREDICTED: translation initiation factor IF-2-like, partial
>gi|1036852527|ref|XP_017139416.1| PREDICTED: uncharacterized protein LOC108153783 [Drosophila
>gi|1036828033|ref|XP_017138037.1| PREDICTED: alsin homolog [Drosophila miranda]
>gi|1036828073|ref|XP_017138039.1| PREDICTED: uncharacterized protein LOC108152904 isoform X2
>gi|1036828054|ref|XP_017138038.1| PREDICTED: uncharacterized protein LOC108152904 isoform X1
>gi|1036764702|ref|XP_017157131.1| PREDICTED: endocuticle structural glycoprotein ABD-5 [Droso;
>gi|1036841558|ref|XP_017138807.1| PREDICTED: ankyrin repeat and BTB/POZ domain-containing prot
>gi|1036841539|ref|XP_017138806.1| PREDICTED: ankyrin repeat and BTB/POZ domain-containing prot
>gi|1036847741|ref|XP_017139131.1| PREDICTED: copper transport protein ATOX1 [Drosophila mirano
>gi|1036851857|ref|XP_017139374.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036851675|ref|XP_017139362.1| PREDICTED: probable fumarate hydratase, mitochondrial [Dros(
>gi|1036811343|ref|XP_017137112.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036811365|ref|XP_017137113.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036811282|ref|XP_017137109.1| PREDICTED: fibroleukin [Drosophila miranda]
>gi|1036811305|ref|XP_017137110.1| PREDICTED: fibroleukin [Drosophila miranda]
>gi|1036811384|ref|XP_017137114.1| PREDICTED: ORM1-like protein [Drosophila miranda]
>gi|1036811324|ref|XP_017137111.1| PREDICTED: asparagine--tRNA ligase, cytoplasmic-like [Drosop
>gi|1036811912|ref|XP_017137141.1| PREDICTED: mediator of RNA polymerase II transcription subu:
>gi|1036811931|ref|XP_017137142.1| PREDICTED: mediator of RNA polymerase II transcription subu:
>gi|1036847271|ref|XP_017139106.1| PREDICTED: uncharacterized protein LOC108153549 [Drosophila
>gi|1036847290|ref|XP_017139107.1| PREDICTED: neuropeptide Y receptor type 6 [Drosophila miranc
>gi|1036847886|ref|XP_017139139.1| PREDICTED: uncharacterized protein LOC108153572 [Drosophila
>gi|1036847904|ref|XP_017139140.1| PREDICTED: uncharacterized protein LOC108153572 [Drosophila
>gi|1036847941|ref|XP_017139143.1| PREDICTED: leucine-rich repeat-containing protein 9 [Drosop]
>gi|1036847922|ref|XP_017139142.1| PREDICTED: glutamate receptor 2 [Drosophila miranda]
>gi|1036767964|ref|XP_017134655.1| PREDICTED: uncharacterized protein LOC108150864 [Drosophila
>gi|1036851890|ref|XP_017139377.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036851906|ref|XP_017139378.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036851925|ref|XP_017139379.1| PREDICTED: phospholipase B1, membrane-associated [Drosophila
>gi|1036834015|ref|XP_017138369.1| PREDICTED: ADP-dependent glucokinase [Drosophila miranda]
>gi|1036823840|ref|XP_017137800.1| PREDICTED: heat shock 70 kDa protein 4 isoform X1 [Drosophil
>gi|1036823859|ref|XP_017137801.1| PREDICTED: heat shock 70 kDa protein 4 isoform X1 [Drosophil
>gi|1036823878|ref|XP_017137802.1| PREDICTED: heat shock 70 kDa protein 4 isoform X2 [Drosophil
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>gi|1036818969|ref|XP_017137537.1| PREDICTED: neuralized-like protein 4 [Drosophila miranda]
>gi|1036838446|ref|XP_017138630.1| PREDICTED: RNA exonuclease 4 [Drosophila miranda]
>gi|1036838463|ref|XP_017138631.1| PREDICTED: M-phase-specific PLK1-interacting protein [Drosop
>gi|1036809572|ref|XP_017137015.1| PREDICTED: formin-like protein CG32138 isoform X5 [Drosophil
>gi|1036809502|ref|XP_017137011.1| PREDICTED: formin-like protein CG32138 isoform X1 [Drosophil
>gi|1036809589|ref|XP_017137016.1| PREDICTED: formin-like protein CG32138 isoform X6 [Drosophil
>gi|1036809521|ref|XP_017137012.1| PREDICTED: formin-like protein CG32138 isoform X2 [Drosophil
>gi|1036809538|ref|XP_017137013.1| PREDICTED: formin-like protein CG32138 isoform X3 [Drosophil
>gi|1036809555|ref|XP_017137014.1| PREDICTED: formin-like protein CG32138 isoform X4 [Drosophil
>gi|1036809464|ref|XP_017137009.1| PREDICTED: peroxisome biogenesis factor 1 [Drosophila miran
>gi|1036809446|ref|XP_017137008.1| PREDICTED: fibroblast growth factor receptor homolog 2 [Dros
>gi|1036809483|ref|XP_017137010.1| PREDICTED: fat-body protein 1 [Drosophila miranda]
>gi|1036846586|ref|XP_017139066.1| PREDICTED: fat-body protein 1 [Drosophila miranda]
>gi|1036835572|ref|XP_017138458.1| PREDICTED: transcription factor Sox-14 [Drosophila miranda]
>gi|1036759200|ref|XP_017156823.1| PREDICTED: BTB/POZ domain-containing protein 2-like [Drosop
>gi|1036835554|ref|XP_017138457.1| PREDICTED: RNA-binding protein fusilli [Drosophila miranda]
>gi|1036841614|ref|XP_017138810.1| PREDICTED: SOX domain-containing protein dichaete [Drosophil
>gi|1036852250|ref|XP_017139400.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036837329|ref|XP_017138562.1| PREDICTED: succinyl-CoA ligase [ADP/GDP-forming] subunit alp
>gi|1036811950|ref|XP_017137143.1| PREDICTED: protein encore isoform X1 [Drosophila miranda]
>gi|1036811988|ref|XP_017137145.1| PREDICTED: protein encore isoform X3 [Drosophila miranda]
>gi|1036811969|ref|XP_017137144.1| PREDICTED: protein encore isoform X2 [Drosophila miranda]
>gi|1036812007|ref|XP_017137146.1| PREDICTED: protein encore isoform X4 [Drosophila miranda]
>gi|1036846199|ref|XP_017139043.1| PREDICTED: LIM/homeobox protein Awh isoform X1 [Drosophila r
>gi|1036846216|ref|XP_017139044.1| PREDICTED: LIM/homeobox protein Awh isoform X2 [Drosophila r
>gi|1036846235|ref|XP_017139045.1| PREDICTED: LIM/homeobox protein Awh isoform X3 [Drosophila r
>gi|1036846256|ref|XP_017139046.1| PREDICTED: LIM/homeobox protein Awh isoform X4 [Drosophila r
>gi|1036763260|ref|XP_017157056.1| PREDICTED: uncharacterized protein LOC108165511 [Drosophila
>gi|1036833374|ref|XP_017138342.1| PREDICTED: uncharacterized protein LOC108153072 [Drosophila
>gi|1036759219|ref|XP_017156824.1| PREDICTED: gustatory and odorant receptor 63a [Drosophila m
>gi|1036833340|ref|XP_017138339.1| PREDICTED: hsp90 co-chaperone Cdc37 [Drosophila miranda]
>gi|1036833356|ref|XP_017138341.1| PREDICTED: dnaJ homolog subfamily B member 13 [Drosophila m
>gi|1036819433|ref|XP_017137560.1| PREDICTED: patj homolog isoform X1 [Drosophila miranda]
>gi|1036819473|ref|XP_017137563.1| PREDICTED: patj homolog isoform X3 [Drosophila miranda]
>gi|1036819454|ref|XP_017137561.1| PREDICTED: patj homolog isoform X2 [Drosophila miranda]
>gi|1036819492|ref|XP_017137564.1| PREDICTED: protein JTB [Drosophila miranda]
>gi|1036832072|ref|XP_017138268.1| PREDICTED: bromodomain-containing protein DDB_G0280777 isofc
>gi|1036832089|ref|XP_017138269.1| PREDICTED: basic-leucine zipper transcription factor A isofc
>gi|1036848164|ref|XP_017139154.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036769686|ref|XP_017134754.1| PREDICTED: uncharacterized protein LOC108150935 [Drosophila
>gi|1036767187|ref|XP_017134617.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1036851618|ref|XP_017139359.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1036850597|ref|XP_017139298.1| PREDICTED: uncharacterized protein LOC108153687 isoform X1
>gi|1036850616|ref|XP_017139299.1| PREDICTED: uncharacterized protein LOC108153687 isoform X2
>gi|1036805293|ref|XP_017136784.1| PREDICTED: CUE domain-containing protein 1 isoform X1 [Dros
>gi|1036805312|ref|XP_017136785.1| PREDICTED: CUE domain-containing protein 1 isoform X2 [Dros
>gi|1036805331|ref|XP_017136786.1| PREDICTED: CUE domain-containing protein 1 isoform X3 [Dros
>gi|1036805348|ref|XP_017136787.1| PREDICTED: ras-like protein 3 [Drosophila miranda]
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>gi|1036805274|ref|XP_017136783.1| PREDICTED: HBS1-like protein [Drosophila miranda]
>gi|1036834916|ref|XP_017138419.1| PREDICTED: transmembrane protein 134 [Drosophila miranda]
>gi|1036823038|ref|XP_017137755.1| PREDICTED: uncharacterized protein LOC108152728 [Drosophila
>gi|1036823019|ref|XP_017137754.1| PREDICTED: FACT complex subunit spt16 isoform X2 [Drosophila
>gi|1036823000|ref|XP_017137752.1| PREDICTED: FACT complex subunit spt16 isoform X1 [Drosophila
>gi|1036840088|ref|XP_017138722.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840107|ref|XP_017138723.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840126|ref|XP_017138724.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840145|ref|XP_017138725.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X1 [Dros
>gi|1036840164|ref|XP_017138726.1| PREDICTED: carbohydrate sulfotransferase 11 isoform X2 [Dros
>gi|1036840183|ref|XP_017138727.1| PREDICTED: adenine phosphoribosyltransferase [Drosophila mi
>gi|1036840204|ref|XP_017138728.1| PREDICTED: adenine phosphoribosyltransferase [Drosophila mi
>gi|1036836729|ref|XP_017138526.1| PREDICTED: lipoma HMGIC fusion partner-like 3 protein [Dros
>gi|1036846935|ref|XP_017139086.1| PREDICTED: lysosome membrane protein 2 isoform X1 [Drosophil
>gi|1036846954|ref|XP_017139087.1| PREDICTED: lysosome membrane protein 2 isoform X1 [Drosophil
>gi|1036846973|ref|XP_017139088.1| PREDICTED: lysosome membrane protein 2 isoform X2 [Drosophil
>gi|1036851944|ref|XP_017139380.1| PREDICTED: alpha-protein kinase 1 [Drosophila miranda]
>gi|1036848126|ref|XP_017139152.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1036771948|ref|XP_017134879.1| PREDICTED: microfibrillar-associated protein 1 [Drosophila r
>gi|1036759238|ref|XP_017156825.1| PREDICTED: uncharacterized protein LOC108165301 [Drosophila
>gi|1036771966|ref|XP_017134881.1| PREDICTED: microtubule-associated protein futsch-like [Dros
>gi|1036771929|ref|XP_017134878.1| PREDICTED: ankyrin repeat domain-containing protein 26 [Dros
>gi|1036771910|ref|XP_017134877.1| PREDICTED: uncharacterized protein LOC108151029 [Drosophila
>gi|1036788535|ref|XP_017135837.1| PREDICTED: RING finger protein 10 [Drosophila miranda]
>gi|1036788606|ref|XP_017135841.1| PREDICTED: choline-phosphate cytidylyltransferase A [Drosopl
>gi|1036788570|ref|XP_017135839.1| PREDICTED: choline-phosphate cytidylyltransferase A [Drosopl
>gi|1036788588|ref|XP_017135840.1| PREDICTED: choline-phosphate cytidylyltransferase A [Drosopl
>gi|1036788519|ref|XP_017135836.1| PREDICTED: DNA polymerase eta [Drosophila miranda]
>gi|1036788550|ref|XP_017135838.1| PREDICTED: solute carrier family 22 member 3 [Drosophila mi
>gi|1036807066|ref|XP_017136880.1| PREDICTED: poly(rC)-binding protein 3 isoform X16 [Drosophil
>gi|1036807048|ref|XP_017136879.1| PREDICTED: poly(rC)-binding protein 3 isoform X15 [Drosophil
>gi|1036807030|ref|XP_017136878.1| PREDICTED: poly(rC)-binding protein 3 isoform X14 [Drosophil
>gi|1036807016|ref|XP_017136876.1| PREDICTED: poly(rC)-binding protein 3 isoform X13 [Drosophil
>gi|1036806997|ref|XP_017136875.1| PREDICTED: poly(rC)-binding protein 3 isoform X12 [Drosophil
>gi|1036806981|ref|XP_017136874.1| PREDICTED: poly(rC)-binding protein 3 isoform X11 [Drosophil
>gi|1036806964|ref|XP_017136873.1| PREDICTED: poly(rC)-binding protein 3 isoform X10 [Drosophil
>gi|1036806927|ref|XP_017136871.1| PREDICTED: poly(rC)-binding protein 3 isoform X8 [Drosophila
>gi|1036806816|ref|XP_017136865.1| PREDICTED: poly(rC)-binding protein 3 isoform X2 [Drosophila
>gi|1036806946|ref|XP_017136872.1| PREDICTED: poly(rC)-binding protein 3 isoform X9 [Drosophila
>gi|1036806909|ref|XP_017136870.1| PREDICTED: poly(rC)-binding protein 3 isoform X7 [Drosophila
>gi|1036806890|ref|XP_017136869.1| PREDICTED: poly(rC)-binding protein 3 isoform X6 [Drosophila
>gi|1036806872|ref|XP_017136868.1| PREDICTED: poly(rC)-binding protein 3 isoform X5 [Drosophila
>gi|1036806853|ref|XP_017136867.1| PREDICTED: poly(rC)-binding protein 3 isoform X4 [Drosophila
>gi|1036806834|ref|XP_017136866.1| PREDICTED: poly(rC)-binding protein 3 isoform X3 [Drosophila
>gi|1036806742|ref|XP_017136861.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036806762|ref|XP_017136862.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036806781|ref|XP_017136863.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
>gi|1036806799|ref|XP_017136864.1| PREDICTED: poly(rC)-binding protein 3 isoform X1 [Drosophila
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>gi|1036768667|ref|XP_017134694.1| PREDICTED: neuropeptide-like protein 29 [Drosophila miranda]
>gi|1036832974|ref|XP_017138318.1| PREDICTED: delta-1-pyrroline-5-carboxylate dehydrogenase, mi
>gi|1036825981|ref|XP_017137917.1| PREDICTED: nucleolar and coiled-body phosphoprotein 1-like
>gi|1036825939|ref|XP_017137915.1| PREDICTED: nucleolin 2-like isoform X1 [Drosophila miranda]
>gi|1036825958|ref|XP_017137916.1| PREDICTED: nucleolin 1-like isoform X2 [Drosophila miranda]
>gi|1036838182|ref|XP_017138614.1| PREDICTED: uncharacterized protein LOC108153263 [Drosophila
>gi|1036826144|ref|XP_017137927.1| PREDICTED: small integral membrane protein 8 [Drosophila mi
>gi|1036826127|ref|XP_017137926.1| PREDICTED: eukaryotic translation initiation factor 2A [Dros
>gi|1036846605|ref|XP_017139067.1| PREDICTED: uncharacterized protein LOC108153520 [Drosophila
>gi|1036799073|ref|XP_017136433.1| PREDICTED: serine/arginine repetitive matrix protein 1 [Dros
>gi|1036799107|ref|XP_017136435.1| PREDICTED: 40S ribosomal protein S4 [Drosophila miranda]
>gi|1036799090|ref|XP_017136434.1| PREDICTED: syntaxin-12 [Drosophila miranda]
>gi|1036799124|ref|XP_017136436.1| PREDICTED: uncharacterized protein LOC108151983 [Drosophila
>gi|1036768442|ref|XP_017134682.1| PREDICTED: uncharacterized protein LOC108150884 [Drosophila
>gi|1036767923|ref|XP_017134653.1| PREDICTED: uncharacterized protein LOC108150862 [Drosophila
>gi|1036759257|ref|XP_017156826.1| PREDICTED: uncharacterized protein LOC108165303 [Drosophila
>gi|1036848278|ref|XP_017139161.1| PREDICTED: serine/threonine-protein phosphatase PP1-like [Dr
>gi|1036835868|ref|XP_017138475.1| PREDICTED: leucine-rich repeat-containing protein 15 [Drosop
>gi|1036852125|ref|XP_017139392.1| PREDICTED: DC-STAMP domain-containing protein 1 [Drosophila
>gi|1036814002|ref|XP_017137255.1| PREDICTED: uncharacterized protein LOC108152434 [Drosophila
>gi|1036813984|ref|XP_017137254.1| PREDICTED: chondroadherin-like protein [Drosophila miranda]
>gi|1036759273|ref|XP_017156827.1| PREDICTED: hydrogenase nickel incorporation protein HypB-lik
>gi|1036766387|ref|XP_017134570.1| PREDICTED: UNC93-like protein [Drosophila miranda]
>gi|1036766138|ref|XP_017134560.1| PREDICTED: collagen alpha-1(VIII) chain-like [Drosophila mi
>gi|1036850436|ref|XP_017139289.1| PREDICTED: cysteine and histidine-rich protein 1-B-like [Dro
>gi|1036850453|ref|XP_017139290.1| PREDICTED: neuropeptide-like 2 [Drosophila miranda]
>gi|1036847960|ref|XP_017139144.1| PREDICTED: uncharacterized protein LOC108153576 [Drosophila
>gi|1036846074|ref|XP_017139037.1| PREDICTED: uncharacterized protein LOC108153497 [Drosophila
>gi|1036759293|ref|XP_017156828.1| PREDICTED: uncharacterized protein LOC108165305, partial [D
>gi|1036846055|ref|XP_017139036.1| PREDICTED: nucleolin [Drosophila miranda]
>gi|1036852417|ref|XP_017139410.1| PREDICTED: uncharacterized protein LOC108153779 [Drosophila
>gi|1036821869|ref|XP_017137689.1| PREDICTED: uncharacterized protein LOC108152694 [Drosophila
>gi|1036821890|ref|XP_017137690.1| PREDICTED: transmembrane protein 19 isoform X1 [Drosophila n
>gi|1036821911|ref|XP_017137691.1| PREDICTED: transmembrane protein 19 isoform X2 [Drosophila n
>gi|1036821850|ref|XP_017137688.1| PREDICTED: POC1 centriolar protein homolog isoform X2 [Drosc
>gi|1036821813|ref|XP_017137685.1| PREDICTED: POC1 centriolar protein homolog isoform X1 [Drosc
>gi|1036821831|ref|XP_017137686.1| PREDICTED: POC1 centriolar protein homolog isoform X1 [Drosc
>gi|1036846624|ref|XP_017139068.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036849657|ref|XP_017139242.1| PREDICTED: tetratricopeptide repeat protein 36 homolog [Dros
>gi|1036849617|ref|XP_017139240.1| PREDICTED: uncharacterized protein LOC108153647 isoform X1
>gi|1036849636|ref|XP_017139241.1| PREDICTED: uncharacterized protein LOC108153647 isoform X2
>gi|1036842712|ref|XP_017138861.1| PREDICTED: whirlin isoform X3 [Drosophila miranda]
>gi|1036842833|ref|XP_017138868.1| PREDICTED: whirlin isoform X10 [Drosophila miranda]
>gi|1036842780|ref|XP_017138865.1| PREDICTED: whirlin isoform X7 [Drosophila miranda]
>gi|1036842728|ref|XP_017138862.1| PREDICTED: whirlin isoform X4 [Drosophila miranda]
>gi|1036842815|ref|XP_017138867.1| PREDICTED: whirlin isoform X9 [Drosophila miranda]
>gi|1036842693|ref|XP_017138859.1| PREDICTED: whirlin isoform X2 [Drosophila miranda]
>gi|1036842797|ref|XP_017138866.1| PREDICTED: whirlin isoform X8 [Drosophila miranda]
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>gi|1036842747|ref|XP_017138863.1| PREDICTED: whirlin isoform X5 [Drosophila miranda]
>gi|1036842764|ref|XP_017138864.1| PREDICTED: whirlin isoform X6 [Drosophila miranda]
>gi|1036842677|ref|XP_017138858.1| PREDICTED: whirlin isoform X1 [Drosophila miranda]
>gi|1036842851|ref|XP_017138869.1| PREDICTED: whirlin isoform X11 [Drosophila miranda]
>gi|1036842864|ref|XP_017138870.1| PREDICTED: uncharacterized protein LOC108153394 [Drosophila
>gi|1036777927|ref|XP_017135223.1| PREDICTED: ral guanine nucleotide dissociation stimulator-l:
>gi|1036777921|ref|XP_017135222.1| PREDICTED: G patch domain-containing protein 1 homolog [Dros
>gi|1036777911|ref|XP_017135220.1| PREDICTED: dynactin subunit 1 [Drosophila miranda]
>gi|1036777915|ref|XP_017135221.1| PREDICTED: dynactin subunit 1 [Drosophila miranda]
>gi|1036777939|ref|XP_017135224.1| PREDICTED: bicaudal D-related protein homolog isoform X1 [D:
>gi|1036777964|ref|XP_017135225.1| PREDICTED: bicaudal D-related protein homolog isoform X1 [D:
>gi|1036777976|ref|XP_017135226.1| PREDICTED: bicaudal D-related protein homolog isoform X2 [D:
>gi|1036777990|ref|XP_017135227.1| PREDICTED: bicaudal D-related protein homolog isoform X2 [D:
>gi|1036838482|ref|XP_017138632.1| PREDICTED: gastrula zinc finger protein XlCGF57.1 [Drosophi:
>gi|1036785178|ref|XP_017135650.1| PREDICTED: ataxin-2 homolog [Drosophila miranda]
>gi|1036785322|ref|XP_017135656.1| PREDICTED: uncharacterized protein LOC108151519 [Drosophila
>gi|1036785197|ref|XP_017135651.1| PREDICTED: heat shock 70 kDa protein cognate 1 [Drosophila m
>gi|1036785286|ref|XP_017135654.1| PREDICTED: uncharacterized protein LOC108151517 isoform X3
>gi|1036785269|ref|XP_017135653.1| PREDICTED: uncharacterized protein LOC108151517 isoform X2
>gi|1036785233|ref|XP_017135652.1| PREDICTED: uncharacterized protein LOC108151517 isoform X1
>gi|1036759312|ref|XP_017156830.1| PREDICTED: cathepsin L-like [Drosophila miranda]
>gi|1036785304|ref|XP_017135655.1| PREDICTED: pre-intermoult gene 1 protein-like [Drosophila m:
>gi|1036759331|ref|XP_017156831.1| PREDICTED: histone deacetylase HDT3-like [Drosophila miranda
>gi|1036839962|ref|XP_017138714.1| PREDICTED: digestive cysteine proteinase 1 [Drosophila mira:
>gi|1036813746|ref|XP_017137240.1| PREDICTED: formin-binding protein 1-like isoform X3 [Drosopl
>gi|1036813710|ref|XP_017137238.1| PREDICTED: formin-binding protein 1-like isoform X1 [Drosopl
>gi|1036813729|ref|XP_017137239.1| PREDICTED: formin-binding protein 1-like isoform X2 [Drosopl
>gi|1036813799|ref|XP_017137243.1| PREDICTED: formin-binding protein 1-like isoform X6 [Drosopl
>gi|1036813818|ref|XP_017137244.1| PREDICTED: formin-binding protein 1-like isoform X7 [Drosopl
>gi|1036813763|ref|XP_017137241.1| PREDICTED: formin-binding protein 1-like isoform X4 [Drosopl
>gi|1036813781|ref|XP_017137242.1| PREDICTED: formin-binding protein 1-like isoform X5 [Drosopl
>gi|1036813837|ref|XP_017137245.1| PREDICTED: formin-binding protein 1-like isoform X8 [Drosopl
>gi|1036813857|ref|XP_017137246.1| PREDICTED: formin-binding protein 1-like isoform X9 [Drosopl
>gi|1036786248|ref|XP_017135709.1| PREDICTED: uncharacterized protein LOC108151553 [Drosophila
>gi|1036786195|ref|XP_017135706.1| PREDICTED: uncharacterized protein LOC108151551 isoform X1
>gi|1036786212|ref|XP_017135707.1| PREDICTED: uncharacterized protein LOC108151551 isoform X2
>gi|1036786029|ref|XP_017135696.1| PREDICTED: semaphorin-5A isoform X2 [Drosophila miranda]
>gi|1036785988|ref|XP_017135694.1| PREDICTED: semaphorin-5A isoform X1 [Drosophila miranda]
>gi|1036786008|ref|XP_017135695.1| PREDICTED: semaphorin-5A isoform X1 [Drosophila miranda]
>gi|1036786231|ref|XP_017135708.1| PREDICTED: uncharacterized protein LOC108151552 [Drosophila
>gi|1036786048|ref|XP_017135697.1| PREDICTED: uncharacterized protein LOC108151547 isoform X1
>gi|1036786065|ref|XP_017135698.1| PREDICTED: uncharacterized protein LOC108151547 isoform X2
>gi|1036786084|ref|XP_017135699.1| PREDICTED: uncharacterized protein LOC108151548 isoform X1
>gi|1036786103|ref|XP_017135700.1| PREDICTED: uncharacterized protein LOC108151548 isoform X2
>gi|1036786159|ref|XP_017135704.1| PREDICTED: uncharacterized protein LOC108151550 isoform X1
>gi|1036786177|ref|XP_017135705.1| PREDICTED: uncharacterized protein LOC108151550 isoform X2
>gi|1036786141|ref|XP_017135703.1| PREDICTED: nuclease-sensitive element-binding protein 1 iso:
>gi|1036786122|ref|XP_017135702.1| PREDICTED: nuclease-sensitive element-binding protein 1 iso:
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>gi|1036759350|ref|XP_017156832.1| PREDICTED: uncharacterized protein LOC108165308 [Drosophila
>gi|1036790950|ref|XP_017135974.1| PREDICTED: sterol regulatory element-binding protein 1 [Dros
>gi|1036790969|ref|XP_017135975.1| PREDICTED: sterol regulatory element-binding protein 1 [Dros
>gi|1036791116|ref|XP_017135984.1| PREDICTED: uncharacterized protein LOC108151719 isoform X3
>gi|1036791040|ref|XP_017135979.1| PREDICTED: uncharacterized protein LOC108151718 isoform X1
>gi|1036791059|ref|XP_017135980.1| PREDICTED: uncharacterized protein LOC108151718 isoform X2
>gi|1036791097|ref|XP_017135983.1| PREDICTED: uncharacterized protein LOC108151719 isoform X2
>gi|1036791078|ref|XP_017135982.1| PREDICTED: uncharacterized protein LOC108151719 isoform X1
>gi|1036791135|ref|XP_017135985.1| PREDICTED: uncharacterized protein LOC108151720 [Drosophila
>gi|1036791154|ref|XP_017135986.1| PREDICTED: coiled-coil domain-containing protein 58 [Drosop
>gi|1036791173|ref|XP_017135987.1| PREDICTED: DNA repair protein SWI5 homolog [Drosophila mira
>gi|1036790984|ref|XP_017135976.1| PREDICTED: importin subunit alpha-7 [Drosophila miranda]
>gi|1036791003|ref|XP_017135977.1| PREDICTED: probable cytochrome P450 305a1 isoform X1 [Droso
>gi|1036791021|ref|XP_017135978.1| PREDICTED: probable cytochrome P450 305a1 isoform X2 [Droso
>gi|1036834413|ref|XP_017138389.1| PREDICTED: protein cycle isoform X2 [Drosophila miranda]
>gi|1036834394|ref|XP_017138387.1| PREDICTED: protein cycle isoform X1 [Drosophila miranda]
>gi|1036834432|ref|XP_017138390.1| PREDICTED: acidic fibroblast growth factor intracellular-bi
>gi|1036834470|ref|XP_017138392.1| PREDICTED: deformed epidermal autoregulatory factor 1 isofor
>gi|1036834489|ref|XP_017138393.1| PREDICTED: deformed epidermal autoregulatory factor 1 isofor
>gi|1036826297|ref|XP_017137936.1| PREDICTED: serine/threonine-protein kinase tricorner [Drosop
>gi|1036826315|ref|XP_017137937.1| PREDICTED: uncharacterized protein LOC108152840 [Drosophila
>gi|1036803956|ref|XP_017136707.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036803974|ref|XP_017136708.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036833484|ref|XP_017138347.1| PREDICTED: bifunctional 3'-phosphoadenosine 5'-phosphosulfa
>gi|1036833448|ref|XP_017138345.1| PREDICTED: bifunctional 3'-phosphoadenosine 5'-phosphosulfa
>gi|1036833467|ref|XP_017138346.1| PREDICTED: bifunctional 3'-phosphoadenosine 5'-phosphosulfa
>gi|1036831476|ref|XP_017138236.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1036831495|ref|XP_017138237.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1036831512|ref|XP_017138238.1| PREDICTED: ras-related protein Rab-8A [Drosophila miranda]
>gi|1036838201|ref|XP_017138616.1| PREDICTED: uncharacterized protein LOC108153264 [Drosophila
>gi|1036803770|ref|XP_017136698.1| PREDICTED: immediate early response 3-interacting protein 1
>gi|1036803751|ref|XP_017136697.1| PREDICTED: biogenesis of lysosome-related organelles comple
>gi|1036803716|ref|XP_017136695.1| PREDICTED: uncharacterized protein YER152C [Drosophila mira
>gi|1036803608|ref|XP_017136689.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803627|ref|XP_017136690.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803645|ref|XP_017136691.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803663|ref|XP_017136692.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803571|ref|XP_017136687.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803590|ref|XP_017136688.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803555|ref|XP_017136686.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803536|ref|XP_017136685.1| PREDICTED: proton-coupled amino acid transporter 1 isoform
>gi|1036803517|ref|XP_017136684.1| PREDICTED: proton-coupled amino acid transporter 4 isoform
>gi|1036803698|ref|XP_017136694.1| PREDICTED: proton-coupled amino acid transporter 1 [Drosophi
>gi|1036803680|ref|XP_017136693.1| PREDICTED: proton-coupled amino acid transporter 3 [Drosophi
>gi|1036803732|ref|XP_017136696.1| PREDICTED: acyl-CoA Delta(11) desaturase [Drosophila miranda
>gi|1036823614|ref|XP_017137789.1| PREDICTED: uncharacterized protein LOC108152746 isoform X4
>gi|1036823499|ref|XP_017137783.1| PREDICTED: uncharacterized protein LOC108152745 [Drosophila
>gi|1036823518|ref|XP_017137784.1| PREDICTED: histidine-rich glycoprotein isoform X1 [Drosophil
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>gi|1036823537|ref|XP_017137785.1| PREDICTED: histidine-rich glycoprotein isoform X2 [Drosophil
>gi|1036823559|ref|XP_017137786.1| PREDICTED: histidine-rich glycoprotein isoform X3 [Drosophil
>gi|1036823577|ref|XP_017137787.1| PREDICTED: histidine-rich glycoprotein isoform X3 [Drosophil
>gi|1036823595|ref|XP_017137788.1| PREDICTED: histidine-rich glycoprotein isoform X3 [Drosophil
>gi|1036825772|ref|XP_017137905.1| PREDICTED: diphosphoinositol polyphosphate phosphohydrolase
>gi|1036825790|ref|XP_017137906.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036764021|ref|XP_017157100.1| PREDICTED: larval cuticle protein LCP-17-like [Drosophila mi
>gi|1036834545|ref|XP_017138397.1| PREDICTED: mitochondrial inner membrane protein OXA1L [Droso
>gi|1036834566|ref|XP_017138398.1| PREDICTED: MIP18 family protein CG7949 [Drosophila miranda]
>gi|1036844391|ref|XP_017138956.1| PREDICTED: histidine-rich protein PFHRP-II isoform X1 [Dros
>gi|1036844409|ref|XP_017138957.1| PREDICTED: sarcoplasmic reticulum histidine-rich calcium-bin
>gi|1036844427|ref|XP_017138958.1| PREDICTED: biogenesis of lysosome-related organelles complex
>gi|1036826109|ref|XP_017137925.1| PREDICTED: ATP-dependent RNA helicase DDX42 [Drosophila mira
>gi|1036804160|ref|XP_017136719.1| PREDICTED: zinc finger MIZ domain-containing protein 2 [Dros
>gi|1036759367|ref|XP_017156833.1| PREDICTED: keratin, type II cytoskeletal 2 epidermal [Drosop
>gi|1036783689|ref|XP_017135562.1| PREDICTED: transmembrane protein 184B isoform X1 [Drosophila
>gi|1036783707|ref|XP_017135564.1| PREDICTED: transmembrane protein 184B isoform X2 [Drosophila
>gi|1036783762|ref|XP_017135567.1| PREDICTED: 40S ribosomal protein S6-like [Drosophila miranda
>gi|1036783726|ref|XP_017135565.1| PREDICTED: rho GTPase-activating protein 17 [Drosophila mira
>gi|1036783610|ref|XP_017135558.1| PREDICTED: uncharacterized protein LOC108151451 [Drosophila
>gi|1036783629|ref|XP_017135559.1| PREDICTED: uncharacterized protein LOC108151451 [Drosophila
>gi|1036783745|ref|XP_017135566.1| PREDICTED: peroxisome biogenesis factor 10 [Drosophila mira
>gi|1036783650|ref|XP_017135560.1| PREDICTED: molybdenum cofactor biosynthesis protein 1 isofo
>gi|1036783671|ref|XP_017135561.1| PREDICTED: molybdenum cofactor biosynthesis protein 1 isofo
>gi|1036783779|ref|XP_017135568.1| PREDICTED: probable prefoldin subunit 2 [Drosophila miranda]
>gi|1036827959|ref|XP_017138034.1| PREDICTED: unc-112-related protein [Drosophila miranda]
>gi|1036830330|ref|XP_017138169.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036830348|ref|XP_017138171.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036830367|ref|XP_017138172.1| PREDICTED: putative uncharacterized protein DDB_G0272516 iso
>gi|1036830385|ref|XP_017138173.1| PREDICTED: probable serine/threonine-protein kinase yakA iso
>gi|1036830404|ref|XP_017138174.1| PREDICTED: adenylate cyclase, terminal-differentiation speci
>gi|1036830423|ref|XP_017138175.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X5
>gi|1036830440|ref|XP_017138176.1| PREDICTED: G-box-binding factor isoform X6 [Drosophila miran
>gi|1036830459|ref|XP_017138177.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X7
>gi|1036830478|ref|XP_017138178.1| PREDICTED: uncharacterized protein LOC108152969 isoform X8
>gi|1036830495|ref|XP_017138179.1| PREDICTED: uncharacterized protein LOC108152969 isoform X9
>gi|1036830514|ref|XP_017138180.1| PREDICTED: uncharacterized protein LOC108152969 isoform X10
>gi|1036842584|ref|XP_017138853.1| PREDICTED: uncharacterized protein LOC108153389 [Drosophila
>gi|1036842530|ref|XP_017138851.1| PREDICTED: glutamate decarboxylase [Drosophila miranda]
>gi|1036842565|ref|XP_017138852.1| PREDICTED: glutamate decarboxylase [Drosophila miranda]
>gi|1036848824|ref|XP_017139192.1| PREDICTED: fumarylacetoacetase [Drosophila miranda]
>gi|1036830702|ref|XP_017138191.1| PREDICTED: bone morphogenetic protein receptor type-2 [Dros
>gi|1036830721|ref|XP_017138192.1| PREDICTED: bone morphogenetic protein receptor type-2 [Dros
>gi|1036830740|ref|XP_017138194.1| PREDICTED: cytochrome P450 302a1, mitochondrial [Drosophila
>gi|1036851838|ref|XP_017139373.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036851819|ref|XP_017139371.1| PREDICTED: 2-hydroxyacylsphingosine 1-beta-galactosyltransfe
>gi|1036846735|ref|XP_017139075.1| PREDICTED: uncharacterized protein LOC108153526 isoform X2
>gi|1036846717|ref|XP_017139074.1| PREDICTED: uncharacterized protein LOC108153526 isoform X1
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>gi|1036846754|ref|XP_017139076.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846771|ref|XP_017139077.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846790|ref|XP_017139078.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846808|ref|XP_017139079.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036846827|ref|XP_017139080.1| PREDICTED: protein dpy-30 homolog [Drosophila miranda]
>gi|1036832162|ref|XP_017138274.1| PREDICTED: microspherule protein 1 [Drosophila miranda]
>gi|1036834451|ref|XP_017138391.1| PREDICTED: BCL2/adenovirus E1B 19 kDa protein-interacting pr
>gi|1036794954|ref|XP_017136201.1| PREDICTED: gustatory receptor for sugar taste 64f isoform X1
>gi|1036794970|ref|XP_017136202.1| PREDICTED: gustatory receptor for sugar taste 64f isoform X2
>gi|1036795027|ref|XP_017136205.1| PREDICTED: gustatory receptor for sugar taste 64e isoform X1
>gi|1036795046|ref|XP_017136206.1| PREDICTED: gustatory receptor for sugar taste 64e isoform X2
>gi|1036795119|ref|XP_017136211.1| PREDICTED: gustatory receptor for sugar taste 64d isoform X4
>gi|1036795081|ref|XP_017136208.1| PREDICTED: gustatory receptor for sugar taste 64c isoform X2
>gi|1036795100|ref|XP_017136209.1| PREDICTED: gustatory receptor for sugar taste 64c isoform X3
>gi|1036795065|ref|XP_017136207.1| PREDICTED: gustatory receptor for sugar taste 64c isoform X1
>gi|1036795213|ref|XP_017136216.1| PREDICTED: gustatory receptor for sugar taste 64c-like isofo
>gi|1036795194|ref|XP_017136215.1| PREDICTED: gustatory receptor for sugar taste 64c-like isofo
>gi|1036795230|ref|XP_017136217.1| PREDICTED: gustatory receptor for sugar taste 64b [Drosophil
>gi|1036795008|ref|XP_017136204.1| PREDICTED: gustatory receptor for sugar taste 64a isoform X2
>gi|1036794988|ref|XP_017136203.1| PREDICTED: gustatory receptor for sugar taste 64a isoform X1
>gi|1036795139|ref|XP_017136212.1| PREDICTED: gustatory receptor for sugar taste 61a isoform X1
>gi|1036795175|ref|XP_017136214.1| PREDICTED: gustatory receptor for sugar taste 61a isoform X3
>gi|1036795156|ref|XP_017136213.1| PREDICTED: gustatory receptor for sugar taste 61a isoform X2
>gi|1036794824|ref|XP_017136193.1| PREDICTED: centrosomal protein of 290 kDa isoform X1 [Drosop
>gi|1036794843|ref|XP_017136194.1| PREDICTED: centrosomal protein of 290 kDa isoform X1 [Drosop
>gi|1036794860|ref|XP_017136195.1| PREDICTED: centrosomal protein of 290 kDa isoform X2 [Drosop
>gi|1036795267|ref|XP_017136219.1| PREDICTED: enoyl-CoA delta isomerase 2, mitochondrial-like
>gi|1036795248|ref|XP_017136218.1| PREDICTED: enoyl-CoA delta isomerase 2, mitochondrial-like
>gi|1036794938|ref|XP_017136200.1| PREDICTED: uncharacterized protein LOC108151835 [Drosophila
>gi|1036794879|ref|XP_017136197.1| PREDICTED: uncharacterized protein LOC108151834 isoform X1
>gi|1036794898|ref|XP_017136198.1| PREDICTED: uncharacterized protein LOC108151834 isoform X1
>gi|1036794918|ref|XP_017136199.1| PREDICTED: uncharacterized protein LOC108151834 isoform X2
>gi|1036781482|ref|XP_017135435.1| PREDICTED: nuclear receptor coactivator 6 [Drosophila miranc
>gi|1036781444|ref|XP_017135433.1| PREDICTED: monocarboxylate transporter 1 [Drosophila miranda
>gi|1036781463|ref|XP_017135434.1| PREDICTED: glycoprotein-N-acetylgalactosamine 3-beta-galacto
>gi|1036805988|ref|XP_017136824.1| PREDICTED: uncharacterized protein LOC108152187 [Drosophila
>gi|1036805970|ref|XP_017136823.1| PREDICTED: gastrula zinc finger protein XlCGF53.1 [Drosophil
>gi|1036806098|ref|XP_017136825.1| PREDICTED: LOW QUALITY PROTEIN: glucose transporter type 1
>gi|1036846991|ref|XP_017139089.1| PREDICTED: protein FAM69C [Drosophila miranda]
>gi|1036838144|ref|XP_017138612.1| PREDICTED: uncharacterized protein LOC108153260 [Drosophila
>gi|1036838125|ref|XP_017138611.1| PREDICTED: myb-like protein AA [Drosophila miranda]
>gi|1036788823|ref|XP_017135855.1| PREDICTED: phosphatase and actin regulator 4 isoform X11 [Dr
>gi|1036788750|ref|XP_017135850.1| PREDICTED: phosphatase and actin regulator 4 isoform X7 [Dro
>gi|1036788716|ref|XP_017135848.1| PREDICTED: neurogenic protein mastermind isoform X5 [Drosoph
>gi|1036788697|ref|XP_017135847.1| PREDICTED: neurogenic protein mastermind isoform X4 [Drosoph
>gi|1036788678|ref|XP_017135846.1| PREDICTED: neurogenic protein mastermind isoform X3 [Drosoph
>gi|1036788659|ref|XP_017135845.1| PREDICTED: neurogenic protein mastermind isoform X2 [Drosoph
>gi|1036788624|ref|XP_017135842.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosoph
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>gi|1036788641|ref|XP_017135844.1| PREDICTED: neurogenic protein mastermind isoform X1 [Drosophila
>gi|1036788767|ref|XP_017135851.1| PREDICTED: phosphatase and actin regulator 4 isoform X8 [Dro
>gi|1036788842|ref|XP_017135856.1| PREDICTED: phosphatase and actin regulator 2 isoform X12 [Di
>gi|1036788786|ref|XP_017135852.1| PREDICTED: phosphatase and actin regulator 2 isoform X9 [Dro
>gi|1036788804|ref|XP_017135853.1| PREDICTED: neurogenic protein mastermind isoform X10 [Drosop
>gi|1036788736|ref|XP_017135849.1| PREDICTED: neurogenic protein mastermind isoform X6 [Drosoph
>gi|1036788861|ref|XP_017135857.1| PREDICTED: sodium channel protein Nach [Drosophila miranda]
>gi|1036788912|ref|XP_017135859.1| PREDICTED: uncharacterized protein LOC108151651 [Drosophila
>gi|1036788879|ref|XP_017135858.1| PREDICTED: protein nutcracker [Drosophila miranda]
>gi|1036809606|ref|XP_017137017.1| PREDICTED: integrator complex subunit 10 [Drosophila miranda
>gi|1036809657|ref|XP_017137020.1| PREDICTED: mpv17-like protein 2 [Drosophila miranda]
>gi|1036809676|ref|XP_017137023.1| PREDICTED: mpv17-like protein 2 [Drosophila miranda]
>gi|1036809623|ref|XP_017137018.1| PREDICTED: GPI mannosyltransferase 3 isoform X1 [Drosophila
>gi|1036809640|ref|XP_017137019.1| PREDICTED: GPI mannosyltransferase 3 isoform X2 [Drosophila
>gi|1036806136|ref|XP_017136827.1| PREDICTED: rab11 family-interacting protein 4B isoform X2 [
>gi|1036806169|ref|XP_017136830.1| PREDICTED: rab11 family-interacting protein 4B isoform X4 [
>gi|1036806187|ref|XP_017136831.1| PREDICTED: rab11 family-interacting protein 4B isoform X5 [
>gi|1036806205|ref|XP_017136832.1| PREDICTED: rab11 family-interacting protein 4B isoform X6 [
>gi|1036806224|ref|XP_017136833.1| PREDICTED: rab11 family-interacting protein 4B isoform X7 [
>gi|1036806279|ref|XP_017136836.1| PREDICTED: peptidyl-prolyl cis-trans isomerase [Drosophila r
>gi|1036806117|ref|XP_017136826.1| PREDICTED: bromodomain-containing protein DDB_G0280777 isofc
>gi|1036806151|ref|XP_017136829.1| PREDICTED: bromodomain-containing protein DDB_G0280777 isofc
>gi|1036806243|ref|XP_017136834.1| PREDICTED: rab11 family-interacting protein 4B isoform X8 [
>gi|1036806262|ref|XP_017136835.1| PREDICTED: MAGE-like protein 2 [Drosophila miranda]
>gi|1036806297|ref|XP_017136837.1| PREDICTED: uncharacterized protein LOC108152194 [Drosophila
>gi|1036764444|ref|XP_017157118.1| PREDICTED: alpha/beta-gliadin A-V [Drosophila miranda]
>gi|1036806412|ref|XP_017136844.1| PREDICTED: frizzled isoform X1 [Drosophila miranda]
>gi|1036806433|ref|XP_017136845.1| PREDICTED: frizzled isoform X2 [Drosophila miranda]
>gi|1036806452|ref|XP_017136846.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036851543|ref|XP_017139355.1| PREDICTED: uncharacterized protein LOC108153736 [Drosophila
>gi|1036839083|ref|XP_017138670.1| PREDICTED: uncharacterized protein LOC108153294 [Drosophila
>gi|1036759386|ref|XP_017156834.1| PREDICTED: digestive cysteine proteinase 1-like, partial [Di
>gi|1036787666|ref|XP_017135789.1| PREDICTED: uncharacterized protein LOC108151609 isoform X1
>gi|1036787681|ref|XP_017135790.1| PREDICTED: uncharacterized protein LOC108151609 isoform X2
>gi|1036787696|ref|XP_017135791.1| PREDICTED: uncharacterized protein LOC108151610 [Drosophila
>gi|1036787592|ref|XP_017135783.1| PREDICTED: uncharacterized protein LOC108151608 isoform X1
>gi|1036787607|ref|XP_017135785.1| PREDICTED: PDZ domain-containing protein 2 isoform X2 [Dros
>gi|1036787622|ref|XP_017135786.1| PREDICTED: PDZ domain-containing protein 2 isoform X3 [Dros
>gi|1036787637|ref|XP_017135787.1| PREDICTED: PDZ domain-containing protein 2 isoform X4 [Dros
>gi|1036787651|ref|XP_017135788.1| PREDICTED: PDZ domain-containing protein 2 isoform X5 [Dros
>gi|1036759403|ref|XP_017156835.1| PREDICTED: uncharacterized protein LOC108165311 [Drosophila
>gi|1036759420|ref|XP_017156836.1| PREDICTED: trypsin-1 [Drosophila miranda]
>gi|1036834033|ref|XP_017138370.1| PREDICTED: phosphate carrier protein, mitochondrial [Drosopl
>gi|1036834049|ref|XP_017138371.1| PREDICTED: phosphate carrier protein, mitochondrial [Drosopl
>gi|1036824733|ref|XP_017137852.1| PREDICTED: protein phosphatase 1 regulatory subunit 3C isofc
>gi|1036824714|ref|XP_017137851.1| PREDICTED: protein phosphatase 1 regulatory subunit 3C isofc
>gi|1036836428|ref|XP_017138509.1| PREDICTED: COMM domain-containing protein 4 [Drosophila mira
>gi|1036836466|ref|XP_017138511.1| PREDICTED: leucokinin [Drosophila miranda]
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>gi|1036836447|ref|XP_017138510.1| PREDICTED: tctex1 domain-containing protein 2 [Drosophila mi
>gi|1036848088|ref|XP_017139150.1| PREDICTED: homeobox protein Nkx-6.1 [Drosophila miranda]
>gi|1036841523|ref|XP_017138804.1| PREDICTED: ecdysone 20-monooxygenase isoform X3 [Drosophila
>gi|1036841469|ref|XP_017138801.1| PREDICTED: ecdysone 20-monooxygenase isoform X1 [Drosophila
>gi|1036841486|ref|XP_017138802.1| PREDICTED: ecdysone 20-monooxygenase isoform X1 [Drosophila
>gi|1036841505|ref|XP_017138803.1| PREDICTED: ecdysone 20-monooxygenase isoform X2 [Drosophila
>gi|1036817367|ref|XP_017137445.1| PREDICTED: uncharacterized protein LOC108152545 isoform X1
>gi|1036817386|ref|XP_017137446.1| PREDICTED: uncharacterized protein LOC108152545 isoform X2
>gi|1036817407|ref|XP_017137447.1| PREDICTED: uncharacterized protein LOC108152545 isoform X2
>gi|1036817348|ref|XP_017137444.1| PREDICTED: ATP-dependent DNA helicase Q5 [Drosophila miranda
>gi|1036801968|ref|XP_017136600.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036802022|ref|XP_017136603.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036801984|ref|XP_017136601.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036802004|ref|XP_017136602.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036802040|ref|XP_017136605.1| PREDICTED: IQ motif and SEC7 domain-containing protein 2 iso
>gi|1036791429|ref|XP_017136001.1| PREDICTED: protein alan shepard isoform X9 [Drosophila miran
>gi|1036791502|ref|XP_017136006.1| PREDICTED: protein alan shepard isoform X13 [Drosophila mira
>gi|1036791447|ref|XP_017136002.1| PREDICTED: protein alan shepard isoform X10 [Drosophila mira
>gi|1036791393|ref|XP_017135999.1| PREDICTED: protein alan shepard isoform X7 [Drosophila miran
>gi|1036791301|ref|XP_017135994.1| PREDICTED: protein alan shepard isoform X2 [Drosophila miran
>gi|1036791412|ref|XP_017136000.1| PREDICTED: protein alan shepard isoform X8 [Drosophila miran
>gi|1036791483|ref|XP_017136005.1| PREDICTED: protein alan shepard isoform X12 [Drosophila mira
>gi|1036791375|ref|XP_017135998.1| PREDICTED: protein alan shepard isoform X6 [Drosophila miran
>gi|1036791356|ref|XP_017135997.1| PREDICTED: protein alan shepard isoform X5 [Drosophila miran
>gi|1036791320|ref|XP_017135995.1| PREDICTED: protein alan shepard isoform X3 [Drosophila miran
>gi|1036791338|ref|XP_017135996.1| PREDICTED: protein alan shepard isoform X4 [Drosophila miran
>gi|1036791266|ref|XP_017135993.1| PREDICTED: protein alan shepard isoform X1 [Drosophila miran
>gi|1036791465|ref|XP_017136004.1| PREDICTED: protein alan shepard isoform X11 [Drosophila mira
>gi|1036850180|ref|XP_017139275.1| PREDICTED: uncharacterized protein LOC108153672 [Drosophila
>gi|1036852180|ref|XP_017139395.1| PREDICTED: glutamate dehydrogenase, mitochondrial-like [Dros
>gi|1036767244|ref|XP_017134620.1| PREDICTED: uncharacterized protein LOC108150824 [Drosophila
>gi|1036767263|ref|XP_017134621.1| PREDICTED: uncharacterized protein LOC108150824 [Drosophila
>gi|1036766739|ref|XP_017134591.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036821581|ref|XP_017137672.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036759439|ref|XP_017156837.1| PREDICTED: uncharacterized protein LOC108165314 [Drosophila
>gi|1036821463|ref|XP_017137666.1| PREDICTED: uncharacterized protein LOC108152680 isoform X1
>gi|1036821484|ref|XP_017137667.1| PREDICTED: uncharacterized protein LOC108152680 isoform X2
>gi|1036821505|ref|XP_017137668.1| PREDICTED: uncharacterized protein LOC108152680 isoform X2
>gi|1036821525|ref|XP_017137669.1| PREDICTED: uncharacterized protein LOC108152681 isoform X1
>gi|1036821544|ref|XP_017137670.1| PREDICTED: uncharacterized protein LOC108152681 isoform X2
>gi|1036821444|ref|XP_017137665.1| PREDICTED: ATP-binding cassette sub-family A member 3 [Dros
>gi|1036821562|ref|XP_017137671.1| PREDICTED: uncharacterized protein LOC108152682 [Drosophila
>gi|1036799479|ref|XP_017136456.1| PREDICTED: junctional adhesion molecule A isoform X2 [Drosop
>gi|1036799460|ref|XP_017136455.1| PREDICTED: fibroblast growth factor receptor-like 1 isoform
>gi|1036799536|ref|XP_017136460.1| PREDICTED: N-alpha-acetyltransferase daf-31 [Drosophila mira
>gi|1036799554|ref|XP_017136461.1| PREDICTED: small integral membrane protein 13 [Drosophila mi
>gi|1036799517|ref|XP_017136458.1| PREDICTED: protein TSSC4 [Drosophila miranda]
>gi|1036799498|ref|XP_017136457.1| PREDICTED: caspase Dronc [Drosophila miranda]
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>gi|1036836151|ref|XP_017138493.1| PREDICTED: uncharacterized protein LOC108153189 isoform X1
>gi|1036836169|ref|XP_017138494.1| PREDICTED: uncharacterized protein LOC108153189 isoform X2
>gi|1036836186|ref|XP_017138495.1| PREDICTED: uncharacterized protein LOC108153189 isoform X3
>gi|1036836132|ref|XP_017138491.1| PREDICTED: SCY1-like protein 2 isoform X2 [Drosophila miranc
>gi|1036836113|ref|XP_017138490.1| PREDICTED: SCY1-like protein 2 isoform X1 [Drosophila miranc
>gi|1036794728|ref|XP_017136188.1| PREDICTED: nuclear distribution protein nudE homolog isoforn
>gi|1036794710|ref|XP_017136187.1| PREDICTED: nuclear distribution protein nudE homolog isoforn
>gi|1036794748|ref|XP_017136189.1| PREDICTED: type 1 phosphatidylinositol 4,5-bisphosphate 4-ph
>gi|1036794768|ref|XP_017136190.1| PREDICTED: type 1 phosphatidylinositol 4,5-bisphosphate 4-ph
>gi|1036794786|ref|XP_017136191.1| PREDICTED: type 1 phosphatidylinositol 4,5-bisphosphate 4-ph
>gi|1036794805|ref|XP_017136192.1| PREDICTED: uncharacterized protein LOC108151831 [Drosophila
>gi|1036794693|ref|XP_017136186.1| PREDICTED: calpain-B [Drosophila miranda]
>gi|1036814019|ref|XP_017137256.1| PREDICTED: transcription initiation factor TFIID subunit 2
>gi|1036814055|ref|XP_017137258.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1036814037|ref|XP_017137257.1| PREDICTED: ataxin-2 homolog isoform X1 [Drosophila miranda]
>gi|1036818985|ref|XP_017137538.1| PREDICTED: glutamic acid-rich protein [Drosophila miranda]
>gi|1036842659|ref|XP_017138857.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036842621|ref|XP_017138855.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036842640|ref|XP_017138856.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036842603|ref|XP_017138854.1| PREDICTED: uncharacterized Golgi apparatus membrane protein-
>gi|1036797036|ref|XP_017136317.1| PREDICTED: spermine oxidase [Drosophila miranda]
>gi|1036797175|ref|XP_017136326.1| PREDICTED: transmembrane protein 60 [Drosophila miranda]
>gi|1036797193|ref|XP_017136327.1| PREDICTED: transmembrane protein 60 [Drosophila miranda]
>gi|1036797017|ref|XP_017136316.1| PREDICTED: proline-rich protein PRCC [Drosophila miranda]
>gi|1036797100|ref|XP_017136321.1| PREDICTED: protein brother isoform X4 [Drosophila miranda]
>gi|1036797052|ref|XP_017136318.1| PREDICTED: protein brother isoform X1 [Drosophila miranda]
>gi|1036797085|ref|XP_017136320.1| PREDICTED: protein brother isoform X3 [Drosophila miranda]
>gi|1036797068|ref|XP_017136319.1| PREDICTED: protein brother isoform X2 [Drosophila miranda]
>gi|1036797118|ref|XP_017136322.1| PREDICTED: protein brother isoform X5 [Drosophila miranda]
>gi|1036797156|ref|XP_017136325.1| PREDICTED: transmembrane protein 35 [Drosophila miranda]
>gi|1036797137|ref|XP_017136324.1| PREDICTED: vacuolar protein sorting-associated protein VTA1
>gi|1036785913|ref|XP_017135692.1| PREDICTED: peroxisomal membrane protein 2 [Drosophila miranc
>gi|1036785929|ref|XP_017135693.1| PREDICTED: peroxisomal membrane protein 2 [Drosophila miranc
>gi|1036785816|ref|XP_017135686.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785778|ref|XP_017135684.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785833|ref|XP_017135687.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785797|ref|XP_017135685.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785852|ref|XP_017135688.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofc
>gi|1036785875|ref|XP_017135690.1| PREDICTED: uncharacterized protein C9orf78 [Drosophila mirar
>gi|1036785894|ref|XP_017135691.1| PREDICTED: 60S ribosomal protein L23a [Drosophila miranda]
>gi|1036764803|ref|XP_017157137.1| PREDICTED: ras-related protein Rab-34 [Drosophila miranda]
>gi|1036836651|ref|XP_017138522.1| PREDICTED: WD repeat-containing protein 78 [Drosophila mirar
>gi|1036836670|ref|XP_017138523.1| PREDICTED: WD repeat-containing protein 78 [Drosophila mirar
>gi|1036836616|ref|XP_017138521.1| PREDICTED: uncharacterized protein LOC108153210 [Drosophila
>gi|1036836689|ref|XP_017138524.1| PREDICTED: uncharacterized protein LOC108153212 [Drosophila
>gi|1036836710|ref|XP_017138525.1| PREDICTED: glycine-rich cell wall structural protein [Drosop
>gi|1036792113|ref|XP_017136041.1| PREDICTED: protein still life, isoform SIF type 1 isoform XS
>gi|1036792035|ref|XP_017136037.1| PREDICTED: protein still life, isoform SIF type 1 isoform X5
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>gi|1036792094|ref|XP_017136040.1| PREDICTED: protein still life, isoform SIF type 1 isoform X8
>gi|1036792187|ref|XP_017136045.1| PREDICTED: protein still life, isoform SIF type 1 isoform X
>gi|1036792168|ref|XP_017136044.1| PREDICTED: protein still life, isoform SIF type 1 isoform X
>gi|1036792054|ref|XP_017136038.1| PREDICTED: protein still life, isoform SIF type 1 isoform X
>gi|1036792150|ref|XP_017136043.1| PREDICTED: protein still life, isoform SIF type 1 isoform X
>gi|1036792132|ref|XP_017136042.1| PREDICTED: protein still life, isoform SIF type 1 isoform X
>gi|1036792073|ref|XP_017136039.1| PREDICTED: protein still life, isoform SIF type 1 isoform X
>gi|1036791927|ref|XP_017136030.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791946|ref|XP_017136031.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791964|ref|XP_017136032.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791983|ref|XP_017136034.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036792001|ref|XP_017136035.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036792018|ref|XP_017136036.1| PREDICTED: protein still life, isoform SIF type 1 isoform X4
>gi|1036791908|ref|XP_017136029.1| PREDICTED: protein still life, isoforms C/SIF type 2 isoform
>gi|1036791890|ref|XP_017136028.1| PREDICTED: protein still life, isoforms C/SIF type 2 isoform
>gi|1036791871|ref|XP_017136027.1| PREDICTED: protein still life, isoforms C/SIF type 2 isoform
>gi|1036792202|ref|XP_017136046.1| PREDICTED: LOW QUALITY PROTEIN: separin [Drosophila miranda]
>gi|1036846308|ref|XP_017139049.1| PREDICTED: protein lin-28 homolog isoform X2 [Drosophila mi
>gi|1036846290|ref|XP_017139048.1| PREDICTED: protein lin-28 homolog isoform X1 [Drosophila mi
>gi|1036768887|ref|XP_017134707.1| PREDICTED: uncharacterized protein LOC108150904 isoform X2
>gi|1036768830|ref|XP_017134704.1| PREDICTED: uncharacterized protein LOC108150904 isoform X1
>gi|1036768849|ref|XP_017134705.1| PREDICTED: uncharacterized protein LOC108150904 isoform X1
>gi|1036768868|ref|XP_017134706.1| PREDICTED: uncharacterized protein LOC108150904 isoform X1
>gi|1036769248|ref|XP_017134729.1| PREDICTED: chromatin modification-related protein MEAF6 iso
>gi|1036769267|ref|XP_017134730.1| PREDICTED: chromatin modification-related protein MEAF6 iso
>gi|1036769286|ref|XP_017134731.1| PREDICTED: chromatin modification-related protein MEAF6 iso
>gi|1036769122|ref|XP_017134721.1| PREDICTED: transmembrane 9 superfamily member 3 [Drosophila
>gi|1036769194|ref|XP_017134726.1| PREDICTED: collagen alpha-2(IX) chain [Drosophila miranda]
>gi|1036769159|ref|XP_017134723.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila
>gi|1036769140|ref|XP_017134722.1| PREDICTED: membrane-bound alkaline phosphatase [Drosophila
>gi|1036759457|ref|XP_017156838.1| PREDICTED: cytochrome P450 307a1 [Drosophila miranda]
>gi|1036768794|ref|XP_017134702.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1036768811|ref|XP_017134703.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1036769086|ref|XP_017134719.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036769105|ref|XP_017134720.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036769305|ref|XP_017134733.1| PREDICTED: bladder cancer-associated protein [Drosophila mi
>gi|1036769322|ref|XP_017134734.1| PREDICTED: bladder cancer-associated protein [Drosophila mi
>gi|1036769213|ref|XP_017134727.1| PREDICTED: CAAX prenyl protease 2 [Drosophila miranda]
>gi|1036769176|ref|XP_017134725.1| PREDICTED: succinyl-CoA ligase [GDP-forming] subunit beta, 
>gi|1036769231|ref|XP_017134728.1| PREDICTED: uncharacterized protein LOC108150914 [Drosophila
>gi|1036768904|ref|XP_017134708.1| PREDICTED: collagen alpha-5(IV) chain isoform X1 [Drosophila
>gi|1036768922|ref|XP_017134709.1| PREDICTED: collagen alpha-5(IV) chain isoform X1 [Drosophila
>gi|1036768941|ref|XP_017134710.1| PREDICTED: collagen alpha-5(IV) chain isoform X2 [Drosophila
>gi|1036768960|ref|XP_017134711.1| PREDICTED: collagen alpha-1(X) chain isoform X3 [Drosophila
>gi|1036769001|ref|XP_017134713.1| PREDICTED: collagen alpha-1(X) chain isoform X5 [Drosophila
>gi|1036768982|ref|XP_017134712.1| PREDICTED: collagen alpha-1(I) chain isoform X4 [Drosophila
>gi|1036769018|ref|XP_017134714.1| PREDICTED: collagen alpha-1(I) chain isoform X6 [Drosophila
>gi|1036769035|ref|XP_017134716.1| PREDICTED: collagen alpha-1(I) chain isoform X6 [Drosophila
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>gi|1036769051|ref|XP_017134717.1| PREDICTED: collagen alpha-1(III) chain isoform X7 [Drosophil
>gi|1036769067|ref|XP_017134718.1| PREDICTED: collagen alpha-1(V) chain isoform X8 [Drosophila
>gi|1036766665|ref|XP_017134586.1| PREDICTED: prisilkin-39 isoform X2 [Drosophila miranda]
>gi|1036766646|ref|XP_017134585.1| PREDICTED: prisilkin-39 isoform X1 [Drosophila miranda]
>gi|1036852398|ref|XP_017139409.1| PREDICTED: neuropeptide Y receptor type 1 [Drosophila miran
>gi|1036768612|ref|XP_017134690.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosop
>gi|1036767126|ref|XP_017134614.1| PREDICTED: uncharacterized protein LOC108150818 [Drosophila
>gi|1036759475|ref|XP_017156839.1| PREDICTED: uncharacterized protein LOC108165316 [Drosophila
>gi|1036784396|ref|XP_017135603.1| PREDICTED: probable prefoldin subunit 4 [Drosophila miranda]
>gi|1036784358|ref|XP_017135601.1| PREDICTED: probable Golgi SNAP receptor complex member 2 iso
>gi|1036784341|ref|XP_017135600.1| PREDICTED: probable Golgi SNAP receptor complex member 2 iso
>gi|1036784377|ref|XP_017135602.1| PREDICTED: ubiquitin-conjugating enzyme E2 variant 2 [Drosop
>gi|1036784322|ref|XP_017135599.1| PREDICTED: cytochrome c1, heme protein, mitochondrial [Droso
>gi|1036784268|ref|XP_017135595.1| PREDICTED: kinesin-like protein KIF3A [Drosophila miranda]
>gi|1036784287|ref|XP_017135597.1| PREDICTED: putative phospholipase B-like lamina ancestor [Dr
>gi|1036784306|ref|XP_017135598.1| PREDICTED: putative phospholipase B-like lamina ancestor [Dr
>gi|1036843783|ref|XP_017138925.1| PREDICTED: protein alan shepard [Drosophila miranda]
>gi|1036843802|ref|XP_017138926.1| PREDICTED: protein alan shepard [Drosophila miranda]
>gi|1036759494|ref|XP_017156841.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036765309|ref|XP_017157163.1| PREDICTED: lachesin [Drosophila miranda]
>gi|1036766794|ref|XP_017134594.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036767206|ref|XP_017134618.1| PREDICTED: uncharacterized protein LOC108150823 [Drosophila
>gi|1036767225|ref|XP_017134619.1| PREDICTED: uncharacterized protein LOC108150823 [Drosophila
>gi|1036843397|ref|XP_017138902.1| PREDICTED: uncharacterized protein LOC108153413 isoform X3
>gi|1036843338|ref|XP_017138897.1| PREDICTED: uncharacterized protein LOC108153413 isoform X1
>gi|1036843357|ref|XP_017138898.1| PREDICTED: uncharacterized protein LOC108153413 isoform X1
>gi|1036843377|ref|XP_017138901.1| PREDICTED: uncharacterized protein LOC108153413 isoform X2
>gi|1036843452|ref|XP_017138905.1| PREDICTED: uncharacterized protein LOC108153415 [Drosophila
>gi|1036843414|ref|XP_017138903.1| PREDICTED: uncharacterized protein LOC108153413 isoform X4
>gi|1036843433|ref|XP_017138904.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1036849873|ref|XP_017139256.1| PREDICTED: nucleolin [Drosophila miranda]
>gi|1036849892|ref|XP_017139257.1| PREDICTED: nucleolin [Drosophila miranda]
>gi|1036830552|ref|XP_017138183.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036830644|ref|XP_017138188.1| PREDICTED: phospholipid scramblase 1-like isoform X2 [Drosop
>gi|1036830624|ref|XP_017138187.1| PREDICTED: phospholipid scramblase 1-like isoform X1 [Drosop
>gi|1036830570|ref|XP_017138184.1| PREDICTED: phospholipid scramblase 2-like isoform X1 [Drosop
>gi|1036830586|ref|XP_017138185.1| PREDICTED: phospholipid scramblase 2-like isoform X2 [Drosop
>gi|1036830605|ref|XP_017138186.1| PREDICTED: phospholipid scramblase 1-like isoform X3 [Drosop
>gi|1036825593|ref|XP_017137894.1| PREDICTED: odorant receptor 67c [Drosophila miranda]
>gi|1036825554|ref|XP_017137891.1| PREDICTED: odorant receptor 67d isoform X1 [Drosophila miran
>gi|1036825572|ref|XP_017137892.1| PREDICTED: odorant receptor 67d isoform X2 [Drosophila miran
>gi|1036825633|ref|XP_017137896.1| PREDICTED: uncharacterized protein LOC108152818 isoform X2
>gi|1036825652|ref|XP_017137897.1| PREDICTED: uncharacterized protein LOC108152818 isoform X3
>gi|1036825614|ref|XP_017137895.1| PREDICTED: uncharacterized protein LOC108152818 isoform X1
>gi|1036851326|ref|XP_017139341.1| PREDICTED: protein ariadne-1 [Drosophila miranda]
>gi|1036759513|ref|XP_017156842.1| PREDICTED: uncharacterized protein LOC108165319 [Drosophila
>gi|1036759532|ref|XP_017156843.1| PREDICTED: uncharacterized protein LOC108165320 [Drosophila
>gi|1036776402|ref|XP_017135128.1| PREDICTED: probable ATP-dependent RNA helicase DDX20 [Drosop
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>gi|1036776855|ref|XP_017135156.1| PREDICTED: uncharacterized protein LOC108151197 [Drosophila
>gi|1036776838|ref|XP_017135155.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1036776687|ref|XP_017135145.1| PREDICTED: AF4/FMR2 family member 4 isoform X14 [Drosophila
>gi|1036776721|ref|XP_017135147.1| PREDICTED: homeotic protein female sterile isoform X16 [Dros
>gi|1036776637|ref|XP_017135142.1| PREDICTED: AF4/FMR2 family member 4 isoform X11 [Drosophila
>gi|1036776668|ref|XP_017135144.1| PREDICTED: AF4/FMR2 family member 4 isoform X13 [Drosophila
>gi|1036776421|ref|XP_017135129.1| PREDICTED: ecdysone-induced protein 74EF isoform X1 [Drosoph
>gi|1036776551|ref|XP_017135136.1| PREDICTED: ecdysone-induced protein 74EF isoform X6 [Drosoph
>gi|1036776532|ref|XP_017135135.1| PREDICTED: ecdysone-induced protein 74EF isoform X5 [Drosoph
>gi|1036776588|ref|XP_017135138.1| PREDICTED: ecdysone-induced protein 74EF isoform X8 [Drosoph
>gi|1036776651|ref|XP_017135143.1| PREDICTED: AF4/FMR2 family member 4 isoform X12 [Drosophila
>gi|1036776440|ref|XP_017135130.1| PREDICTED: ecdysone-induced protein 74EF isoform X2 [Drosoph
>gi|1036776570|ref|XP_017135137.1| PREDICTED: ecdysone-induced protein 74EF isoform X7 [Drosoph
>gi|1036776607|ref|XP_017135139.1| PREDICTED: ecdysone-induced protein 74EF isoform X9 [Drosoph
>gi|1036776623|ref|XP_017135140.1| PREDICTED: ecdysone-induced protein 74EF isoform X10 [Drosop
>gi|1036776804|ref|XP_017135153.1| PREDICTED: ecdysone-induced protein 74EF isoform X19 [Drosop
>gi|1036776821|ref|XP_017135154.1| PREDICTED: ecdysone-induced protein 74EF isoform X20 [Drosop
>gi|1036776459|ref|XP_017135131.1| PREDICTED: ecdysone-induced protein 74EF isoform X3 [Drosoph
>gi|1036776738|ref|XP_017135148.1| PREDICTED: homeotic protein female sterile isoform X17 [Dros
>gi|1036776755|ref|XP_017135149.1| PREDICTED: homeotic protein female sterile isoform X17 [Dros
>gi|1036776771|ref|XP_017135150.1| PREDICTED: homeotic protein female sterile isoform X17 [Dros
>gi|1036776787|ref|XP_017135151.1| PREDICTED: homeotic protein female sterile isoform X18 [Dros
>gi|1036776478|ref|XP_017135132.1| PREDICTED: ecdysone-induced protein 74EF isoform X4 [Drosoph
>gi|1036776497|ref|XP_017135133.1| PREDICTED: ecdysone-induced protein 74EF isoform X4 [Drosoph
>gi|1036776513|ref|XP_017135134.1| PREDICTED: ecdysone-induced protein 74EF isoform X4 [Drosoph
>gi|1036776703|ref|XP_017135146.1| PREDICTED: ecdysone-induced protein 74EF isoform X15 [Drosop
>gi|1036776872|ref|XP_017135158.1| PREDICTED: uncharacterized protein LOC108151198 [Drosophila
>gi|1036759551|ref|XP_017156844.1| PREDICTED: cytosolic non-specific dipeptidase-like [Drosophi
>gi|1036852379|ref|XP_017139407.1| PREDICTED: UNC93-like protein isoform X2 [Drosophila miranda
>gi|1036852360|ref|XP_017139406.1| PREDICTED: UNC93-like protein isoform X1 [Drosophila miranda
>gi|1036766939|ref|XP_017134603.1| PREDICTED: uncharacterized protein LOC108150809 isoform X2
>gi|1036766920|ref|XP_017134602.1| PREDICTED: uncharacterized protein LOC108150809 isoform X1
>gi|1036768649|ref|XP_017134693.1| PREDICTED: uncharacterized protein LOC108150894 [Drosophila
>gi|1036767484|ref|XP_017134632.1| PREDICTED: uncharacterized protein LOC108150836 [Drosophila
>gi|1036765364|ref|XP_017157166.1| PREDICTED: uncharacterized protein LOC108165603 isoform X1
>gi|1036765378|ref|XP_017157167.1| PREDICTED: uncharacterized protein LOC108165603 isoform X2
>gi|1036765393|ref|XP_017157168.1| PREDICTED: uncharacterized protein LOC108165603 isoform X3
>gi|1036759570|ref|XP_017156845.1| PREDICTED: uncharacterized protein LOC108165322, partial [D
>gi|1036782761|ref|XP_017135506.1| PREDICTED: chromosome transmission fidelity protein 8 homolo
>gi|1036782691|ref|XP_017135502.1| PREDICTED: DNA excision repair protein haywire [Drosophila m
>gi|1036782709|ref|XP_017135503.1| PREDICTED: histone-lysine N-methyltransferase E(z) [Drosophi
>gi|1036782779|ref|XP_017135508.1| PREDICTED: nuclear envelope phosphatase-regulatory subunit 1
>gi|1036782728|ref|XP_017135504.1| PREDICTED: protein FAM49B isoform X1 [Drosophila miranda]
>gi|1036782745|ref|XP_017135505.1| PREDICTED: protein FAM49B isoform X2 [Drosophila miranda]
>gi|1036804980|ref|XP_017136766.1| PREDICTED: ankyrin repeat and MYND domain-containing protei
>gi|1036804999|ref|XP_017136767.1| PREDICTED: 1,2-dihydroxy-3-keto-5-methylthiopentene dioxygen
>gi|1036804942|ref|XP_017136764.1| PREDICTED: ras guanine nucleotide exchange factor P isoform
>gi|1036804961|ref|XP_017136765.1| PREDICTED: ras guanine nucleotide exchange factor P isoform
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>gi|1036805034|ref|XP_017136769.1| PREDICTED: U4/U6 small nuclear ribonucleoprotein Prp4 [Dros
>gi|1036805013|ref|XP_017136768.1| PREDICTED: solute carrier organic anion transporter family n
>gi|1036805088|ref|XP_017136772.1| PREDICTED: uncharacterized protein LOC108152164 [Drosophila
>gi|1036759588|ref|XP_017156846.1| PREDICTED: uncharacterized protein LOC108165323 [Drosophila
>gi|1036805071|ref|XP_017136771.1| PREDICTED: uncharacterized protein LOC108152163 [Drosophila
>gi|1036805053|ref|XP_017136770.1| PREDICTED: cytochrome b-c1 complex subunit 8 [Drosophila mir
>gi|1036848950|ref|XP_017139200.1| PREDICTED: glycine receptor subunit alphaZ1 [Drosophila mira
>gi|1036847401|ref|XP_017139113.1| PREDICTED: MICOS complex subunit MIC13 homolog QIL1 [Drosoph
>gi|1036795780|ref|XP_017136248.1| PREDICTED: uncharacterized protein LOC108151867 isoform X2
>gi|1036795727|ref|XP_017136245.1| PREDICTED: uncharacterized protein LOC108151867 isoform X1
>gi|1036795745|ref|XP_017136246.1| PREDICTED: uncharacterized protein LOC108151867 isoform X1
>gi|1036795763|ref|XP_017136247.1| PREDICTED: uncharacterized protein LOC108151867 isoform X1
>gi|1036795799|ref|XP_017136249.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1036795817|ref|XP_017136250.1| PREDICTED: uncharacterized protein LOC108151870 [Drosophila
>gi|1036795836|ref|XP_017136251.1| PREDICTED: UDP-sugar transporter UST74c [Drosophila miranda]
>gi|1036795874|ref|XP_017136254.1| PREDICTED: ubiquinone biosynthesis protein COQ4 homolog, mit
>gi|1036795893|ref|XP_017136255.1| PREDICTED: ubiquinone biosynthesis protein COQ4 homolog, mit
>gi|1036795912|ref|XP_017136256.1| PREDICTED: ubiquinone biosynthesis protein COQ4 homolog, mit
>gi|1036795855|ref|XP_017136252.1| PREDICTED: RAB6-interacting golgin [Drosophila miranda]
>gi|1036769992|ref|XP_017134770.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC7 [Dro
>gi|1036770030|ref|XP_017134772.1| PREDICTED: uncharacterized protein LOC108150950 isoform X1
>gi|1036770047|ref|XP_017134773.1| PREDICTED: uncharacterized protein LOC108150950 isoform X2
>gi|1036769895|ref|XP_017134765.1| PREDICTED: exocyst complex component 1 [Drosophila miranda]
>gi|1036769857|ref|XP_017134762.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036769876|ref|XP_017134764.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036769916|ref|XP_017134766.1| PREDICTED: LOW QUALITY PROTEIN: proline-rich extensin-like p
>gi|1036759606|ref|XP_017156847.1| PREDICTED: uncharacterized protein LOC108165324 [Drosophila
>gi|1036769741|ref|XP_017134756.1| PREDICTED: cadherin-87A [Drosophila miranda]
>gi|1036769760|ref|XP_017134757.1| PREDICTED: cadherin-87A [Drosophila miranda]
>gi|1036769954|ref|XP_017134768.1| PREDICTED: basic helix-loop-helix neural transcription facto
>gi|1036769799|ref|XP_017134759.1| PREDICTED: uncharacterized protein LOC108150939 isoform X2
>gi|1036769779|ref|XP_017134758.1| PREDICTED: uncharacterized protein LOC108150939 isoform X1
>gi|1036769818|ref|XP_017134760.1| PREDICTED: uncharacterized protein LOC108150939 isoform X3
>gi|1036769837|ref|XP_017134761.1| PREDICTED: uncharacterized protein LOC108150939 isoform X4
>gi|1036770011|ref|XP_017134771.1| PREDICTED: allatostatins MIP [Drosophila miranda]
>gi|1036759625|ref|XP_017156848.1| PREDICTED: polycystic kidney disease 2-like 2 protein [Dros
>gi|1036759642|ref|XP_017156849.1| PREDICTED: uncharacterized protein LOC108165327 [Drosophila
>gi|1036769935|ref|XP_017134767.1| PREDICTED: transmembrane protein 209 [Drosophila miranda]
>gi|1036769973|ref|XP_017134769.1| PREDICTED: odorant receptor 74a [Drosophila miranda]
>gi|1036759658|ref|XP_017156850.1| PREDICTED: uncharacterized protein LOC108165328 [Drosophila
>gi|1036766761|ref|XP_017134592.1| PREDICTED: NADH-quinone oxidoreductase subunit E [Drosophila
>gi|1036766443|ref|XP_017134575.1| PREDICTED: uncharacterized protein LOC108150785 [Drosophila
>gi|1036832514|ref|XP_017138291.1| PREDICTED: UPF0687 protein C20orf27 homolog [Drosophila mira
>gi|1036792425|ref|XP_017136059.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792504|ref|XP_017136064.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792388|ref|XP_017136057.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792406|ref|XP_017136058.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792484|ref|XP_017136063.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
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>gi|1036792582|ref|XP_017136068.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792627|ref|XP_017136071.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792465|ref|XP_017136062.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792616|ref|XP_017136070.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792563|ref|XP_017136067.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
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>gi|1036792444|ref|XP_017136060.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792523|ref|XP_017136065.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792542|ref|XP_017136066.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792682|ref|XP_017136074.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036792646|ref|XP_017136072.1| PREDICTED: arginine-glutamic acid dipeptide repeats protein
>gi|1036818174|ref|XP_017137491.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila
>gi|1036818193|ref|XP_017137492.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila
>gi|1036818212|ref|XP_017137493.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila
>gi|1036818231|ref|XP_017137494.1| PREDICTED: tetratricopeptide repeat protein 28 [Drosophila
>gi|1036773858|ref|XP_017134991.1| PREDICTED: zinc finger CCCH domain-containing protein 3 [Dro
>gi|1036773747|ref|XP_017134984.1| PREDICTED: pseudouridylate synthase 7 homolog [Drosophila mi
>gi|1036773982|ref|XP_017134998.1| PREDICTED: uncharacterized protein LOC108151109 [Drosophila
>gi|1036773802|ref|XP_017134987.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036773821|ref|XP_017134988.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036773839|ref|XP_017134989.1| PREDICTED: protein glass [Drosophila miranda]
>gi|1036773963|ref|XP_017134997.1| PREDICTED: pyrimidodiazepine synthase [Drosophila miranda]
>gi|1036773875|ref|XP_017134992.1| PREDICTED: pyrimidodiazepine synthase [Drosophila miranda]
>gi|1036773946|ref|XP_017134996.1| PREDICTED: pyrimidodiazepine synthase isoform X2 [Drosophila
>gi|1036773928|ref|XP_017134995.1| PREDICTED: pyrimidodiazepine synthase isoform X1 [Drosophila
>gi|1036773894|ref|XP_017134993.1| PREDICTED: pyrimidodiazepine synthase isoform X1 [Drosophila
>gi|1036773909|ref|XP_017134994.1| PREDICTED: pyrimidodiazepine synthase isoform X2 [Drosophila
>gi|1036773766|ref|XP_017134985.1| PREDICTED: zinc transporter foi [Drosophila miranda]
>gi|1036773785|ref|XP_017134986.1| PREDICTED: zinc transporter foi [Drosophila miranda]
>gi|1036790045|ref|XP_017135925.1| PREDICTED: zinc finger protein 345 [Drosophila miranda]
>gi|1036789994|ref|XP_017135922.1| PREDICTED: E3 ubiquitin-protein ligase SHPRH isoform X1 [Dro
>gi|1036790011|ref|XP_017135923.1| PREDICTED: E3 ubiquitin-protein ligase SHPRH isoform X2 [Dro
>gi|1036790027|ref|XP_017135924.1| PREDICTED: zinc finger protein 91 [Drosophila miranda]
>gi|1036801261|ref|XP_017136558.1| PREDICTED: serine/threonine-protein phosphatase 6 regulatory
>gi|1036801280|ref|XP_017136559.1| PREDICTED: serine/threonine-protein phosphatase 6 regulatory
>gi|1036801301|ref|XP_017136560.1| PREDICTED: serine/threonine-protein phosphatase 6 regulatory
>gi|1036764770|ref|XP_017157135.1| PREDICTED: Krueppel-like factor 7 [Drosophila miranda]
>gi|1036764787|ref|XP_017157136.1| PREDICTED: pupal cuticle protein 20 [Drosophila miranda]
>gi|1036768630|ref|XP_017134692.1| PREDICTED: uncharacterized protein LOC108150893 [Drosophila
>gi|1036768004|ref|XP_017134658.1| PREDICTED: uncharacterized protein LOC108150866 [Drosophila
>gi|1036765832|ref|XP_017134542.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036765852|ref|XP_017134543.1| PREDICTED: endocuticle structural glycoprotein ABD-5 [Drosop
>gi|1036767559|ref|XP_017134636.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036764187|ref|XP_017157105.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036764205|ref|XP_017157106.1| PREDICTED: larval cuticle protein 5-like [Drosophila miranda
>gi|1036766577|ref|XP_017134583.1| PREDICTED: larval cuticle protein 8 [Drosophila miranda]
>gi|1036767795|ref|XP_017134645.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
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>gi|1036766175|ref|XP_017134562.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036767353|ref|XP_017134625.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036766902|ref|XP_017134601.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036766777|ref|XP_017134593.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036766721|ref|XP_017134590.1| PREDICTED: endocuticle structural protein SgAbd-6 [Drosophil
>gi|1036764353|ref|XP_017157115.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036764373|ref|XP_017157116.1| PREDICTED: larval cuticle protein 8-like [Drosophila miranda
>gi|1036849855|ref|XP_017139255.1| PREDICTED: protein lethal(3)malignant blood neoplasm 1 [Dros
>gi|1036849193|ref|XP_017139214.1| PREDICTED: DNA-binding protein D-ETS-3 isoform X1 [Drosophil
>gi|1036849211|ref|XP_017139215.1| PREDICTED: DNA-binding protein D-ETS-3 isoform X2 [Drosophil
>gi|1036759675|ref|XP_017156852.1| PREDICTED: DNA-binding protein D-ETS-3-like [Drosophila mira
>gi|1036842968|ref|XP_017138877.1| PREDICTED: transmembrane emp24 domain-containing protein B
>gi|1036797790|ref|XP_017136360.1| PREDICTED: aldose 1-epimerase [Drosophila miranda]
>gi|1036797810|ref|XP_017136361.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036797677|ref|XP_017136354.1| PREDICTED: protein SCAI isoform X4 [Drosophila miranda]
>gi|1036797644|ref|XP_017136352.1| PREDICTED: protein SCAI isoform X2 [Drosophila miranda]
>gi|1036797661|ref|XP_017136353.1| PREDICTED: protein SCAI isoform X3 [Drosophila miranda]
>gi|1036797613|ref|XP_017136350.1| PREDICTED: protein SCAI isoform X1 [Drosophila miranda]
>gi|1036797628|ref|XP_017136351.1| PREDICTED: protein SCAI isoform X1 [Drosophila miranda]
>gi|1036797576|ref|XP_017136348.1| PREDICTED: uncharacterized protein LOC108151934 [Drosophila
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>gi|1036797826|ref|XP_017136362.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm5 [Drosop
>gi|1036797710|ref|XP_017136357.1| PREDICTED: T-cell activation inhibitor, mitochondrial isofor
>gi|1036797694|ref|XP_017136356.1| PREDICTED: T-cell activation inhibitor, mitochondrial isofor
>gi|1036797752|ref|XP_017136358.1| PREDICTED: T-cell activation inhibitor, mitochondrial isofor
>gi|1036797771|ref|XP_017136359.1| PREDICTED: uncharacterized protein LOC108151939 [Drosophila
>gi|1036815454|ref|XP_017137337.1| PREDICTED: organic cation transporter protein isoform X1 [Dr
>gi|1036815473|ref|XP_017137338.1| PREDICTED: organic cation transporter protein isoform X2 [Dr
>gi|1036815503|ref|XP_017137339.1| PREDICTED: organic cation transporter protein isoform X3 [Dr
>gi|1036815567|ref|XP_017137342.1| PREDICTED: brachyurin-like [Drosophila miranda]
>gi|1036759695|ref|XP_017156853.1| PREDICTED: LOW QUALITY PROTEIN: brachyurin-like [Drosophila
>gi|1036815525|ref|XP_017137340.1| PREDICTED: transmembrane protease serine 12 [Drosophila mir
>gi|1036815548|ref|XP_017137341.1| PREDICTED: protein tantalus [Drosophila miranda]
>gi|1036815646|ref|XP_017137347.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036815628|ref|XP_017137346.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036815605|ref|XP_017137345.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036815588|ref|XP_017137344.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036759714|ref|XP_017156854.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036850286|ref|XP_017139280.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036759732|ref|XP_017156855.1| PREDICTED: brachyurin [Drosophila miranda]
>gi|1036846510|ref|XP_017139062.1| PREDICTED: tensin-1 isoform X2 [Drosophila miranda]
>gi|1036846472|ref|XP_017139060.1| PREDICTED: protein sprint isoform X1 [Drosophila miranda]
>gi|1036846491|ref|XP_017139061.1| PREDICTED: protein sprint isoform X1 [Drosophila miranda]
>gi|1036846529|ref|XP_017139063.1| PREDICTED: serine protease 3-like [Drosophila miranda]
>gi|1036846548|ref|XP_017139064.1| PREDICTED: serine protease 3-like [Drosophila miranda]
>gi|1036846567|ref|XP_017139065.1| PREDICTED: serine protease 3-like [Drosophila miranda]
>gi|1036828928|ref|XP_017138088.1| PREDICTED: tensin-4 isoform X1 [Drosophila miranda]
>gi|1036828947|ref|XP_017138089.1| PREDICTED: uncharacterized protein LOC108152920 isoform X2
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>gi|1036833857|ref|XP_017138359.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036833874|ref|XP_017138360.1| PREDICTED: DNA N6-methyl adenine demethylase [Drosophila mi
>gi|1036836748|ref|XP_017138527.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036836765|ref|XP_017138529.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036836838|ref|XP_017138533.1| PREDICTED: peptidoglycan-recognition protein LD isoform X4
>gi|1036836819|ref|XP_017138532.1| PREDICTED: peptidoglycan-recognition protein LD isoform X3
>gi|1036836801|ref|XP_017138531.1| PREDICTED: peptidoglycan-recognition protein LD isoform X2
>gi|1036836783|ref|XP_017138530.1| PREDICTED: peptidoglycan-recognition protein LD isoform X1
>gi|1036836857|ref|XP_017138534.1| PREDICTED: peptidoglycan-recognition protein LD isoform X5
>gi|1036836878|ref|XP_017138535.1| PREDICTED: transmembrane protein 11 homolog, mitochondrial i
>gi|1036836897|ref|XP_017138536.1| PREDICTED: transmembrane protein 11 homolog, mitochondrial i
>gi|1036837237|ref|XP_017138556.1| PREDICTED: membrane-associated tyrosine- and threonine-spec
>gi|1036837256|ref|XP_017138557.1| PREDICTED: membrane-associated tyrosine- and threonine-spec
>gi|1036803499|ref|XP_017136683.1| PREDICTED: nucleolar protein 12 [Drosophila miranda]
>gi|1036803442|ref|XP_017136680.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 36 isoform
>gi|1036803460|ref|XP_017136681.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 36 isoform
>gi|1036817291|ref|XP_017137441.1| PREDICTED: bifunctional coenzyme A synthase [Drosophila mira
>gi|1036817329|ref|XP_017137443.1| PREDICTED: thioredoxin-like protein 1 [Drosophila miranda]
>gi|1036817310|ref|XP_017137442.1| PREDICTED: proliferation-associated protein 2G4 [Drosophila
>gi|1036805557|ref|XP_017136799.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 64E [Drosop
>gi|1036805576|ref|XP_017136800.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 64E [Drosop
>gi|1036805595|ref|XP_017136801.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 64E [Drosop
>gi|1036844482|ref|XP_017138961.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036844499|ref|XP_017138962.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036829057|ref|XP_017138096.1| PREDICTED: R3H and coiled-coil domain-containing protein 1
>gi|1036829074|ref|XP_017138097.1| PREDICTED: R3H and coiled-coil domain-containing protein 1
>gi|1036826962|ref|XP_017137975.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036826981|ref|XP_017137976.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036827000|ref|XP_017137977.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036826923|ref|XP_017137973.1| PREDICTED: phospholipid scramblase 1 isoform X1 [Drosophila
>gi|1036826941|ref|XP_017137974.1| PREDICTED: phospholipid scramblase 2 isoform X2 [Drosophila
>gi|1036839630|ref|XP_017138696.1| PREDICTED: ankyrin repeat domain-containing protein 54 [Dros
>gi|1036827038|ref|XP_017137979.1| PREDICTED: 85/88 kDa calcium-independent phospholipase A2 is
>gi|1036827019|ref|XP_017137978.1| PREDICTED: 85/88 kDa calcium-independent phospholipase A2 is
>gi|1036808148|ref|XP_017136940.1| PREDICTED: protein phosphatase inhibitor 2 isoform X1 [Droso
>gi|1036808167|ref|XP_017136941.1| PREDICTED: protein phosphatase inhibitor 2 isoform X2 [Droso
>gi|1036808075|ref|XP_017136936.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1036808020|ref|XP_017136933.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808039|ref|XP_017136934.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808001|ref|XP_017136932.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808111|ref|XP_017136938.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808056|ref|XP_017136935.1| PREDICTED: putative uncharacterized protein DDB_G0271606 iso
>gi|1036808092|ref|XP_017136937.1| PREDICTED: uncharacterized protein LOC108152243 isoform X6
>gi|1036808271|ref|XP_017136947.1| PREDICTED: probable insulin-like peptide 5 [Drosophila miran
>gi|1036808130|ref|XP_017136939.1| PREDICTED: acid sphingomyelinase-like phosphodiesterase 3b
>gi|1036808220|ref|XP_017136944.1| PREDICTED: probable insulin-like peptide 4 isoform X1 [Droso
>gi|1036808236|ref|XP_017136945.1| PREDICTED: probable insulin-like peptide 4 isoform X2 [Droso
>gi|1036808253|ref|XP_017136946.1| PREDICTED: probable insulin-like peptide 4 isoform X3 [Droso
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>gi|1036808290|ref|XP_017136948.1| PREDICTED: probable insulin-like peptide 3 [Drosophila miran
>gi|1036808202|ref|XP_017136943.1| PREDICTED: probable insulin-like peptide 2 [Drosophila miran
>gi|1036808184|ref|XP_017136942.1| PREDICTED: probable insulin-like peptide 1 [Drosophila miran
>gi|1036807925|ref|XP_017136927.1| PREDICTED: synaptopodin 2-like protein isoform X1 [Drosophil
>gi|1036807944|ref|XP_017136929.1| PREDICTED: synaptopodin 2-like protein isoform X2 [Drosophil
>gi|1036807963|ref|XP_017136930.1| PREDICTED: synaptopodin 2-like protein isoform X3 [Drosophil
>gi|1036807981|ref|XP_017136931.1| PREDICTED: synaptopodin 2-like protein isoform X4 [Drosophil
>gi|1036822438|ref|XP_017137720.1| PREDICTED: anion exchange protein 3 isoform X2 [Drosophila r
>gi|1036822402|ref|XP_017137718.1| PREDICTED: anion exchange protein 2 isoform X1 [Drosophila r
>gi|1036822419|ref|XP_017137719.1| PREDICTED: anion exchange protein 2 isoform X1 [Drosophila r
>gi|1036820285|ref|XP_017137608.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1036820302|ref|XP_017137609.1| PREDICTED: homeobox protein prospero isoform X2 [Drosophila
>gi|1036820318|ref|XP_017137610.1| PREDICTED: homeobox protein prospero isoform X3 [Drosophila
>gi|1036820335|ref|XP_017137611.1| PREDICTED: homeobox protein prospero isoform X4 [Drosophila
>gi|1036820388|ref|XP_017137614.1| PREDICTED: homeobox protein prospero isoform X7 [Drosophila
>gi|1036820405|ref|XP_017137615.1| PREDICTED: homeobox protein prospero isoform X8 [Drosophila
>gi|1036820424|ref|XP_017137616.1| PREDICTED: homeobox protein prospero isoform X9 [Drosophila
>gi|1036820353|ref|XP_017137612.1| PREDICTED: homeobox protein prospero isoform X5 [Drosophila
>gi|1036820463|ref|XP_017137619.1| PREDICTED: homeobox protein prospero isoform X11 [Drosophila
>gi|1036820369|ref|XP_017137613.1| PREDICTED: homeobox protein prospero isoform X6 [Drosophila
>gi|1036820443|ref|XP_017137617.1| PREDICTED: homeobox protein prospero isoform X10 [Drosophila
>gi|1036820482|ref|XP_017137620.1| PREDICTED: insulin-like growth factor-binding protein comple
>gi|1036820499|ref|XP_017137621.1| PREDICTED: ATP synthase subunit b, mitochondrial [Drosophila
>gi|1036837200|ref|XP_017138554.1| PREDICTED: N-alpha-acetyltransferase 60 isoform X2 [Drosophi
>gi|1036837164|ref|XP_017138552.1| PREDICTED: N-alpha-acetyltransferase 60 isoform X1 [Drosophi
>gi|1036837182|ref|XP_017138553.1| PREDICTED: N-alpha-acetyltransferase 60 isoform X1 [Drosophi
>gi|1036759752|ref|XP_017156856.1| PREDICTED: integrator complex subunit 7 [Drosophila miranda]
>gi|1036817195|ref|XP_017137435.1| PREDICTED: nibrin [Drosophila miranda]
>gi|1036817252|ref|XP_017137438.1| PREDICTED: uncharacterized protein LOC108152539 [Drosophila
>gi|1036817270|ref|XP_017137439.1| PREDICTED: adrenodoxin [Drosophila miranda]
>gi|1036817214|ref|XP_017137436.1| PREDICTED: uncharacterized protein LOC108152538 isoform X1
>gi|1036817233|ref|XP_017137437.1| PREDICTED: uncharacterized protein LOC108152538 isoform X2
>gi|1036793701|ref|XP_017136128.1| PREDICTED: claspin [Drosophila miranda]
>gi|1036793757|ref|XP_017136131.1| PREDICTED: peroxisomal membrane protein PMP34 [Drosophila mi
>gi|1036793739|ref|XP_017136130.1| PREDICTED: THUMP domain-containing protein 1 homolog [Drosop
>gi|1036793720|ref|XP_017136129.1| PREDICTED: lysine-specific demethylase 8 [Drosophila miranda
>gi|1036793773|ref|XP_017136133.1| PREDICTED: B-cell receptor-associated protein 31 [Drosophila
>gi|1036793795|ref|XP_017136134.1| PREDICTED: B-cell receptor-associated protein 31 [Drosophila
>gi|1036793812|ref|XP_017136135.1| PREDICTED: protein DPCD [Drosophila miranda]
>gi|1036793868|ref|XP_017136136.1| PREDICTED: uncharacterized protein LOC108151797 [Drosophila
>gi|1036842984|ref|XP_017138878.1| PREDICTED: UPF0415 protein C7orf25 homolog [Drosophila miran
>gi|1036834602|ref|XP_017138400.1| PREDICTED: uncharacterized protein LOC108153123 [Drosophila
>gi|1036834585|ref|XP_017138399.1| PREDICTED: dimethyladenosine transferase 1, mitochondrial [D
>gi|1036834621|ref|XP_017138402.1| PREDICTED: cytochrome c oxidase assembly factor 3, mitochond
>gi|1036816225|ref|XP_017137379.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816244|ref|XP_017137381.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816263|ref|XP_017137382.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
>gi|1036816282|ref|XP_017137383.1| PREDICTED: F-box only protein 32 [Drosophila miranda]
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>gi|1036816152|ref|XP_017137375.1| PREDICTED: dynein intermediate chain 3, ciliary [Drosophila
>gi|1036816171|ref|XP_017137376.1| PREDICTED: diphthamide biosynthesis protein 1 [Drosophila mi
>gi|1036816188|ref|XP_017137377.1| PREDICTED: diphthamide biosynthesis protein 1 [Drosophila mi
>gi|1036816206|ref|XP_017137378.1| PREDICTED: diphthamide biosynthesis protein 1 [Drosophila mi
>gi|1036801688|ref|XP_017136583.1| PREDICTED: uncharacterized protein LOC108152071 [Drosophila
>gi|1036801706|ref|XP_017136584.1| PREDICTED: uncharacterized protein LOC108152071 [Drosophila
>gi|1036801724|ref|XP_017136585.1| PREDICTED: biogenesis of lysosome-related organelles comple
>gi|1036801653|ref|XP_017136581.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801634|ref|XP_017136580.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801672|ref|XP_017136582.1| PREDICTED: general transcription factor IIE subunit 1 [Dros
>gi|1036839157|ref|XP_017138675.1| PREDICTED: F-box/LRR-repeat protein 16 isoform X1 [Drosophil
>gi|1036839176|ref|XP_017138676.1| PREDICTED: F-box/LRR-repeat protein 16 isoform X2 [Drosophil
>gi|1036839194|ref|XP_017138677.1| PREDICTED: uncharacterized protein LOC108153298 [Drosophila
>gi|1036816592|ref|XP_017137401.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC8 [Dr
>gi|1036816573|ref|XP_017137400.1| PREDICTED: serine/threonine-protein kinase RIO1 [Drosophila
>gi|1036816534|ref|XP_017137397.1| PREDICTED: ATP-binding cassette sub-family G member 1 isofor
>gi|1036816553|ref|XP_017137399.1| PREDICTED: ATP-binding cassette sub-family G member 1 isofor
>gi|1036833084|ref|XP_017138324.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833102|ref|XP_017138325.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833121|ref|XP_017138326.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833138|ref|XP_017138328.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833156|ref|XP_017138329.1| PREDICTED: acyl-protein thioesterase 1 isoform X1 [Drosophil
>gi|1036833229|ref|XP_017138333.1| PREDICTED: acyl-protein thioesterase 1 isoform X3 [Drosophil
>gi|1036833174|ref|XP_017138330.1| PREDICTED: acyl-protein thioesterase 1 isoform X2 [Drosophil
>gi|1036833193|ref|XP_017138331.1| PREDICTED: acyl-protein thioesterase 1 isoform X2 [Drosophil
>gi|1036833212|ref|XP_017138332.1| PREDICTED: acyl-protein thioesterase 1 isoform X2 [Drosophil
>gi|1036838610|ref|XP_017138643.1| PREDICTED: PCI domain-containing protein 2 homolog [Drosoph
>gi|1036848678|ref|XP_017139184.1| PREDICTED: uncharacterized protein LOC108153603 isoform X2
>gi|1036848659|ref|XP_017139183.1| PREDICTED: uncharacterized protein LOC108153603 isoform X1
>gi|1036848640|ref|XP_017139182.1| PREDICTED: alcohol dehydrogenase [NADP(+)] [Drosophila mira
>gi|1036842273|ref|XP_017138839.1| PREDICTED: aldose reductase isoform X1 [Drosophila miranda]
>gi|1036842328|ref|XP_017138840.1| PREDICTED: aldose reductase isoform X2 [Drosophila miranda]
>gi|1036849429|ref|XP_017139229.1| PREDICTED: putative protein TPRXL [Drosophila miranda]
>gi|1036849413|ref|XP_017139227.1| PREDICTED: uncharacterized serine-rich protein C215.13-like
>gi|1036830884|ref|XP_017138202.1| PREDICTED: uncharacterized protein LOC108152986 isoform X1
>gi|1036830922|ref|XP_017138204.1| PREDICTED: uncharacterized protein LOC108152986 isoform X3
>gi|1036830903|ref|XP_017138203.1| PREDICTED: uncharacterized protein LOC108152986 isoform X2
>gi|1036830943|ref|XP_017138205.1| PREDICTED: uncharacterized protein LOC108152986 isoform X4
>gi|1036823934|ref|XP_017137805.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036823953|ref|XP_017137806.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036823971|ref|XP_017137807.1| PREDICTED: zinc finger protein sens [Drosophila miranda]
>gi|1036824063|ref|XP_017137813.1| PREDICTED: uncharacterized protein LOC108152760 [Drosophila
>gi|1036824046|ref|XP_017137812.1| PREDICTED: protein new-glue 3-like [Drosophila miranda]
>gi|1036823897|ref|XP_017137803.1| PREDICTED: protein O-mannosyltransferase 1 [Drosophila mira
>gi|1036823916|ref|XP_017137804.1| PREDICTED: protein O-mannosyltransferase 1 [Drosophila mira
>gi|1036823990|ref|XP_017137809.1| PREDICTED: uncharacterized protein LOC108152757 [Drosophila
>gi|1036824029|ref|XP_017137811.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036824010|ref|XP_017137810.1| PREDICTED: mitochondrial import inner membrane translocase s
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>gi|1036799366|ref|XP_017136450.1| PREDICTED: MOB kinase activator-like 2 isoform X4 [Drosophil
>gi|1036799328|ref|XP_017136448.1| PREDICTED: MOB kinase activator-like 2 isoform X2 [Drosophil
>gi|1036799309|ref|XP_017136447.1| PREDICTED: MOB kinase activator-like 2 isoform X1 [Drosophil
>gi|1036799422|ref|XP_017136453.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1036799347|ref|XP_017136449.1| PREDICTED: MOB kinase activator-like 2 isoform X3 [Drosophil
>gi|1036799385|ref|XP_017136451.1| PREDICTED: putative alpha-L-fucosidase [Drosophila miranda]
>gi|1036799404|ref|XP_017136452.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1036759771|ref|XP_017156857.1| PREDICTED: uncharacterized protein LOC108165334 [Drosophila
>gi|1036799441|ref|XP_017136454.1| PREDICTED: salivary glue protein Sgs-3-like [Drosophila mira
>gi|1036806637|ref|XP_017136857.1| PREDICTED: protein charybde isoform X2 [Drosophila miranda]
>gi|1036806619|ref|XP_017136856.1| PREDICTED: protein charybde isoform X1 [Drosophila miranda]
>gi|1036847028|ref|XP_017139091.1| PREDICTED: uncharacterized protein LOC108153537 isoform X2
>gi|1036847009|ref|XP_017139090.1| PREDICTED: uncharacterized protein LOC108153537 isoform X1
>gi|1036759788|ref|XP_017156858.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036850578|ref|XP_017139297.1| PREDICTED: ketohexokinase [Drosophila miranda]
>gi|1036806526|ref|XP_017136850.1| PREDICTED: uncharacterized protein LOC108152202 [Drosophila
>gi|1036806582|ref|XP_017136853.1| PREDICTED: androgen-induced gene 1 protein [Drosophila mirar
>gi|1036806489|ref|XP_017136848.1| PREDICTED: methylenetetrahydrofolate reductase 1 [Drosophila
>gi|1036806601|ref|XP_017136855.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036806545|ref|XP_017136851.1| PREDICTED: eukaryotic translation initiation factor 4H [Dros
>gi|1036806471|ref|XP_017136847.1| PREDICTED: YLP motif-containing protein 1 [Drosophila mirano
>gi|1036806563|ref|XP_017136852.1| PREDICTED: uncharacterized protein LOC108152205 [Drosophila
>gi|1036806507|ref|XP_017136849.1| PREDICTED: glutaminyl-peptide cyclotransferase-like protein
>gi|1036759807|ref|XP_017156859.1| PREDICTED: CAAX prenyl protease 1 homolog [Drosophila mirano
>gi|1036840850|ref|XP_017138764.1| PREDICTED: AF4/FMR2 family member 4 [Drosophila miranda]
>gi|1036848257|ref|XP_017139160.1| PREDICTED: gamma-tubulin complex component 2 homolog isoforr
>gi|1036848238|ref|XP_017139159.1| PREDICTED: gamma-tubulin complex component 2 homolog isoforr
>gi|1036833947|ref|XP_017138364.1| PREDICTED: protein scylla [Drosophila miranda]
>gi|1036814350|ref|XP_017137274.1| PREDICTED: uncharacterized protein LOC108152442 isoform X1
>gi|1036814369|ref|XP_017137277.1| PREDICTED: uncharacterized protein LOC108152442 isoform X2
>gi|1036814213|ref|XP_017137267.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814232|ref|XP_017137268.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814253|ref|XP_017137269.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814272|ref|XP_017137270.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814291|ref|XP_017137271.1| PREDICTED: uncharacterized protein LOC108152440 isoform X1
>gi|1036814329|ref|XP_017137273.1| PREDICTED: uncharacterized protein LOC108152440 isoform X3
>gi|1036814310|ref|XP_017137272.1| PREDICTED: uncharacterized protein LOC108152440 isoform X2
>gi|1036810546|ref|XP_017137070.1| PREDICTED: protein FAM91A1 [Drosophila miranda]
>gi|1036810491|ref|XP_017137067.1| PREDICTED: ubiquitin-protein ligase E3A isoform X1 [Drosophi
>gi|1036810509|ref|XP_017137068.1| PREDICTED: ubiquitin-protein ligase E3A isoform X2 [Drosophi
>gi|1036810528|ref|XP_017137069.1| PREDICTED: ubiquitin-protein ligase E3A isoform X3 [Drosophi
>gi|1036825126|ref|XP_017137872.1| PREDICTED: procollagen-lysine,2-oxoglutarate 5-dioxygenase 
>gi|1036825144|ref|XP_017137873.1| PREDICTED: immunoglobulin domain-containing protein oig-1 [
>gi|1036825164|ref|XP_017137874.1| PREDICTED: immunoglobulin domain-containing protein oig-1 [
>gi|1036824867|ref|XP_017137858.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucurc
>gi|1036824904|ref|XP_017137859.1| PREDICTED: galactosylgalactosylxylosylprotein 3-beta-glucurc
>gi|1036767759|ref|XP_017134644.1| PREDICTED: protein FAM188B2 [Drosophila miranda]
>gi|1036793284|ref|XP_017136107.1| PREDICTED: uncharacterized aarF domain-containing protein ki
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>gi|1036793339|ref|XP_017136109.1| PREDICTED: uncharacterized protein LOC108151775 [Drosophila
>gi|1036793265|ref|XP_017136106.1| PREDICTED: protein wntless [Drosophila miranda]
>gi|1036793305|ref|XP_017136108.1| PREDICTED: putative Dol-P-Glc:Glc(2)Man(9)GlcNAc(2)-PP-Dol a
>gi|1036793246|ref|XP_017136105.1| PREDICTED: brain-specific angiogenesis inhibitor 1-associate
>gi|1036793210|ref|XP_017136104.1| PREDICTED: putative uncharacterized protein DDB_G0277255 iso
>gi|1036793358|ref|XP_017136110.1| PREDICTED: uncharacterized protein LOC108151776 [Drosophila
>gi|1036823224|ref|XP_017137766.1| PREDICTED: sodium-dependent phosphate transporter 1-B [Droso
>gi|1036823243|ref|XP_017137767.1| PREDICTED: sodium-dependent phosphate transporter 1-B [Droso
>gi|1036843637|ref|XP_017138916.1| PREDICTED: uncharacterized protein LOC108153424 [Drosophila
>gi|1036813914|ref|XP_017137250.1| PREDICTED: forkhead box protein K2 isoform X2 [Drosophila mi
>gi|1036813876|ref|XP_017137247.1| PREDICTED: forkhead box protein K2 isoform X1 [Drosophila mi
>gi|1036813894|ref|XP_017137248.1| PREDICTED: forkhead box protein K2 isoform X1 [Drosophila mi
>gi|1036813932|ref|XP_017137251.1| PREDICTED: 39S ribosomal protein L2, mitochondrial [Drosophi
>gi|1036813950|ref|XP_017137252.1| PREDICTED: ubiquitin fusion degradation protein 1 homolog [D
>gi|1036829033|ref|XP_017138095.1| PREDICTED: superoxide dismutase [Cu-Zn] isoform X2 [Drosophi
>gi|1036829015|ref|XP_017138094.1| PREDICTED: superoxide dismutase [Cu-Zn] isoform X1 [Drosophi
>gi|1036829003|ref|XP_017138092.1| PREDICTED: transmembrane protein 161B [Drosophila miranda]
>gi|1036851458|ref|XP_017139349.1| PREDICTED: protein new-glue 2-like [Drosophila miranda]
>gi|1036851472|ref|XP_017139350.1| PREDICTED: uncharacterized protein LOC108153732 [Drosophila
>gi|1036769705|ref|XP_017134755.1| PREDICTED: uncharacterized protein LOC108150936 [Drosophila
>gi|1036759825|ref|XP_017156860.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036819553|ref|XP_017137567.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036819513|ref|XP_017137565.1| PREDICTED: nucleobindin-2 [Drosophila miranda]
>gi|1036819534|ref|XP_017137566.1| PREDICTED: probable ATP-dependent RNA helicase DDX52 [Drosop
>gi|1036845261|ref|XP_017138998.1| PREDICTED: uncharacterized protein LOC108153475 [Drosophila
>gi|1036845244|ref|XP_017138997.1| PREDICTED: arylsulfatase B [Drosophila miranda]
>gi|1036849909|ref|XP_017139258.1| PREDICTED: battenin [Drosophila miranda]
>gi|1036849928|ref|XP_017139259.1| PREDICTED: battenin [Drosophila miranda]
>gi|1036849947|ref|XP_017139261.1| PREDICTED: battenin [Drosophila miranda]
>gi|1036765695|ref|XP_017134534.1| PREDICTED: glutamate receptor ionotropic, kainate 4 [Drosoph
>gi|1036765727|ref|XP_017134535.1| PREDICTED: glutamate receptor ionotropic, kainate 4 [Drosoph
>gi|1036759844|ref|XP_017156861.1| PREDICTED: glutamate receptor ionotropic, kainate 5 [Drosoph
>gi|1036759863|ref|XP_017156862.1| PREDICTED: gamma-aminobutyric acid receptor alpha-like [Dros
>gi|1036778515|ref|XP_017135258.1| PREDICTED: EF-hand domain-containing family member B-like [D
>gi|1036778554|ref|XP_017135260.1| PREDICTED: glycerol-3-phosphate phosphatase [Drosophila mira
>gi|1036778535|ref|XP_017135259.1| PREDICTED: glycerol-3-phosphate phosphatase [Drosophila mira
>gi|1036778665|ref|XP_017135267.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036778646|ref|XP_017135266.1| PREDICTED: dihydrolipoyl dehydrogenase, mitochondrial [Dros
>gi|1036778573|ref|XP_017135261.1| PREDICTED: tryptophan--tRNA ligase [Drosophila miranda]
>gi|1036778592|ref|XP_017135262.1| PREDICTED: cationic amino acid transporter 2 [Drosophila mi
>gi|1036778610|ref|XP_017135263.1| PREDICTED: arylsulfatase B isoform X1 [Drosophila miranda]
>gi|1036778629|ref|XP_017135265.1| PREDICTED: arylsulfatase B isoform X2 [Drosophila miranda]
>gi|1036778684|ref|XP_017135268.1| PREDICTED: uncharacterized protein LOC108151261 [Drosophila
>gi|1036766561|ref|XP_017134581.1| PREDICTED: uncharacterized protein LOC108150790 [Drosophila
>gi|1036759882|ref|XP_017156863.1| PREDICTED: arylsulfatase B [Drosophila miranda]
>gi|1036844991|ref|XP_017138982.1| PREDICTED: CD151 antigen [Drosophila miranda]
>gi|1036845010|ref|XP_017138983.1| PREDICTED: CD151 antigen [Drosophila miranda]
>gi|1036841752|ref|XP_017138819.1| PREDICTED: LOW QUALITY PROTEIN: solute carrier family 26 mem
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>gi|1036821183|ref|XP_017137650.1| PREDICTED: WD repeat-containing protein 92 [Drosophila mira
>gi|1036821199|ref|XP_017137651.1| PREDICTED: WD repeat-containing protein 92 [Drosophila mira
>gi|1036821163|ref|XP_017137649.1| PREDICTED: LOW QUALITY PROTEIN: tetratricopeptide repeat pro
>gi|1036849999|ref|XP_017139264.1| PREDICTED: uncharacterized protein LOC108153663 [Drosophila
>gi|1036850019|ref|XP_017139266.1| PREDICTED: uncharacterized protein LOC108153663 [Drosophila
>gi|1036759901|ref|XP_017156864.1| PREDICTED: dihydrolipoyl dehydrogenase-like [Drosophila mira
>gi|1036801175|ref|XP_017136553.1| PREDICTED: ecdysone-induced protein 75B isoform X1 [Drosophi
>gi|1036801225|ref|XP_017136556.1| PREDICTED: ecdysone-induced protein 75B, isoforms C/D isofor
>gi|1036801208|ref|XP_017136555.1| PREDICTED: ecdysone-induced protein 75B, isoforms C/D isofor
>gi|1036801192|ref|XP_017136554.1| PREDICTED: ecdysone-induced protein 75B isoform X2 [Drosophi
>gi|1036801242|ref|XP_017136557.1| PREDICTED: ecdysone-induced protein 75B, isoforms C/D isofor
>gi|1036759920|ref|XP_017156865.1| PREDICTED: uncharacterized protein LOC108165343 [Drosophila
>gi|1036849767|ref|XP_017139249.1| PREDICTED: uncharacterized protein LOC108153654 [Drosophila
>gi|1036849785|ref|XP_017139250.1| PREDICTED: uncharacterized protein LOC108153655 isoform X1
>gi|1036849803|ref|XP_017139252.1| PREDICTED: uncharacterized protein LOC108153655 isoform X2
>gi|1036765659|ref|XP_017157176.1| PREDICTED: uncharacterized protein LOC108165611 [Drosophila
>gi|1036829193|ref|XP_017138104.1| PREDICTED: uncharacterized protein LOC108152929 [Drosophila
>gi|1036829212|ref|XP_017138105.1| PREDICTED: uncharacterized protein LOC108152929 [Drosophila
>gi|1036829231|ref|XP_017138106.1| PREDICTED: probable 28S ribosomal protein S26, mitochondrial
>gi|1036829172|ref|XP_017138103.1| PREDICTED: DNA-directed RNA polymerase III subunit RPC4 [Dro
>gi|1036787005|ref|XP_017135751.1| PREDICTED: ATR-interacting protein mus304 [Drosophila mirand
>gi|1036787043|ref|XP_017135753.1| PREDICTED: uncharacterized protein LOC108151585 [Drosophila
>gi|1036787024|ref|XP_017135752.1| PREDICTED: probable cytochrome P450 312a1 [Drosophila mirand
>gi|1036787061|ref|XP_017135754.1| PREDICTED: uncharacterized protein LOC108151586 isoform X1
>gi|1036787080|ref|XP_017135755.1| PREDICTED: uncharacterized protein LOC108151586 isoform X2
>gi|1036787116|ref|XP_017135757.1| PREDICTED: uncharacterized protein LOC108151589 [Drosophila
>gi|1036787135|ref|XP_017135758.1| PREDICTED: uncharacterized protein LOC108151590 [Drosophila
>gi|1036787153|ref|XP_017135759.1| PREDICTED: uncharacterized protein LOC108151591 [Drosophila
>gi|1036787099|ref|XP_017135756.1| PREDICTED: uncharacterized protein LOC108151587 [Drosophila
>gi|1036831774|ref|XP_017138251.1| PREDICTED: cell death protein hid [Drosophila miranda]
>gi|1036850056|ref|XP_017139268.1| PREDICTED: larval serum protein 1 gamma chain [Drosophila mi
>gi|1036767164|ref|XP_017134616.1| PREDICTED: uncharacterized protein LOC108150820 [Drosophila
>gi|1036850670|ref|XP_017139303.1| PREDICTED: uncharacterized protein LOC108153690 [Drosophila
>gi|1036849494|ref|XP_017139232.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036763349|ref|XP_017157060.1| PREDICTED: cell death protein Grim [Drosophila miranda]
>gi|1036766326|ref|XP_017134566.1| PREDICTED: uncharacterized protein LOC108150780 isoform X1
>gi|1036766347|ref|XP_017134568.1| PREDICTED: uncharacterized protein LOC108150780 isoform X2
>gi|1036849580|ref|XP_017139238.1| PREDICTED: uncharacterized protein LOC108153645 [Drosophila
>gi|1036764169|ref|XP_017157104.1| PREDICTED: coactosin-like protein [Drosophila miranda]
>gi|1036765778|ref|XP_017134539.1| PREDICTED: mitochondrial dicarboxylate carrier [Drosophila m
>gi|1036763136|ref|XP_017157051.1| PREDICTED: cell death protein rpr [Drosophila miranda]
>gi|1036832217|ref|XP_017138277.1| PREDICTED: uncharacterized protein LOC108153032 [Drosophila
>gi|1036832198|ref|XP_017138276.1| PREDICTED: gamma-glutamylcyclotransferase [Drosophila miranc
>gi|1036832181|ref|XP_017138275.1| PREDICTED: gamma-glutamylcyclotransferase [Drosophila miranc
>gi|1036835683|ref|XP_017138464.1| PREDICTED: angiopoietin-related protein 1-like [Drosophila r
>gi|1036835630|ref|XP_017138461.1| PREDICTED: somatostatin receptor type 2 [Drosophila miranda]
>gi|1036835701|ref|XP_017138465.1| PREDICTED: uncharacterized protein LOC108153171 [Drosophila
>gi|1036835647|ref|XP_017138462.1| PREDICTED: somatostatin receptor type 5 [Drosophila miranda]
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>gi|1036835666|ref|XP_017138463.1| PREDICTED: somatostatin receptor type 5 [Drosophila miranda]
>gi|1036838106|ref|XP_017138610.1| PREDICTED: protein terminus [Drosophila miranda]
>gi|1036825411|ref|XP_017137889.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2-like [Droso
>gi|1036825392|ref|XP_017137888.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Droso
>gi|1036825375|ref|XP_017137886.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1036825356|ref|XP_017137885.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 [Drosophila
>gi|1036825430|ref|XP_017137890.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1-like [Droso
>gi|1036825337|ref|XP_017137884.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-2 [Drosophila
>gi|1036825299|ref|XP_017137882.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 isoform X1
>gi|1036825318|ref|XP_017137883.1| PREDICTED: prolyl 4-hydroxylase subunit alpha-1 isoform X2
>gi|1036842403|ref|XP_017138843.1| PREDICTED: uncharacterized protein LOC108153381 [Drosophila
>gi|1036808882|ref|XP_017136980.1| PREDICTED: LOW QUALITY PROTEIN: ubiquitin carboxyl-terminal
>gi|1036808901|ref|XP_017136982.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase nonstop-li
>gi|1036808863|ref|XP_017136979.1| PREDICTED: LOW QUALITY PROTEIN: protein aurora borealis [Dro
>gi|1036819061|ref|XP_017137541.1| PREDICTED: uncharacterized protein LOC108152597 [Drosophila
>gi|1036819080|ref|XP_017137543.1| PREDICTED: protein phosphatase 1 regulatory subunit 12A [Dro
>gi|1036763295|ref|XP_017157057.1| PREDICTED: uncharacterized protein LOC108165512 [Drosophila
>gi|1036788386|ref|XP_017135827.1| PREDICTED: caprin homolog [Drosophila miranda]
>gi|1036788454|ref|XP_017135832.1| PREDICTED: gram-negative bacteria-binding protein 2 [Drosop
>gi|1036788404|ref|XP_017135828.1| PREDICTED: gram-negative bacteria-binding protein 1 isoform
>gi|1036788420|ref|XP_017135829.1| PREDICTED: gram-negative bacteria-binding protein 2 isoform
>gi|1036788437|ref|XP_017135830.1| PREDICTED: gram-negative bacteria-binding protein 2-like [D
>gi|1036788471|ref|XP_017135833.1| PREDICTED: SAGA-associated factor 11 homolog [Drosophila mi
>gi|1036788503|ref|XP_017135835.1| PREDICTED: uncharacterized protein LOC108151640 [Drosophila
>gi|1036788487|ref|XP_017135834.1| PREDICTED: charged multivesicular body protein 1b [Drosophi
>gi|1036847180|ref|XP_017139100.1| PREDICTED: probable cytochrome P450 12c1, mitochondrial [Dro
>gi|1036759939|ref|XP_017156866.1| PREDICTED: mitochondrial dicarboxylate carrier, partial [Dro
>gi|1036787986|ref|XP_017135807.1| PREDICTED: uncharacterized protein LOC108151619 [Drosophila
>gi|1036787821|ref|XP_017135799.1| PREDICTED: uncharacterized protein LOC108151613 [Drosophila
>gi|1036787859|ref|XP_017135801.1| PREDICTED: uncharacterized protein LOC108151615 [Drosophila
>gi|1036787840|ref|XP_017135800.1| PREDICTED: equilibrative nucleoside transporter 4 [Drosophil
>gi|1036787915|ref|XP_017135804.1| PREDICTED: WW domain-binding protein 2 isoform X1 [Drosophil
>gi|1036787968|ref|XP_017135806.1| PREDICTED: WW domain-binding protein 2 isoform X3 [Drosophil
>gi|1036787932|ref|XP_017135805.1| PREDICTED: WW domain-binding protein 2 isoform X2 [Drosophil
>gi|1036787710|ref|XP_017135792.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X1
>gi|1036787727|ref|XP_017135793.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 1 isoform X2
>gi|1036787783|ref|XP_017135796.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X5
>gi|1036787802|ref|XP_017135797.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X6
>gi|1036787764|ref|XP_017135795.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 1 isoform X4
>gi|1036787746|ref|XP_017135794.1| PREDICTED: ecto-NOX disulfide-thiol exchanger 2 isoform X3
>gi|1036787877|ref|XP_017135802.1| PREDICTED: facilitated trehalose transporter Tret1-2 homolog
>gi|1036787896|ref|XP_017135803.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1036810362|ref|XP_017137059.1| PREDICTED: titin [Drosophila miranda]
>gi|1036818506|ref|XP_017137510.1| PREDICTED: splicing factor 3A subunit 2 [Drosophila miranda]
>gi|1036818487|ref|XP_017137509.1| PREDICTED: mucin-19 isoform X4 [Drosophila miranda]
>gi|1036818469|ref|XP_017137507.1| PREDICTED: mucin-19 isoform X3 [Drosophila miranda]
>gi|1036818450|ref|XP_017137506.1| PREDICTED: mucin-19 isoform X2 [Drosophila miranda]
>gi|1036818432|ref|XP_017137505.1| PREDICTED: mucin-19 isoform X1 [Drosophila miranda]
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>gi|1036809408|ref|XP_017137006.1| PREDICTED: serine/threonine-protein kinase unc-51 [Drosophil
>gi|1036809427|ref|XP_017137007.1| PREDICTED: leucine-rich repeat and transmembrane domain-cont
>gi|1036850037|ref|XP_017139267.1| PREDICTED: uncharacterized protein LOC108153664 [Drosophila
>gi|1036759958|ref|XP_017156867.1| PREDICTED: RNA polymerase II degradation factor 1 [Drosophil
>gi|1036816362|ref|XP_017137387.1| PREDICTED: uncharacterized protein LOC108152508 [Drosophila
>gi|1036816381|ref|XP_017137388.1| PREDICTED: AP2-associated protein kinase 1 [Drosophila miran
>gi|1036759975|ref|XP_017156868.1| PREDICTED: uncharacterized protein LOC108165346 [Drosophila
>gi|1036816479|ref|XP_017137394.1| PREDICTED: uncharacterized protein C16orf52 homolog A [Droso
>gi|1036816322|ref|XP_017137385.1| PREDICTED: adenylate cyclase type 2 isoform X2 [Drosophila n
>gi|1036816303|ref|XP_017137384.1| PREDICTED: adenylate cyclase type 2 isoform X1 [Drosophila n
>gi|1036816341|ref|XP_017137386.1| PREDICTED: adenylate cyclase type 2 isoform X3 [Drosophila n
>gi|1036816498|ref|XP_017137395.1| PREDICTED: uncharacterized protein LOC108152513 [Drosophila
>gi|1036816400|ref|XP_017137390.1| PREDICTED: uncharacterized protein LOC108152510 isoform X1
>gi|1036816421|ref|XP_017137391.1| PREDICTED: uncharacterized protein LOC108152510 isoform X2
>gi|1036816440|ref|XP_017137392.1| PREDICTED: uncharacterized protein LOC108152510 isoform X3
>gi|1036816458|ref|XP_017137393.1| PREDICTED: uncharacterized protein LOC108152511 [Drosophila
>gi|1036816516|ref|XP_017137396.1| PREDICTED: uncharacterized protein LOC108152514 [Drosophila
>gi|1036804217|ref|XP_017136723.1| PREDICTED: rho GDP-dissociation inhibitor 1 [Drosophila mira
>gi|1036804236|ref|XP_017136724.1| PREDICTED: rho GDP-dissociation inhibitor 1 [Drosophila mira
>gi|1036804198|ref|XP_017136722.1| PREDICTED: metaxin-2 [Drosophila miranda]
>gi|1036804254|ref|XP_017136725.1| PREDICTED: larval/pupal cuticle protein H1C-like [Drosophila
>gi|1036804179|ref|XP_017136720.1| PREDICTED: Golgi reassembly-stacking protein 2 [Drosophila n
>gi|1036804273|ref|XP_017136726.1| PREDICTED: probable prefoldin subunit 6 [Drosophila miranda]
>gi|1036819395|ref|XP_017137558.1| PREDICTED: polycomb protein Su(z)12 isoform X1 [Drosophila n
>gi|1036819414|ref|XP_017137559.1| PREDICTED: polycomb protein Su(z)12 isoform X2 [Drosophila n
>gi|1036820778|ref|XP_017137628.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 2 is
>gi|1036820759|ref|XP_017137627.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 2 is
>gi|1036829903|ref|XP_017138142.1| PREDICTED: exosome complex component MTR3 [Drosophila miranc
>gi|1036829882|ref|XP_017138141.1| PREDICTED: U4/U6 small nuclear ribonucleoprotein Prp3 [Droso
>gi|1036770726|ref|XP_017134810.1| PREDICTED: chromodomain-helicase-DNA-binding protein Mi-2 ho
>gi|1036770747|ref|XP_017134811.1| PREDICTED: chromodomain-helicase-DNA-binding protein Mi-2 ho
>gi|1036770822|ref|XP_017134816.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770840|ref|XP_017134817.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770865|ref|XP_017134818.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770882|ref|XP_017134819.1| PREDICTED: RNA polymerase II elongation factor Ell [Drosophi
>gi|1036770785|ref|XP_017134813.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 32 isoform
>gi|1036770766|ref|XP_017134812.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 32 isoform
>gi|1036770804|ref|XP_017134814.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 32 isoform
>gi|1036792369|ref|XP_017136056.1| PREDICTED: uncharacterized protein LOC108151745 [Drosophila
>gi|1036792219|ref|XP_017136047.1| PREDICTED: protein suppressor of underreplication [Drosophil
>gi|1036792328|ref|XP_017136054.1| PREDICTED: uncharacterized protein LOC108151743 [Drosophila
>gi|1036792348|ref|XP_017136055.1| PREDICTED: uncharacterized protein LOC108151744 [Drosophila
>gi|1036792235|ref|XP_017136049.1| PREDICTED: succinyl-CoA:3-ketoacid coenzyme A transferase 1
>gi|1036792290|ref|XP_017136052.1| PREDICTED: uncharacterized protein LOC108151742 isoform X3
>gi|1036792273|ref|XP_017136051.1| PREDICTED: uncharacterized protein LOC108151742 isoform X2
>gi|1036792254|ref|XP_017136050.1| PREDICTED: uncharacterized protein LOC108151742 isoform X1
>gi|1036792309|ref|XP_017136053.1| PREDICTED: uncharacterized protein LOC108151742 isoform X4
>gi|1036852052|ref|XP_017139387.1| PREDICTED: proton-coupled amino acid transporter 4 [Drosophi
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>gi|1036759995|ref|XP_017156869.1| PREDICTED: uncharacterized protein LOC108165347 [Drosophila
>gi|1036766366|ref|XP_017134569.1| PREDICTED: uncharacterized protein LOC108150781 [Drosophila
>gi|1036760013|ref|XP_017156870.1| PREDICTED: uncharacterized protein LOC108165348 [Drosophila
>gi|1036772467|ref|XP_017134910.1| PREDICTED: LOW QUALITY PROTEIN: titin-like [Drosophila mira
>gi|1036772444|ref|XP_017134909.1| PREDICTED: uncharacterized protein DDB_G0285291-like [Droso
>gi|1036842460|ref|XP_017138846.1| PREDICTED: SEC14 domain and spectrin repeat-containing prote
>gi|1036767013|ref|XP_017134607.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036847199|ref|XP_017139101.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036849598|ref|XP_017139239.1| PREDICTED: uncharacterized protein LOC108153646 [Drosophila
>gi|1036848107|ref|XP_017139151.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036763227|ref|XP_017157055.1| PREDICTED: uncharacterized protein LOC108165509 [Drosophila
>gi|1036814604|ref|XP_017137290.1| PREDICTED: uncharacterized protein LOC108152447 [Drosophila
>gi|1036814407|ref|XP_017137279.1| PREDICTED: intraflagellar transport protein 172 homolog iso
>gi|1036814426|ref|XP_017137280.1| PREDICTED: intraflagellar transport protein 172 homolog iso
>gi|1036814622|ref|XP_017137291.1| PREDICTED: regulation of nuclear pre-mRNA domain-containing
>gi|1036814446|ref|XP_017137281.1| PREDICTED: adenylate cyclase type 8 isoform X1 [Drosophila r
>gi|1036814465|ref|XP_017137282.1| PREDICTED: adenylate cyclase type 8 isoform X1 [Drosophila r
>gi|1036814485|ref|XP_017137283.1| PREDICTED: adenylate cyclase type 8 isoform X2 [Drosophila r
>gi|1036814504|ref|XP_017137284.1| PREDICTED: adenylate cyclase type 8 isoform X3 [Drosophila r
>gi|1036814546|ref|XP_017137286.1| PREDICTED: adenylate cyclase type 8 isoform X5 [Drosophila r
>gi|1036814583|ref|XP_017137288.1| PREDICTED: adenylate cyclase type 8 isoform X7 [Drosophila r
>gi|1036814564|ref|XP_017137287.1| PREDICTED: adenylate cyclase type 8 isoform X6 [Drosophila r
>gi|1036814527|ref|XP_017137285.1| PREDICTED: adenylate cyclase type 2 isoform X4 [Drosophila r
>gi|1036786354|ref|XP_017135716.1| PREDICTED: adenylate cyclase type 2-like isoform X1 [Drosopl
>gi|1036786373|ref|XP_017135717.1| PREDICTED: adenylate cyclase type 8-like isoform X2 [Drosopl
>gi|1036786507|ref|XP_017135724.1| PREDICTED: cytochrome b reductase 1 isoform X3 [Drosophila r
>gi|1036786430|ref|XP_017135720.1| PREDICTED: cytochrome b reductase 1 isoform X1 [Drosophila r
>gi|1036786449|ref|XP_017135721.1| PREDICTED: cytochrome b reductase 1 isoform X2 [Drosophila r
>gi|1036786470|ref|XP_017135722.1| PREDICTED: cytochrome b reductase 1 isoform X2 [Drosophila r
>gi|1036786489|ref|XP_017135723.1| PREDICTED: cytochrome b reductase 1 isoform X2 [Drosophila r
>gi|1036786526|ref|XP_017135725.1| PREDICTED: cytochrome b reductase 1 isoform X4 [Drosophila r
>gi|1036786545|ref|XP_017135727.1| PREDICTED: uncharacterized protein LOC108151563 [Drosophila
>gi|1036786392|ref|XP_017135718.1| PREDICTED: xylosyltransferase oxt [Drosophila miranda]
>gi|1036786564|ref|XP_017135728.1| PREDICTED: 5'-3' exoribonuclease 2 homolog [Drosophila mirar
>gi|1036786411|ref|XP_017135719.1| PREDICTED: protein ecdysoneless [Drosophila miranda]
>gi|1036835407|ref|XP_017138448.1| PREDICTED: uncharacterized protein LOC108153156 [Drosophila
>gi|1036835388|ref|XP_017138447.1| PREDICTED: peritrophin-48-like [Drosophila miranda]
>gi|1036835332|ref|XP_017138443.1| PREDICTED: major royal jelly protein 1 [Drosophila miranda]
>gi|1036835350|ref|XP_017138444.1| PREDICTED: major royal jelly protein 1 [Drosophila miranda]
>gi|1036835313|ref|XP_017138442.1| PREDICTED: sex peptide receptor [Drosophila miranda]
>gi|1036835294|ref|XP_017138441.1| PREDICTED: uncharacterized protein LOC108153150 [Drosophila
>gi|1036835423|ref|XP_017138449.1| PREDICTED: rab-like protein 3 [Drosophila miranda]
>gi|1036835369|ref|XP_017138446.1| PREDICTED: choline-phosphate cytidylyltransferase A-like [D
>gi|1036763770|ref|XP_017157085.1| PREDICTED: E3 ubiquitin-protein ligase MARCH6 [Drosophila m
>gi|1036763985|ref|XP_017157098.1| PREDICTED: thioredoxin, mitochondrial [Drosophila miranda]
>gi|1036763967|ref|XP_017157097.1| PREDICTED: 39S ribosomal protein L23, mitochondrial [Drosopl
>gi|1036763788|ref|XP_017157086.1| PREDICTED: nucleolar MIF4G domain-containing protein 1 homol
>gi|1036763842|ref|XP_017157089.1| PREDICTED: uncharacterized protein C7orf50 homolog [Drosophi
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>gi|1036763494|ref|XP_017157069.1| PREDICTED: uncharacterized protein LOC108165524 isoform X1
>gi|1036763512|ref|XP_017157070.1| PREDICTED: uncharacterized protein LOC108165524 isoform X2
>gi|1036763552|ref|XP_017157072.1| PREDICTED: uncharacterized protein LOC108165524 isoform X4
>gi|1036763533|ref|XP_017157071.1| PREDICTED: uncharacterized protein LOC108165524 isoform X3
>gi|1036763682|ref|XP_017157079.1| PREDICTED: uncharacterized protein LOC108165524 isoform X10
>gi|1036763860|ref|XP_017157090.1| PREDICTED: surface protein [Drosophila miranda]
>gi|1036763931|ref|XP_017157094.1| PREDICTED: uncharacterized protein LOC108165533 isoform X2
>gi|1036763949|ref|XP_017157095.1| PREDICTED: uncharacterized protein LOC108165533 isoform X3
>gi|1036763895|ref|XP_017157092.1| PREDICTED: uncharacterized protein LOC108165533 isoform X1
>gi|1036763913|ref|XP_017157093.1| PREDICTED: uncharacterized protein LOC108165533 isoform X1
>gi|1036763611|ref|XP_017157075.1| PREDICTED: uncharacterized protein LOC108165524 isoform X6
>gi|1036763586|ref|XP_017157073.1| PREDICTED: uncharacterized protein LOC108165524 isoform X5
>gi|1036763700|ref|XP_017157080.1| PREDICTED: microtubule-associated protein futsch isoform X1
>gi|1036763736|ref|XP_017157082.1| PREDICTED: serine/arginine repetitive matrix protein 2 isofo
>gi|1036763718|ref|XP_017157081.1| PREDICTED: mucin-5AC isoform X12 [Drosophila miranda]
>gi|1036763630|ref|XP_017157076.1| PREDICTED: supervillin isoform X7 [Drosophila miranda]
>gi|1036763754|ref|XP_017157083.1| PREDICTED: AF4/FMR2 family member 4 isoform X14 [Drosophila
>gi|1036763877|ref|XP_017157091.1| PREDICTED: uncharacterized protein LOC108165531 [Drosophila
>gi|1036763806|ref|XP_017157087.1| PREDICTED: probable cytochrome P450 4d20 [Drosophila miranda
>gi|1036763646|ref|XP_017157077.1| PREDICTED: supervillin isoform X8 [Drosophila miranda]
>gi|1036763824|ref|XP_017157088.1| PREDICTED: uncharacterized protein LOC108165528 [Drosophila
>gi|1036763664|ref|XP_017157078.1| PREDICTED: supervillin isoform X9 [Drosophila miranda]
>gi|1036832665|ref|XP_017138300.1| PREDICTED: purine nucleoside phosphorylase isoform X3 [Dros
>gi|1036832646|ref|XP_017138299.1| PREDICTED: purine nucleoside phosphorylase isoform X2 [Dros
>gi|1036832627|ref|XP_017138298.1| PREDICTED: purine nucleoside phosphorylase isoform X1 [Dros
>gi|1036832701|ref|XP_017138302.1| PREDICTED: purine nucleoside phosphorylase isoform X5 [Dros
>gi|1036832684|ref|XP_017138301.1| PREDICTED: purine nucleoside phosphorylase isoform X4 [Dros
>gi|1036789463|ref|XP_017135890.1| PREDICTED: AF4/FMR2 family member 4 isoform X18 [Drosophila
>gi|1036789294|ref|XP_017135880.1| PREDICTED: uncharacterized protein LOC108151662 isoform X9
>gi|1036789201|ref|XP_017135875.1| PREDICTED: uncharacterized protein LOC108151662 isoform X4
>gi|1036789387|ref|XP_017135885.1| PREDICTED: AF4/FMR2 family member 4 isoform X14 [Drosophila
>gi|1036789349|ref|XP_017135883.1| PREDICTED: AF4/FMR2 family member 4 isoform X12 [Drosophila
>gi|1036789259|ref|XP_017135878.1| PREDICTED: uncharacterized protein LOC108151662 isoform X7
>gi|1036789165|ref|XP_017135873.1| PREDICTED: uncharacterized protein LOC108151662 isoform X2
>gi|1036789311|ref|XP_017135881.1| PREDICTED: uncharacterized protein LOC108151662 isoform X10
>gi|1036789276|ref|XP_017135879.1| PREDICTED: uncharacterized protein LOC108151662 isoform X8
>gi|1036789368|ref|XP_017135884.1| PREDICTED: AF4/FMR2 family member 4 isoform X13 [Drosophila
>gi|1036789330|ref|XP_017135882.1| PREDICTED: AF4/FMR2 family member 4 isoform X11 [Drosophila
>gi|1036789183|ref|XP_017135874.1| PREDICTED: uncharacterized protein LOC108151662 isoform X3
>gi|1036789220|ref|XP_017135876.1| PREDICTED: uncharacterized protein LOC108151662 isoform X5
>gi|1036789144|ref|XP_017135872.1| PREDICTED: uncharacterized protein LOC108151662 isoform X1
>gi|1036789238|ref|XP_017135877.1| PREDICTED: uncharacterized protein LOC108151662 isoform X6
>gi|1036789406|ref|XP_017135886.1| PREDICTED: uncharacterized protein LOC108151662 isoform X15
>gi|1036789425|ref|XP_017135887.1| PREDICTED: uncharacterized protein LOC108151662 isoform X16
>gi|1036789482|ref|XP_017135891.1| PREDICTED: uncharacterized protein LOC108151662 isoform X19
>gi|1036789444|ref|XP_017135888.1| PREDICTED: AF4/FMR2 family member 4 isoform X17 [Drosophila
>gi|1036809813|ref|XP_017137030.1| PREDICTED: altered inheritance of mitochondria protein 3 iso
>gi|1036809788|ref|XP_017137029.1| PREDICTED: extensin isoform X1 [Drosophila miranda]
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>gi|1036809943|ref|XP_017137036.1| PREDICTED: O-acetyl-ADP-ribose deacetylase 1 [Drosophila mi
>gi|1036809922|ref|XP_017137035.1| PREDICTED: O-acetyl-ADP-ribose deacetylase 1 [Drosophila mi
>gi|1036809903|ref|XP_017137033.1| PREDICTED: uncharacterized protein LOC108152312 [Drosophila
>gi|1036809884|ref|XP_017137032.1| PREDICTED: uncharacterized oxidoreductase YjmC [Drosophila r
>gi|1036809866|ref|XP_017137031.1| PREDICTED: uncharacterized protein LOC108152310 [Drosophila
>gi|1036809695|ref|XP_017137024.1| PREDICTED: uncharacterized protein LOC108152307 isoform X1
>gi|1036809751|ref|XP_017137027.1| PREDICTED: uncharacterized protein LOC108152307 isoform X4
>gi|1036809714|ref|XP_017137025.1| PREDICTED: uncharacterized protein LOC108152307 isoform X2
>gi|1036809769|ref|XP_017137028.1| PREDICTED: uncharacterized protein LOC108152307 isoform X5
>gi|1036809733|ref|XP_017137026.1| PREDICTED: GTPase-activating protein BEM2/IPL2 isoform X3 [
>gi|1036809959|ref|XP_017137037.1| PREDICTED: multifunctional methyltransferase subunit TRM112-
>gi|1036836316|ref|XP_017138502.1| PREDICTED: integrin-linked protein kinase homolog pat-4 [Dro
>gi|1036841245|ref|XP_017138788.1| PREDICTED: ecdysone-induced protein 78C isoform X1 [Drosophi
>gi|1036841264|ref|XP_017138789.1| PREDICTED: ecdysone-induced protein 78C isoform X2 [Drosophi
>gi|1036841300|ref|XP_017138792.1| PREDICTED: peritrophin-48 [Drosophila miranda]
>gi|1036841282|ref|XP_017138790.1| PREDICTED: ecdysone-induced protein 78C isoform X3 [Drosophi
>gi|1036844209|ref|XP_017138947.1| PREDICTED: inositol monophosphatase 1 [Drosophila miranda]
>gi|1036849525|ref|XP_017139234.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1036789773|ref|XP_017135909.1| PREDICTED: acetyl-coenzyme A synthetase [Drosophila miranda]
>gi|1036789813|ref|XP_017135912.1| PREDICTED: glycine cleavage system H protein, mitochondrial
>gi|1036789794|ref|XP_017135910.1| PREDICTED: uncharacterized protein LOC108151679 [Drosophila
>gi|1036789754|ref|XP_017135908.1| PREDICTED: uncharacterized protein LOC108151677 [Drosophila
>gi|1036767612|ref|XP_017134639.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036767577|ref|XP_017134637.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036808488|ref|XP_017136959.1| PREDICTED: pupal cuticle protein Edg-78E-like [Drosophila mi
>gi|1036808450|ref|XP_017136957.1| PREDICTED: pupal cuticle protein Edg-78E [Drosophila miranda
>gi|1036808469|ref|XP_017136958.1| PREDICTED: pupal cuticle protein Edg-78E-like [Drosophila mi
>gi|1036808414|ref|XP_017136955.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036808432|ref|XP_017136956.1| PREDICTED: probable serine hydrolase [Drosophila miranda]
>gi|1036808396|ref|XP_017136954.1| PREDICTED: polycomb group protein Pc [Drosophila miranda]
>gi|1036847834|ref|XP_017139136.1| PREDICTED: ras-related protein Rab-26 isoform X2 [Drosophila
>gi|1036847853|ref|XP_017139137.1| PREDICTED: ras-related protein Rab-26 isoform X3 [Drosophila
>gi|1036847817|ref|XP_017139135.1| PREDICTED: ras-related protein Rab-26 isoform X1 [Drosophila
>gi|1036804826|ref|XP_017136757.1| PREDICTED: pre-rRNA-processing protein TSR1 homolog [Drosoph
>gi|1036804807|ref|XP_017136756.1| PREDICTED: TBC1 domain family member 9 isoform X2 [Drosophi
>gi|1036804789|ref|XP_017136755.1| PREDICTED: TBC1 domain family member 9 isoform X1 [Drosophi
>gi|1036838677|ref|XP_017138647.1| PREDICTED: uncharacterized protein LOC108153286 [Drosophila
>gi|1036838696|ref|XP_017138648.1| PREDICTED: uncharacterized protein LOC108153287 [Drosophila
>gi|1036838712|ref|XP_017138649.1| PREDICTED: uncharacterized protein LOC108153287 [Drosophila
>gi|1036846881|ref|XP_017139083.1| PREDICTED: nuclear transcription factor Y subunit beta [Dros
>gi|1036846900|ref|XP_017139084.1| PREDICTED: uncharacterized protein DDB_G0283697 isoform X1
>gi|1036846919|ref|XP_017139085.1| PREDICTED: uncharacterized protein LOC108153533 isoform X2
>gi|1036760031|ref|XP_017156871.1| PREDICTED: uncharacterized protein LOC108165349, partial [D
>gi|1036815665|ref|XP_017137348.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1036815684|ref|XP_017137349.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1036815703|ref|XP_017137350.1| PREDICTED: trichohyalin isoform X2 [Drosophila miranda]
>gi|1036815724|ref|XP_017137351.1| PREDICTED: HIV Tat-specific factor 1 [Drosophila miranda]
>gi|1036815743|ref|XP_017137352.1| PREDICTED: uncharacterized protein LOC108152488 [Drosophila
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>gi|1036780912|ref|XP_017135400.1| PREDICTED: adult enhancer factor 1 isoform X1 [Drosophila mi
>gi|1036780930|ref|XP_017135402.1| PREDICTED: adult enhancer factor 1 isoform X2 [Drosophila mi
>gi|1036780879|ref|XP_017135398.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036781019|ref|XP_017135408.1| PREDICTED: pro-resilin [Drosophila miranda]
>gi|1036760050|ref|XP_017156872.1| PREDICTED: cuticle protein 8 [Drosophila miranda]
>gi|1036780983|ref|XP_017135406.1| PREDICTED: uncharacterized protein LOC108151358 [Drosophila
>gi|1036781037|ref|XP_017135409.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036781001|ref|XP_017135407.1| PREDICTED: peroxisome biogenesis factor 2 [Drosophila mirand
>gi|1036780964|ref|XP_017135404.1| PREDICTED: UPF0183 protein CG7083 [Drosophila miranda]
>gi|1036780948|ref|XP_017135403.1| PREDICTED: DNA primase small subunit [Drosophila miranda]
>gi|1036780896|ref|XP_017135399.1| PREDICTED: heat shock 70 kDa protein 14 [Drosophila miranda]
>gi|1036799591|ref|XP_017136462.1| PREDICTED: rab GTPase-activating protein 1-like isoform X1
>gi|1036799612|ref|XP_017136463.1| PREDICTED: rab GTPase-activating protein 1-like isoform X2
>gi|1036799630|ref|XP_017136464.1| PREDICTED: uncharacterized protein LOC108152004 isoform X1
>gi|1036799649|ref|XP_017136465.1| PREDICTED: uncharacterized protein LOC108152004 isoform X2
>gi|1036799726|ref|XP_017136469.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799708|ref|XP_017136468.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799687|ref|XP_017136467.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799668|ref|XP_017136466.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799744|ref|XP_017136470.1| PREDICTED: isocitrate dehydrogenase [NADP] cytoplasmic isofo
>gi|1036799762|ref|XP_017136471.1| PREDICTED: fibronectin-binding protein A-like [Drosophila mi
>gi|1036843726|ref|XP_017138921.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036843743|ref|XP_017138922.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036843762|ref|XP_017138924.1| PREDICTED: uncharacterized protein LOC108153430 isoform X2
>gi|1036763458|ref|XP_017157067.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036821930|ref|XP_017137692.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036821949|ref|XP_017137693.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036821968|ref|XP_017137694.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036821987|ref|XP_017137695.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036822006|ref|XP_017137696.1| PREDICTED: uncharacterized protein LOC108152696 isoform X1
>gi|1036822023|ref|XP_017137697.1| PREDICTED: uncharacterized protein LOC108152696 isoform X2
>gi|1036822042|ref|XP_017137698.1| PREDICTED: uncharacterized protein LOC108152696 isoform X3
>gi|1036822061|ref|XP_017137699.1| PREDICTED: uncharacterized protein LOC108152698 [Drosophila
>gi|1036795692|ref|XP_017136243.1| PREDICTED: NADPH-dependent diflavin oxidoreductase 1 [Drosop
>gi|1036795619|ref|XP_017136238.1| PREDICTED: intraflagellar transport protein 122 homolog [Dro
>gi|1036795637|ref|XP_017136239.1| PREDICTED: exocyst complex component 7 [Drosophila miranda]
>gi|1036795656|ref|XP_017136240.1| PREDICTED: exocyst complex component 7 [Drosophila miranda]
>gi|1036795711|ref|XP_017136244.1| PREDICTED: protein matrimony [Drosophila miranda]
>gi|1036795673|ref|XP_017136242.1| PREDICTED: agrin [Drosophila miranda]
>gi|1036821777|ref|XP_017137683.1| PREDICTED: E3 ubiquitin-protein ligase CBL-B-B [Drosophila m
>gi|1036818077|ref|XP_017137486.1| PREDICTED: cold shock domain-containing protein E1 isoform X
>gi|1036818060|ref|XP_017137485.1| PREDICTED: cold shock domain-containing protein E1 isoform X
>gi|1036804608|ref|XP_017136745.1| PREDICTED: LOW QUALITY PROTEIN: AFG3-like protein 2 [Drosoph
>gi|1036804645|ref|XP_017136747.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804662|ref|XP_017136748.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804699|ref|XP_017136751.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036804626|ref|XP_017136746.1| PREDICTED: hormone receptor 4 isoform X1 [Drosophila miranda
>gi|1036804681|ref|XP_017136750.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
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>gi|1036804733|ref|XP_017136752.1| PREDICTED: RNA-binding protein Musashi homolog Rbp6 isoform
>gi|1036764499|ref|XP_017157120.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-:
>gi|1036764517|ref|XP_017157121.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-:
>gi|1036764543|ref|XP_017157122.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-:
>gi|1036764563|ref|XP_017157123.1| PREDICTED: 3 beta-hydroxysteroid dehydrogenase/Delta 5--_4-:
>gi|1036830757|ref|XP_017138195.1| PREDICTED: protein rogdi [Drosophila miranda]
>gi|1036830776|ref|XP_017138196.1| PREDICTED: troponin C [Drosophila miranda]
>gi|1036805538|ref|XP_017136798.1| PREDICTED: uncharacterized protein LOC108152178 [Drosophila
>gi|1036805500|ref|XP_017136796.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805479|ref|XP_017136795.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805460|ref|XP_017136793.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805439|ref|XP_017136792.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805421|ref|XP_017136791.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805386|ref|XP_017136789.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805403|ref|XP_017136790.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805367|ref|XP_017136788.1| PREDICTED: multiple epidermal growth factor-like domains pro
>gi|1036805519|ref|XP_017136797.1| PREDICTED: uncharacterized protein LOC108152177 [Drosophila
>gi|1036760069|ref|XP_017156873.1| PREDICTED: uncharacterized protein LOC108165352 [Drosophila
>gi|1036781772|ref|XP_017135452.1| PREDICTED: putative RNA-binding protein Luc7-like 1 [Drosopl
>gi|1036781754|ref|XP_017135450.1| PREDICTED: enhancer of mRNA-decapping protein 3 [Drosophila
>gi|1036781519|ref|XP_017135437.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X1 [D:
>gi|1036781590|ref|XP_017135441.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X5 [D:
>gi|1036781573|ref|XP_017135440.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X4 [D:
>gi|1036781645|ref|XP_017135444.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X8 [D:
>gi|1036781681|ref|XP_017135446.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X10 [I
>gi|1036781538|ref|XP_017135438.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X2 [D:
>gi|1036781737|ref|XP_017135449.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X13 [I
>gi|1036781664|ref|XP_017135445.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X9 [D:
>gi|1036781554|ref|XP_017135439.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X3 [D:
>gi|1036781607|ref|XP_017135442.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X6 [D:
>gi|1036781626|ref|XP_017135443.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X7 [D:
>gi|1036781699|ref|XP_017135447.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X11 [I
>gi|1036781717|ref|XP_017135448.1| PREDICTED: E3 ubiquitin-protein ligase Nedd-4 isoform X12 [I
>gi|1036823803|ref|XP_017137798.1| PREDICTED: uncharacterized protein LOC108152751 [Drosophila
>gi|1036823761|ref|XP_017137796.1| PREDICTED: insulin-degrading enzyme [Drosophila miranda]
>gi|1036823780|ref|XP_017137797.1| PREDICTED: insulin-degrading enzyme [Drosophila miranda]
>gi|1036820849|ref|XP_017137633.1| PREDICTED: rho-related BTB domain-containing protein 1 isofc
>gi|1036820887|ref|XP_017137635.1| PREDICTED: rho-related BTB domain-containing protein 1 isofc
>gi|1036820868|ref|XP_017137634.1| PREDICTED: rho-related BTB domain-containing protein 1 isofc
>gi|1036820905|ref|XP_017137636.1| PREDICTED: charged multivesicular body protein 7 [Drosophila
>gi|1036814073|ref|XP_017137259.1| PREDICTED: pantothenate kinase 3 isoform X1 [Drosophila mira
>gi|1036814091|ref|XP_017137260.1| PREDICTED: pantothenate kinase 3 isoform X2 [Drosophila mira
>gi|1036814138|ref|XP_017137263.1| PREDICTED: pantothenate kinase 3 isoform X4 [Drosophila mira
>gi|1036814176|ref|XP_017137265.1| PREDICTED: pantothenate kinase 3 isoform X6 [Drosophila mira
>gi|1036814117|ref|XP_017137262.1| PREDICTED: pantothenate kinase 3 isoform X3 [Drosophila mira
>gi|1036814157|ref|XP_017137264.1| PREDICTED: pantothenate kinase 3 isoform X5 [Drosophila mira
>gi|1036814194|ref|XP_017137266.1| PREDICTED: F-box/WD repeat-containing protein 4 [Drosophila
>gi|1036827193|ref|XP_017137988.1| PREDICTED: CBL-interacting serine/threonine-protein kinase
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>gi|1036847567|ref|XP_017139121.1| PREDICTED: cationic amino acid transporter 4 [Drosophila mi
>gi|1036843599|ref|XP_017138914.1| PREDICTED: uncharacterized protein LOC108153422 [Drosophila
>gi|1036843618|ref|XP_017138915.1| PREDICTED: dipeptidase 3 [Drosophila miranda]
>gi|1036838554|ref|XP_017138636.1| PREDICTED: uncharacterized protein C7orf26 homolog [Drosoph
>gi|1036838161|ref|XP_017138613.1| PREDICTED: transmembrane protein 104 homolog [Drosophila mi
>gi|1036781297|ref|XP_017135424.1| PREDICTED: lipoyl synthase, mitochondrial [Drosophila miran
>gi|1036781351|ref|XP_017135427.1| PREDICTED: neuronal membrane glycoprotein M6-a isoform X1 [I
>gi|1036781369|ref|XP_017135428.1| PREDICTED: neuronal membrane glycoprotein M6-a isoform X2 [I
>gi|1036781335|ref|XP_017135426.1| PREDICTED: serine/threonine-protein kinase PLK4 [Drosophila
>gi|1036781316|ref|XP_017135425.1| PREDICTED: cyclin-dependent kinase 12 [Drosophila miranda]
>gi|1036808324|ref|XP_017136950.1| PREDICTED: WD repeat-containing protein 26 homolog [Drosoph
>gi|1036808342|ref|XP_017136951.1| PREDICTED: WD repeat-containing protein 26 homolog [Drosoph
>gi|1036808360|ref|XP_017136952.1| PREDICTED: WD repeat-containing protein 26 homolog [Drosoph
>gi|1036808378|ref|XP_017136953.1| PREDICTED: protein ARV1 [Drosophila miranda]
>gi|1036827172|ref|XP_017137987.1| PREDICTED: probable E3 ubiquitin-protein ligase makorin-1 [I
>gi|1036767050|ref|XP_017134610.1| PREDICTED: pickpocket protein 28 isoform X1 [Drosophila mira
>gi|1036767071|ref|XP_017134611.1| PREDICTED: pickpocket protein 28 isoform X2 [Drosophila mira
>gi|1036833247|ref|XP_017138334.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 4 [I
>gi|1036833265|ref|XP_017138335.1| PREDICTED: uncharacterized protein LOC108153066 [Drosophila
>gi|1036847420|ref|XP_017139115.1| PREDICTED: V-type proton ATPase subunit e 2 [Drosophila mira
>gi|1036802904|ref|XP_017136648.1| PREDICTED: serine/threonine-protein kinase WNK1 [Drosophila
>gi|1036802885|ref|XP_017136647.1| PREDICTED: N-acylneuraminate cytidylyltransferase [Drosophil
>gi|1036830998|ref|XP_017138208.1| PREDICTED: neuronal calcium sensor 2 isoform X2 [Drosophila
>gi|1036830981|ref|XP_017138207.1| PREDICTED: neuronal calcium sensor 2 isoform X1 [Drosophila
>gi|1036830962|ref|XP_017138206.1| PREDICTED: neurocalcin homolog [Drosophila miranda]
>gi|1036840070|ref|XP_017138721.1| PREDICTED: 4-aminobutyrate aminotransferase, mitochondrial
>gi|1036840388|ref|XP_017138738.1| PREDICTED: mitochondrial import receptor subunit TOM20 homol
>gi|1036775759|ref|XP_017135093.1| PREDICTED: fat-like cadherin-related tumor suppressor homolo
>gi|1036775778|ref|XP_017135094.1| PREDICTED: uncharacterized protein LOC108151174 [Drosophila
>gi|1036816945|ref|XP_017137423.1| PREDICTED: GDP-Man:Man(3)GlcNAc(2)-PP-Dol alpha-1,2-mannosyl
>gi|1036816964|ref|XP_017137424.1| PREDICTED: uncharacterized protein LOC108152530 [Drosophila
>gi|1036816844|ref|XP_017137417.1| PREDICTED: IQ and AAA domain-containing protein 1-like isof
>gi|1036816863|ref|XP_017137418.1| PREDICTED: IQ and AAA domain-containing protein 1-like isof
>gi|1036816903|ref|XP_017137421.1| PREDICTED: IQ and AAA domain-containing protein 1-like isof
>gi|1036816882|ref|XP_017137420.1| PREDICTED: IQ and AAA domain-containing protein 1-like isof
>gi|1036816926|ref|XP_017137422.1| PREDICTED: opioid-binding protein/cell adhesion molecule-li
>gi|1036835442|ref|XP_017138450.1| PREDICTED: uncharacterized protein LOC108153158 [Drosophila
>gi|1036831455|ref|XP_017138235.1| PREDICTED: membrane-bound transcription factor site-1 prote
>gi|1036845372|ref|XP_017139005.1| PREDICTED: transcription factor AP-2-epsilon isoform X6 [Dr
>gi|1036845336|ref|XP_017139003.1| PREDICTED: transcription factor AP-2-epsilon isoform X4 [Dr
>gi|1036845317|ref|XP_017139001.1| PREDICTED: transcription factor AP-2-epsilon isoform X3 [Dr
>gi|1036845280|ref|XP_017138999.1| PREDICTED: transcription factor AP-2-epsilon isoform X1 [Dr
>gi|1036845355|ref|XP_017139004.1| PREDICTED: transcription factor AP-2-epsilon isoform X5 [Dr
>gi|1036845298|ref|XP_017139000.1| PREDICTED: transcription factor AP-2-epsilon isoform X2 [Dr
>gi|1036841786|ref|XP_017138821.1| PREDICTED: ER membrane protein complex subunit 10 [Drosophil
>gi|1036814985|ref|XP_017137310.1| PREDICTED: uncharacterized protein LOC108152457 [Drosophila
>gi|1036815006|ref|XP_017137311.1| PREDICTED: alanine--glyoxylate aminotransferase 2, mitochond
>gi|1036815027|ref|XP_017137312.1| PREDICTED: alanine--glyoxylate aminotransferase 2, mitochond
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>gi|1036815048|ref|XP_017137313.1| PREDICTED: transmembrane protein 64 [Drosophila miranda]
>gi|1036844266|ref|XP_017138950.1| PREDICTED: uncharacterized protein LOC108153445 isoform X1
>gi|1036844285|ref|XP_017138951.1| PREDICTED: uncharacterized protein LOC108153445 isoform X1
>gi|1036844304|ref|XP_017138952.1| PREDICTED: uncharacterized protein LOC108153445 isoform X2
>gi|1036834508|ref|XP_017138394.1| PREDICTED: transcription elongation factor, mitochondrial [I
>gi|1036834527|ref|XP_017138395.1| PREDICTED: protein maelstrom 1 [Drosophila miranda]
>gi|1036817710|ref|XP_017137465.1| PREDICTED: serine/arginine repetitive matrix protein 1 [Dros
>gi|1036817674|ref|XP_017137463.1| PREDICTED: ADP-ribosylation factor GTPase-activating protei
>gi|1036817691|ref|XP_017137464.1| PREDICTED: ADP-ribosylation factor GTPase-activating protei
>gi|1036817782|ref|XP_017137469.1| PREDICTED: translation initiation factor IF-2 isoform X4 [D]
>gi|1036817763|ref|XP_017137468.1| PREDICTED: salt stress root protein RS1 isoform X3 [Drosophi
>gi|1036817746|ref|XP_017137467.1| PREDICTED: translation initiation factor IF-2 isoform X2 [D]
>gi|1036817729|ref|XP_017137466.1| PREDICTED: salt stress root protein RS1 isoform X1 [Drosophi
>gi|1036760087|ref|XP_017156874.1| PREDICTED: cell wall protein TIR3-like [Drosophila miranda]
>gi|1036769608|ref|XP_017134749.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
>gi|1036760104|ref|XP_017156877.1| PREDICTED: protein new-glue 2-like [Drosophila miranda]
>gi|1036768041|ref|XP_017134660.1| PREDICTED: uncharacterized protein LOC108150868 [Drosophila
>gi|1036760124|ref|XP_017156878.1| PREDICTED: uncharacterized protein LOC108165355 [Drosophila
>gi|1036760145|ref|XP_017156879.1| PREDICTED: circadian clock-controlled protein [Drosophila m
>gi|1036839884|ref|XP_017138710.1| PREDICTED: ribonuclease P protein subunit p29-like [Drosophi
>gi|1036839903|ref|XP_017138711.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosopl
>gi|1036839922|ref|XP_017138712.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosopl
>gi|1036839943|ref|XP_017138713.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosopl
>gi|1036763368|ref|XP_017157062.1| PREDICTED: S-phase kinase-associated protein 1-like [Drosopl
>gi|1036770903|ref|XP_017134820.1| PREDICTED: uncharacterized protein LOC108150985 [Drosophila
>gi|1036760162|ref|XP_017156880.1| PREDICTED: uncharacterized protein LOC108165357 [Drosophila
>gi|1036760180|ref|XP_017156881.1| PREDICTED: uncharacterized protein LOC108165358 [Drosophila
>gi|1036760200|ref|XP_017156882.1| PREDICTED: uncharacterized protein LOC108165359 [Drosophila
>gi|1036770922|ref|XP_017134821.1| PREDICTED: uncharacterized protein LOC108150986 [Drosophila
>gi|1036760218|ref|XP_017156884.1| PREDICTED: uncharacterized protein LOC108165360 [Drosophila
>gi|1036760237|ref|XP_017156885.1| PREDICTED: uncharacterized protein LOC108165361 [Drosophila
>gi|1036760256|ref|XP_017156886.1| PREDICTED: uncharacterized protein LOC108165362 [Drosophila
>gi|1036760275|ref|XP_017156887.1| PREDICTED: uncharacterized protein LOC108165363 [Drosophila
>gi|1036760296|ref|XP_017156888.1| PREDICTED: uncharacterized protein LOC108165364 [Drosophila
>gi|1036760315|ref|XP_017156889.1| PREDICTED: putative uncharacterized protein DDB_G0282129 [D]
>gi|1036852232|ref|XP_017139398.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036850112|ref|XP_017139271.1| PREDICTED: thioredoxin reductase 2, mitochondrial [Drosophil
>gi|1036766462|ref|XP_017134576.1| PREDICTED: uncharacterized protein LOC108150786 [Drosophila
>gi|1036760334|ref|XP_017156890.1| PREDICTED: uncharacterized protein LOC108165366 [Drosophila
>gi|1036760376|ref|XP_017156892.1| PREDICTED: uncharacterized protein LOC108165367 [Drosophila
>gi|1036852195|ref|XP_017139396.1| PREDICTED: uncharacterized protein LOC108153768 [Drosophila
>gi|1036852214|ref|XP_017139397.1| PREDICTED: uncharacterized protein LOC108153770 [Drosophila
>gi|1036766504|ref|XP_017134578.1| PREDICTED: uncharacterized protein LOC108150788 [Drosophila
>gi|1036760395|ref|XP_017156893.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036851018|ref|XP_017139322.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840279|ref|XP_017138733.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840335|ref|XP_017138736.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840314|ref|XP_017138735.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
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>gi|1036840296|ref|XP_017138734.1| PREDICTED: putative phosphatidate phosphatase [Drosophila m
>gi|1036840261|ref|XP_017138732.1| PREDICTED: cilia- and flagella-associated protein 45 [Droso
>gi|1036773303|ref|XP_017134957.1| PREDICTED: teneurin-m isoform X1 [Drosophila miranda]
>gi|1036773322|ref|XP_017134958.1| PREDICTED: teneurin-m isoform X2 [Drosophila miranda]
>gi|1036773341|ref|XP_017134959.1| PREDICTED: teneurin-m isoform X3 [Drosophila miranda]
>gi|1036773360|ref|XP_017134960.1| PREDICTED: teneurin-m isoform X4 [Drosophila miranda]
>gi|1036819799|ref|XP_017137581.1| PREDICTED: isoleucine--tRNA ligase, cytoplasmic [Drosophila
>gi|1036821237|ref|XP_017137654.1| PREDICTED: DDB1- and CUL4-associated factor 8 [Drosophila m
>gi|1036821218|ref|XP_017137652.1| PREDICTED: dynein heavy chain 12, axonemal [Drosophila mira
>gi|1036772576|ref|XP_017134916.1| PREDICTED: ankyrin-1 [Drosophila miranda]
>gi|1036772595|ref|XP_017134917.1| PREDICTED: DNA polymerase delta small subunit [Drosophila m
>gi|1036772637|ref|XP_017134920.1| PREDICTED: U6 snRNA-associated Sm-like protein LSm8 [Droso
>gi|1036772616|ref|XP_017134918.1| PREDICTED: 39S ribosomal protein L46, mitochondrial [Droso
>gi|1036772538|ref|XP_017134914.1| PREDICTED: spectrin alpha chain isoform X4 [Drosophila mira
>gi|1036772521|ref|XP_017134913.1| PREDICTED: spectrin alpha chain isoform X3 [Drosophila mira
>gi|1036772503|ref|XP_017134912.1| PREDICTED: spectrin alpha chain isoform X2 [Drosophila mira
>gi|1036772486|ref|XP_017134911.1| PREDICTED: spectrin alpha chain isoform X1 [Drosophila mira
>gi|1036772557|ref|XP_017134915.1| PREDICTED: protein disks lost [Drosophila miranda]
>gi|1036836935|ref|XP_017138538.1| PREDICTED: LOW QUALITY PROTEIN: ragulator complex protein L
>gi|1036836916|ref|XP_017138537.1| PREDICTED: vacuolar fusion protein CCZ1 homolog [Drosophila
>gi|1036844517|ref|XP_017138963.1| PREDICTED: polyubiquitin [Drosophila miranda]
>gi|1036821406|ref|XP_017137662.1| PREDICTED: very-long-chain enoyl-CoA reductase [Drosophila
>gi|1036821385|ref|XP_017137661.1| PREDICTED: anaphase-promoting complex subunit 5 [Drosophila
>gi|1036821425|ref|XP_017137664.1| PREDICTED: mitochondrial import receptor subunit TOM22 homo
>gi|1036839718|ref|XP_017138701.1| PREDICTED: calcium-binding protein E63-1 isoform X2 [Droso
>gi|1036839699|ref|XP_017138700.1| PREDICTED: calcium-binding protein E63-1 isoform X1 [Droso
>gi|1036839735|ref|XP_017138702.1| PREDICTED: calcium-binding protein E63-1 isoform X3 [Droso
>gi|1036839792|ref|XP_017138705.1| PREDICTED: calcium-binding protein E63-1 isoform X6 [Droso
>gi|1036839773|ref|XP_017138704.1| PREDICTED: calcium-binding protein E63-1 isoform X5 [Droso
>gi|1036839754|ref|XP_017138703.1| PREDICTED: calcium-binding protein E63-1 isoform X4 [Droso
>gi|1036845762|ref|XP_017139019.1| PREDICTED: transmembrane protein C5orf28 homolog [Drosophila
>gi|1036843132|ref|XP_017138885.1| PREDICTED: aldose reductase [Drosophila miranda]
>gi|1036843077|ref|XP_017138882.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036843095|ref|XP_017138883.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036843114|ref|XP_017138884.1| PREDICTED: filaggrin [Drosophila miranda]
>gi|1036770382|ref|XP_017134792.1| PREDICTED: F-box/LRR-repeat protein 17 [Drosophila miranda]
>gi|1036766809|ref|XP_017134595.1| PREDICTED: uncharacterized protein LOC108150802 [Drosophila
>gi|1036841357|ref|XP_017138795.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841374|ref|XP_017138796.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841393|ref|XP_017138797.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841412|ref|XP_017138798.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841432|ref|XP_017138799.1| PREDICTED: zinc finger protein 37 isoform X1 [Drosophila mi
>gi|1036841451|ref|XP_017138800.1| PREDICTED: zinc finger protein 2 isoform X2 [Drosophila mira
>gi|1036838087|ref|XP_017138609.1| PREDICTED: zinc finger protein rotund [Drosophila miranda]
>gi|1036844972|ref|XP_017138980.1| PREDICTED: MIT domain-containing protein 1 [Drosophila mira
>gi|1036824637|ref|XP_017137846.1| PREDICTED: fork head domain-containing protein FD2 isoform
>gi|1036824658|ref|XP_017137848.1| PREDICTED: fork head domain-containing protein FD2 isoform
>gi|1036824677|ref|XP_017137849.1| PREDICTED: mitochondrial ubiquitin ligase activator of NFKB
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>gi|1036824617|ref|XP_017137845.1| PREDICTED: FGGY carbohydrate kinase domain-containing prote:
>gi|1036800785|ref|XP_017136530.1| PREDICTED: uncharacterized protein LOC108152042 [Drosophila
>gi|1036800745|ref|XP_017136528.1| PREDICTED: adenosine 3'-phospho 5'-phosphosulfate transporte
>gi|1036800692|ref|XP_017136525.1| PREDICTED: calcium homeostasis endoplasmic reticulum protei:
>gi|1036800728|ref|XP_017136527.1| PREDICTED: calcium homeostasis endoplasmic reticulum protei:
>gi|1036800710|ref|XP_017136526.1| PREDICTED: calcium homeostasis endoplasmic reticulum protei:
>gi|1036800766|ref|XP_017136529.1| PREDICTED: probable malonyl-CoA-acyl carrier protein transac
>gi|1036842385|ref|XP_017138842.1| PREDICTED: uncharacterized protein LOC108153380 [Drosophila
>gi|1036842367|ref|XP_017138841.1| PREDICTED: uncharacterized protein LOC108153379 [Drosophila
>gi|1036795458|ref|XP_017136229.1| PREDICTED: unc-112-related protein [Drosophila miranda]
>gi|1036795476|ref|XP_017136230.1| PREDICTED: unc-112-related protein [Drosophila miranda]
>gi|1036795494|ref|XP_017136231.1| PREDICTED: uncharacterized protein LOC108151858 [Drosophila
>gi|1036795439|ref|XP_017136228.1| PREDICTED: ribosome biogenesis protein BMS1 homolog [Drosopl
>gi|1036833429|ref|XP_017138344.1| PREDICTED: coiled-coil domain-containing protein 6 [Drosophi
>gi|1036824982|ref|XP_017137863.1| PREDICTED: acetyl-coenzyme A transporter 1 [Drosophila mira:
>gi|1036825001|ref|XP_017137864.1| PREDICTED: eukaryotic translation initiation factor 3 subun:
>gi|1036811610|ref|XP_017137128.1| PREDICTED: glutamate synthase [NADH], amyloplastic isoform 2
>gi|1036811592|ref|XP_017137126.1| PREDICTED: glutamate synthase [NADH], amyloplastic isoform 2
>gi|1036811629|ref|XP_017137129.1| PREDICTED: glutamate synthase 1 [NADH], chloroplastic isofo:
>gi|1036811648|ref|XP_017137130.1| PREDICTED: glutamate synthase 1 [NADH], chloroplastic isofo:
>gi|1036811667|ref|XP_017137131.1| PREDICTED: glutamate synthase 1 [NADH], chloroplastic isofo:
>gi|1036831019|ref|XP_017138210.1| PREDICTED: nuclear migration protein nudC [Drosophila miran
>gi|1036831038|ref|XP_017138211.1| PREDICTED: vitellogenin-2 isoform X1 [Drosophila miranda]
>gi|1036831057|ref|XP_017138212.1| PREDICTED: vitellogenin-2 isoform X2 [Drosophila miranda]
>gi|1036831075|ref|XP_017138213.1| PREDICTED: vitellogenin-2 isoform X2 [Drosophila miranda]
>gi|1036796359|ref|XP_017136279.1| PREDICTED: zinc finger protein CG2199 [Drosophila miranda]
>gi|1036796378|ref|XP_017136280.1| PREDICTED: uncharacterized protein LOC108151890 [Drosophila
>gi|1036796397|ref|XP_017136281.1| PREDICTED: uncharacterized protein LOC108151891 [Drosophila
>gi|1036796418|ref|XP_017136282.1| PREDICTED: uncharacterized protein LOC108151892 [Drosophila
>gi|1036796488|ref|XP_017136286.1| PREDICTED: uncharacterized protein LOC108151895 [Drosophila
>gi|1036796456|ref|XP_017136284.1| PREDICTED: acyl carrier protein, mitochondrial isoform X2 [I
>gi|1036796471|ref|XP_017136285.1| PREDICTED: acyl carrier protein, mitochondrial isoform X3 [I
>gi|1036796437|ref|XP_017136283.1| PREDICTED: acyl carrier protein, mitochondrial isoform X1 [I
>gi|1036776981|ref|XP_017135164.1| PREDICTED: myosin-IB isoform X1 [Drosophila miranda]
>gi|1036776998|ref|XP_017135165.1| PREDICTED: myosin-IB isoform X2 [Drosophila miranda]
>gi|1036777069|ref|XP_017135170.1| PREDICTED: annexin A11 isoform X1 [Drosophila miranda]
>gi|1036777087|ref|XP_017135171.1| PREDICTED: annexin A7 isoform X2 [Drosophila miranda]
>gi|1036776963|ref|XP_017135163.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofor
>gi|1036776945|ref|XP_017135162.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofor
>gi|1036776926|ref|XP_017135161.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofor
>gi|1036776908|ref|XP_017135160.1| PREDICTED: probable E3 ubiquitin-protein ligase HERC4 isofor
>gi|1036777050|ref|XP_017135168.1| PREDICTED: lipid droplet-associated hydrolase [Drosophila m:
>gi|1036776889|ref|XP_017135159.1| PREDICTED: RNA polymerase-associated protein CTR9 homolog [I
>gi|1036777012|ref|XP_017135166.1| PREDICTED: 5'-nucleotidase domain-containing protein 1 [Dros
>gi|1036777031|ref|XP_017135167.1| PREDICTED: acetyl-CoA acetyltransferase, cytosolic [Drosophi
>gi|1036835217|ref|XP_017138436.1| PREDICTED: ras-related protein Rac1 [Drosophila miranda]
>gi|1036841651|ref|XP_017138812.1| PREDICTED: calumenin [Drosophila miranda]
>gi|1036836224|ref|XP_017138497.1| PREDICTED: rab5 GDP/GTP exchange factor [Drosophila miranda]
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>gi|1036852560|ref|XP_017139418.1| PREDICTED: E3 ubiquitin-protein ligase msl-2-like [Drosophi
>gi|1036828205|ref|XP_017138047.1| PREDICTED: uncharacterized protein LOC108152907 isoform X2
>gi|1036828148|ref|XP_017138044.1| PREDICTED: uncharacterized protein LOC108152907 isoform X1
>gi|1036828167|ref|XP_017138045.1| PREDICTED: uncharacterized protein LOC108152907 isoform X1
>gi|1036828186|ref|XP_017138046.1| PREDICTED: uncharacterized protein LOC108152907 isoform X1
>gi|1036828224|ref|XP_017138048.1| PREDICTED: uncharacterized protein LOC108152907 isoform X3
>gi|1036850398|ref|XP_017139286.1| PREDICTED: uncharacterized protein LOC108153681 isoform X1
>gi|1036850417|ref|XP_017139288.1| PREDICTED: perlucin isoform X2 [Drosophila miranda]
>gi|1036827133|ref|XP_017137985.1| PREDICTED: uncharacterized protein LOC108152866 [Drosophila
>gi|1036827114|ref|XP_017137983.1| PREDICTED: uncharacterized protein LOC108152865 [Drosophila
>gi|1036827057|ref|XP_017137980.1| PREDICTED: phosphatidylinositide phosphatase SAC1 isoform X
>gi|1036827076|ref|XP_017137981.1| PREDICTED: phosphatidylinositide phosphatase SAC1 isoform X
>gi|1036827095|ref|XP_017137982.1| PREDICTED: phosphatidylinositide phosphatase SAC1 isoform X
>gi|1036807280|ref|XP_017136893.1| PREDICTED: kinesin-like protein Klp61F isoform X2 [Drosophi
>gi|1036807263|ref|XP_017136892.1| PREDICTED: kinesin-like protein Klp61F isoform X1 [Drosophi
>gi|1036807299|ref|XP_017136894.1| PREDICTED: uncharacterized protein LOC108152223 [Drosophila
>gi|1036807317|ref|XP_017136895.1| PREDICTED: uncharacterized protein LOC108152223 [Drosophila
>gi|1036826333|ref|XP_017137938.1| PREDICTED: UBX domain-containing protein 4 [Drosophila mira
>gi|1036826352|ref|XP_017137939.1| PREDICTED: ribonuclease Oy [Drosophila miranda]
>gi|1036847978|ref|XP_017139145.1| PREDICTED: homeotic protein empty spiracles [Drosophila mira
>gi|1036810752|ref|XP_017137081.1| PREDICTED: signal recognition particle 9 kDa protein [Drosop
>gi|1036810564|ref|XP_017137071.1| PREDICTED: JNK-interacting protein 3 isoform X1 [Drosophila
>gi|1036810583|ref|XP_017137072.1| PREDICTED: JNK-interacting protein 3 isoform X2 [Drosophila
>gi|1036810602|ref|XP_017137073.1| PREDICTED: JNK-interacting protein 3 isoform X3 [Drosophila
>gi|1036810677|ref|XP_017137077.1| PREDICTED: JNK-interacting protein 3 isoform X7 [Drosophila
>gi|1036810640|ref|XP_017137075.1| PREDICTED: JNK-interacting protein 3 isoform X5 [Drosophila
>gi|1036810621|ref|XP_017137074.1| PREDICTED: JNK-interacting protein 3 isoform X4 [Drosophila
>gi|1036810696|ref|XP_017137078.1| PREDICTED: JNK-interacting protein 3 isoform X8 [Drosophila
>gi|1036810714|ref|XP_017137079.1| PREDICTED: JNK-interacting protein 3 isoform X9 [Drosophila
>gi|1036810659|ref|XP_017137076.1| PREDICTED: JNK-interacting protein 3 isoform X6 [Drosophila
>gi|1036810733|ref|XP_017137080.1| PREDICTED: JNK-interacting protein 3 isoform X10 [Drosophila
>gi|1036836561|ref|XP_017138518.1| PREDICTED: transmembrane protein 43 homolog [Drosophila mira
>gi|1036836579|ref|XP_017138519.1| PREDICTED: uncharacterized protein LOC108153208 [Drosophila
>gi|1036798749|ref|XP_017136415.1| PREDICTED: protein ECT2 [Drosophila miranda]
>gi|1036798768|ref|XP_017136416.1| PREDICTED: eukaryotic translation initiation factor 4E [Dros
>gi|1036760408|ref|XP_017156894.1| PREDICTED: uncharacterized protein LOC108165369 [Drosophila
>gi|1036810295|ref|XP_017137055.1| PREDICTED: hepatic leukemia factor isoform X11 [Drosophila r
>gi|1036810312|ref|XP_017137056.1| PREDICTED: hepatic leukemia factor isoform X12 [Drosophila r
>gi|1036810329|ref|XP_017137058.1| PREDICTED: hepatic leukemia factor isoform X13 [Drosophila r
>gi|1036810113|ref|XP_017137045.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X1
>gi|1036810150|ref|XP_017137047.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X3
>gi|1036810222|ref|XP_017137051.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X7
>gi|1036810132|ref|XP_017137046.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X2
>gi|1036810169|ref|XP_017137048.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X4
>gi|1036810187|ref|XP_017137049.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X5
>gi|1036810205|ref|XP_017137050.1| PREDICTED: uncharacterized protein DDB_G0283357 isoform X6
>gi|1036810278|ref|XP_017137054.1| PREDICTED: DNA N6-methyl adenine demethylase isoform X10 [Dr
>gi|1036810242|ref|XP_017137052.1| PREDICTED: transcription factor mef2A isoform X8 [Drosophila
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>gi|1036810261|ref|XP_017137053.1| PREDICTED: transcription factor mef2A isoform X9 [Drosophil
>gi|1036832589|ref|XP_017138296.1| PREDICTED: uncharacterized protein LOC108153045 [Drosophila
>gi|1036832606|ref|XP_017138297.1| PREDICTED: uncharacterized protein LOC108153045 [Drosophila
>gi|1036760426|ref|XP_017156895.1| PREDICTED: circadian locomoter output cycles protein kaput
>gi|1036837419|ref|XP_017138568.1| PREDICTED: protein henna [Drosophila miranda]
>gi|1036837401|ref|XP_017138567.1| PREDICTED: uncharacterized protein LOC108153239 isoform X2
>gi|1036837382|ref|XP_017138566.1| PREDICTED: fibrillin-1 isoform X1 [Drosophila miranda]
>gi|1036837435|ref|XP_017138569.1| PREDICTED: heat shock protein 27 [Drosophila miranda]
>gi|1036760445|ref|XP_017156896.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036848732|ref|XP_017139187.1| PREDICTED: sensory neuron membrane protein 2 isoform X2 [Dro
>gi|1036848695|ref|XP_017139185.1| PREDICTED: sensory neuron membrane protein 2 isoform X1 [Dro
>gi|1036848713|ref|XP_017139186.1| PREDICTED: sensory neuron membrane protein 2 isoform X1 [Dro
>gi|1036848750|ref|XP_017139188.1| PREDICTED: sensory neuron membrane protein 2 isoform X3 [Dro
>gi|1036848771|ref|XP_017139189.1| PREDICTED: sensory neuron membrane protein 2 isoform X4 [Dro
>gi|1036847236|ref|XP_017139103.1| PREDICTED: uncharacterized protein LOC108153548 isoform X1
>gi|1036847253|ref|XP_017139104.1| PREDICTED: uncharacterized protein LOC108153548 isoform X2
>gi|1036760462|ref|XP_017156897.1| PREDICTED: probable serine/threonine-protein kinase DDB_G02
>gi|1036806315|ref|XP_017136838.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806333|ref|XP_017136839.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806375|ref|XP_017136841.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806394|ref|XP_017136843.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036806352|ref|XP_017136840.1| PREDICTED: transcription initiation factor TFIID subunit 4
>gi|1036822288|ref|XP_017137712.1| PREDICTED: WASH complex subunit strumpellin homolog [Drosoph
>gi|1036822307|ref|XP_017137713.1| PREDICTED: phosphoglucomutase [Drosophila miranda]
>gi|1036799926|ref|XP_017136480.1| PREDICTED: LOW QUALITY PROTEIN: nuclear pore complex protein
>gi|1036799944|ref|XP_017136481.1| PREDICTED: translocon-associated protein subunit beta [Droso
>gi|1036799960|ref|XP_017136482.1| PREDICTED: probable trafficking protein particle complex sub
>gi|1036799907|ref|XP_017136479.1| PREDICTED: ATPase family AAA domain-containing protein 5 [Dr
>gi|1036799977|ref|XP_017136484.1| PREDICTED: cytochrome b5 [Drosophila miranda]
>gi|1036779161|ref|XP_017135297.1| PREDICTED: P17/29C-like protein DDB_G0287399 [Drosophila mir
>gi|1036779181|ref|XP_017135298.1| PREDICTED: P17/29C-like protein DDB_G0287399 [Drosophila mir
>gi|1036779144|ref|XP_017135296.1| PREDICTED: LOW QUALITY PROTEIN: protein arginine N-methyltra
>gi|1036779443|ref|XP_017135313.1| PREDICTED: uncharacterized protein LOC108151290 isoform X4
>gi|1036779425|ref|XP_017135312.1| PREDICTED: uncharacterized protein LOC108151290 isoform X3
>gi|1036779459|ref|XP_017135314.1| PREDICTED: uncharacterized protein LOC108151290 isoform X5
>gi|1036779406|ref|XP_017135311.1| PREDICTED: uncharacterized protein LOC108151290 isoform X2
>gi|1036779387|ref|XP_017135310.1| PREDICTED: uncharacterized protein LOC108151290 isoform X1
>gi|1036760478|ref|XP_017156898.1| PREDICTED: uncharacterized protein LOC108165373 [Drosophila
>gi|1036760495|ref|XP_017156900.1| PREDICTED: homeotic protein female sterile [Drosophila miran
>gi|1036779107|ref|XP_017135294.1| PREDICTED: U3 small nucleolar RNA-associated protein 4 homol
>gi|1036779199|ref|XP_017135299.1| PREDICTED: protein disulfide-isomerase TMX3 isoform X1 [Dros
>gi|1036779218|ref|XP_017135300.1| PREDICTED: protein disulfide-isomerase TMX3 isoform X2 [Dros
>gi|1036779604|ref|XP_017135323.1| PREDICTED: programmed cell death protein 5 [Drosophila miran
>gi|1036779641|ref|XP_017135325.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036779348|ref|XP_017135308.1| PREDICTED: iron-sulfur cluster co-chaperone protein HscB, mi
>gi|1036779237|ref|XP_017135301.1| PREDICTED: gamma-glutamyl hydrolase isoform X1 [Drosophila m
>gi|1036779256|ref|XP_017135303.1| PREDICTED: gamma-glutamyl hydrolase A isoform X2 [Drosophila
>gi|1036779312|ref|XP_017135306.1| PREDICTED: gamma-glutamyl hydrolase isoform X1 [Drosophila m
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>gi|1036779329|ref|XP_017135307.1| PREDICTED: gamma-glutamyl hydrolase A isoform X2 [Drosophila
>gi|1036779293|ref|XP_017135305.1| PREDICTED: homeobox protein 5 [Drosophila miranda]
>gi|1036779274|ref|XP_017135304.1| PREDICTED: dentin sialophosphoprotein [Drosophila miranda]
>gi|1036779819|ref|XP_017135336.1| PREDICTED: uncharacterized protein LOC108151309 isoform X2
>gi|1036779804|ref|XP_017135335.1| PREDICTED: uncharacterized protein LOC108151309 isoform X1
>gi|1036779909|ref|XP_017135342.1| PREDICTED: uncharacterized protein LOC108151314 [Drosophila
>gi|1036779928|ref|XP_017135343.1| PREDICTED: uncharacterized protein LOC108151315 [Drosophila
>gi|1036760511|ref|XP_017156901.1| PREDICTED: uncharacterized protein LOC108165375 [Drosophila
>gi|1036779854|ref|XP_017135339.1| PREDICTED: cuticle protein 38 [Drosophila miranda]
>gi|1036779124|ref|XP_017135295.1| PREDICTED: leucine-rich repeat-containing G-protein coupled
>gi|1036779660|ref|XP_017135327.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mi
>gi|1036779837|ref|XP_017135337.1| PREDICTED: EMI domain-containing protein 1 [Drosophila mira
>gi|1036779945|ref|XP_017135344.1| PREDICTED: uncharacterized protein LOC108151316 [Drosophila
>gi|1036779513|ref|XP_017135318.1| PREDICTED: cuticle protein 10.6 [Drosophila miranda]
>gi|1036779549|ref|XP_017135320.1| PREDICTED: glycine-rich protein [Drosophila miranda]
>gi|1036779873|ref|XP_017135340.1| PREDICTED: uncharacterized protein LOC108151312 [Drosophila
>gi|1036779476|ref|XP_017135316.1| PREDICTED: cuticle protein 38 [Drosophila miranda]
>gi|1036760529|ref|XP_017156902.1| PREDICTED: pollen-specific leucine-rich repeat extensin-like
>gi|1036760545|ref|XP_017156903.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mi
>gi|1036779368|ref|XP_017135309.1| PREDICTED: transcription factor btd [Drosophila miranda]
>gi|1036760563|ref|XP_017156904.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036779568|ref|XP_017135321.1| PREDICTED: cuticle protein 10.6 [Drosophila miranda]
>gi|1036779714|ref|XP_017135330.1| PREDICTED: uncharacterized protein LOC108151305 [Drosophila
>gi|1036779532|ref|XP_017135319.1| PREDICTED: pupal cuticle protein C1B [Drosophila miranda]
>gi|1036779587|ref|XP_017135322.1| PREDICTED: cuticle protein LPCP-23 [Drosophila miranda]
>gi|1036779733|ref|XP_017135331.1| PREDICTED: pupal cuticle protein C1B [Drosophila miranda]
>gi|1036779752|ref|XP_017135332.1| PREDICTED: uncharacterized protein LOC108151307 [Drosophila
>gi|1036779771|ref|XP_017135333.1| PREDICTED: uncharacterized protein LOC108151307 [Drosophila
>gi|1036779789|ref|XP_017135334.1| PREDICTED: uncharacterized protein LOC108151307 [Drosophila
>gi|1036779697|ref|XP_017135329.1| PREDICTED: uncharacterized histidine-rich protein DDB_G02745
>gi|1036779891|ref|XP_017135341.1| PREDICTED: neuropeptide-like 3 [Drosophila miranda]
>gi|1036779679|ref|XP_017135328.1| PREDICTED: uncharacterized protein LOC108151303 [Drosophila
>gi|1036779622|ref|XP_017135324.1| PREDICTED: pupal cuticle protein Edg-84A-like [Drosophila mi
>gi|1036779964|ref|XP_017135346.1| PREDICTED: uncharacterized protein LOC108151317 [Drosophila
>gi|1036779494|ref|XP_017135317.1| PREDICTED: cuticle protein LPCP-23 [Drosophila miranda]
>gi|1036768478|ref|XP_017134684.1| PREDICTED: cuticle protein [Drosophila miranda]
>gi|1036766977|ref|XP_017134605.1| PREDICTED: uncharacterized protein LOC108150811 [Drosophila
>gi|1036767983|ref|XP_017134657.1| PREDICTED: uncharacterized protein LOC108150865 [Drosophila
>gi|1036851525|ref|XP_017139354.1| PREDICTED: E3 ubiquitin-protein ligase TRIM33 [Drosophila mi
>gi|1036851506|ref|XP_017139352.1| PREDICTED: trimeric intracellular cation channel type B [Dro
>gi|1036760581|ref|XP_017156905.1| PREDICTED: uncharacterized protein LOC108165379 [Drosophila
>gi|1036851180|ref|XP_017139332.1| PREDICTED: uncharacterized protein LOC108153716 [Drosophila
>gi|1036851199|ref|XP_017139333.1| PREDICTED: uncharacterized protein LOC108153717 [Drosophila
>gi|1036827712|ref|XP_017138019.1| PREDICTED: uncharacterized protein LOC108152891 isoform X2
>gi|1036827694|ref|XP_017138018.1| PREDICTED: uncharacterized protein LOC108152891 isoform X1
>gi|1036760600|ref|XP_017156907.1| PREDICTED: protein anoxia up-regulated [Drosophila miranda]
>gi|1036827750|ref|XP_017138022.1| PREDICTED: uncharacterized protein LOC108152893 [Drosophila
>gi|1036827731|ref|XP_017138020.1| PREDICTED: coiled-coil domain-containing protein 186 [Drosop
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>gi|1036837536|ref|XP_017138575.1| PREDICTED: probable tRNA N6-adenosine threonylcarbamoyltrans
>gi|1036837573|ref|XP_017138578.1| PREDICTED: alcohol dehydrogenase isoform X2 [Drosophila mira
>gi|1036837554|ref|XP_017138576.1| PREDICTED: alcohol dehydrogenase isoform X1 [Drosophila mira
>gi|1036837594|ref|XP_017138579.1| PREDICTED: TPPP family protein CG45057 isoform X1 [Drosophil
>gi|1036837611|ref|XP_017138580.1| PREDICTED: TPPP family protein CG45057 isoform X2 [Drosophil
>gi|1036767887|ref|XP_017134651.1| PREDICTED: histidine-rich glycoprotein [Drosophila miranda]
>gi|1036770141|ref|XP_017134777.1| PREDICTED: adult-specific rigid cuticular protein 15.7 [Dros
>gi|1036760618|ref|XP_017156908.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036851237|ref|XP_017139336.1| PREDICTED: alcohol dehydrogenase-like [Drosophila miranda]
>gi|1036779038|ref|XP_017135289.1| PREDICTED: roquin-1 [Drosophila miranda]
>gi|1036779055|ref|XP_017135291.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036779071|ref|XP_017135292.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036778929|ref|XP_017135283.1| PREDICTED: uncharacterized protein LOC108151271 isoform X1
>gi|1036778981|ref|XP_017135286.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036778999|ref|XP_017135287.1| PREDICTED: 1-acyl-sn-glycerol-3-phosphate acyltransferase ga
>gi|1036779017|ref|XP_017135288.1| PREDICTED: bromodomain-containing protein 4 [Drosophila mira
>gi|1036778945|ref|XP_017135284.1| PREDICTED: uncharacterized protein LOC108151271 isoform X2
>gi|1036778963|ref|XP_017135285.1| PREDICTED: uncharacterized protein LOC108151271 isoform X3
>gi|1036779088|ref|XP_017135293.1| PREDICTED: uncharacterized protein LOC108151276 [Drosophila
>gi|1036811725|ref|XP_017137133.1| PREDICTED: failed axon connections isoform X2 [Drosophila mi
>gi|1036811706|ref|XP_017137132.1| PREDICTED: failed axon connections isoform X1 [Drosophila mi
>gi|1036775285|ref|XP_017135068.1| PREDICTED: serine incorporator 1 isoform X1 [Drosophila mira
>gi|1036775304|ref|XP_017135069.1| PREDICTED: serine incorporator 1 isoform X2 [Drosophila mira
>gi|1036775322|ref|XP_017135070.1| PREDICTED: serine incorporator 1 isoform X3 [Drosophila mira
>gi|1036775251|ref|XP_017135065.1| PREDICTED: probable glutamate--tRNA ligase, mitochondrial [
>gi|1036775233|ref|XP_017135064.1| PREDICTED: tyrosine--tRNA ligase, cytoplasmic [Drosophila mi
>gi|1036775161|ref|XP_017135060.1| PREDICTED: elongation factor-like GTPase 1 [Drosophila miran
>gi|1036775178|ref|XP_017135061.1| PREDICTED: uncharacterized protein LOC108151146 [Drosophila
>gi|1036775269|ref|XP_017135066.1| PREDICTED: protein giant-lens [Drosophila miranda]
>gi|1036775215|ref|XP_017135063.1| PREDICTED: proton-coupled folate transporter [Drosophila mir
>gi|1036775379|ref|XP_017135073.1| PREDICTED: pupal cuticle protein C1B-like [Drosophila miranc
>gi|1036775398|ref|XP_017135074.1| PREDICTED: pupal cuticle protein C1B-like [Drosophila miranc
>gi|1036775196|ref|XP_017135062.1| PREDICTED: protein scarlet [Drosophila miranda]
>gi|1036775086|ref|XP_017135056.1| PREDICTED: uncharacterized protein LOC108151144 isoform X1
>gi|1036775105|ref|XP_017135057.1| PREDICTED: adenylate cyclase type 5 isoform X2 [Drosophila r
>gi|1036775124|ref|XP_017135058.1| PREDICTED: adenylate cyclase type 5 isoform X2 [Drosophila r
>gi|1036775143|ref|XP_017135059.1| PREDICTED: adenylate cyclase type 5 isoform X2 [Drosophila r
>gi|1036775339|ref|XP_017135071.1| PREDICTED: uncharacterized protein LOC108151154 [Drosophila
>gi|1036775358|ref|XP_017135072.1| PREDICTED: uncharacterized protein LOC108151154 [Drosophila
>gi|1036760636|ref|XP_017156909.1| PREDICTED: uncharacterized protein LOC108165382 [Drosophila
>gi|1036789499|ref|XP_017135893.1| PREDICTED: multiple inositol polyphosphate phosphatase 1 [Dr
>gi|1036789588|ref|XP_017135898.1| PREDICTED: LOW QUALITY PROTEIN: endothelial differentiation-
>gi|1036789606|ref|XP_017135900.1| PREDICTED: nuclear RNA export factor 2 [Drosophila miranda]
>gi|1036789553|ref|XP_017135896.1| PREDICTED: survival motor neuron protein [Drosophila miranda
>gi|1036789537|ref|XP_017135895.1| PREDICTED: 26S proteasome non-ATPase regulatory subunit 8 [
>gi|1036789518|ref|XP_017135894.1| PREDICTED: cytochrome b-c1 complex subunit 2, mitochondrial
>gi|1036789571|ref|XP_017135897.1| PREDICTED: syntaxin-8 [Drosophila miranda]
>gi|1036835974|ref|XP_017138482.1| PREDICTED: polycomb group RING finger protein 3 [Drosophila
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>gi|1036835989|ref|XP_017138483.1| PREDICTED: female-specific protein transformer [Drosophila r
>gi|1036780737|ref|XP_017135389.1| PREDICTED: probable GPI-anchored adhesin-like protein PGA55
>gi|1036780755|ref|XP_017135390.1| PREDICTED: importin-4 [Drosophila miranda]
>gi|1036780826|ref|XP_017135395.1| PREDICTED: ADP-ribose pyrophosphatase, mitochondrial isoform
>gi|1036780808|ref|XP_017135394.1| PREDICTED: ADP-ribose pyrophosphatase, mitochondrial isoform
>gi|1036780843|ref|XP_017135396.1| PREDICTED: proteasome subunit beta type-1 [Drosophila mirand
>gi|1036780861|ref|XP_017135397.1| PREDICTED: protein twisted gastrulation [Drosophila miranda]
>gi|1036780772|ref|XP_017135391.1| PREDICTED: RNA-binding protein 45 isoform X1 [Drosophila mir
>gi|1036780790|ref|XP_017135392.1| PREDICTED: RNA-binding protein 45 isoform X2 [Drosophila mir
>gi|1036811554|ref|XP_017137124.1| PREDICTED: general transcription factor IIE subunit 2 [Dros
>gi|1036811496|ref|XP_017137121.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036811515|ref|XP_017137122.1| PREDICTED: chitooligosaccharidolytic beta-N-acetylglucosamin
>gi|1036811573|ref|XP_017137125.1| PREDICTED: transmembrane protein 50A [Drosophila miranda]
>gi|1036811535|ref|XP_017137123.1| PREDICTED: nucleolin 2 [Drosophila miranda]
>gi|1036795399|ref|XP_017136226.1| PREDICTED: LOW QUALITY PROTEIN: salivary glue protein Sgs-3
>gi|1036795380|ref|XP_017136225.1| PREDICTED: sorting nexin lst-4 [Drosophila miranda]
>gi|1036795418|ref|XP_017136227.1| PREDICTED: COPII coat assembly protein SEC16 [Drosophila mi
>gi|1036822794|ref|XP_017137740.1| PREDICTED: uncharacterized protein LOC108152717 isoform X1
>gi|1036822813|ref|XP_017137742.1| PREDICTED: uncharacterized protein LOC108152717 isoform X2
>gi|1036822870|ref|XP_017137745.1| PREDICTED: trafficking protein particle complex subunit 3 [
>gi|1036822832|ref|XP_017137743.1| PREDICTED: nuclear transcription factor Y subunit alpha [Dr
>gi|1036822851|ref|XP_017137744.1| PREDICTED: nuclear transcription factor Y subunit alpha [Dr
>gi|1036829711|ref|XP_017138131.1| PREDICTED: TOM1-like protein 2 isoform X1 [Drosophila miran
>gi|1036829789|ref|XP_017138136.1| PREDICTED: TOM1-like protein 2 isoform X5 [Drosophila miran
>gi|1036829730|ref|XP_017138132.1| PREDICTED: TOM1-like protein 2 isoform X2 [Drosophila miran
>gi|1036829749|ref|XP_017138133.1| PREDICTED: TOM1-like protein 2 isoform X3 [Drosophila miran
>gi|1036829770|ref|XP_017138134.1| PREDICTED: TOM1-like protein 2 isoform X4 [Drosophila miran
>gi|1036829808|ref|XP_017138137.1| PREDICTED: uncharacterized protein LOC108152949 [Drosophila
>gi|1036778100|ref|XP_017135233.1| PREDICTED: uncharacterized protein LOC108151243 [Drosophila
>gi|1036778020|ref|XP_017135228.1| PREDICTED: uncharacterized protein LOC108151239 [Drosophila
>gi|1036778084|ref|XP_017135232.1| PREDICTED: SAC3 domain-containing protein 1 [Drosophila mir
>gi|1036778068|ref|XP_017135231.1| PREDICTED: GDP-D-glucose phosphorylase 1 [Drosophila miranda
>gi|1036778051|ref|XP_017135230.1| PREDICTED: queuine tRNA-ribosyltransferase subunit QTRTD1 h
>gi|1036778132|ref|XP_017135235.1| PREDICTED: uncharacterized protein LOC108151244 isoform X2
>gi|1036778116|ref|XP_017135234.1| PREDICTED: probable serine/threonine-protein kinase nek3 iso
>gi|1036787226|ref|XP_017135762.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase isozyme L5
>gi|1036787207|ref|XP_017135761.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036787243|ref|XP_017135763.1| PREDICTED: F-box only protein 28 [Drosophila miranda]
>gi|1036760655|ref|XP_017156910.1| PREDICTED: uncharacterized protein LOC108165383 [Drosophila
>gi|1036795286|ref|XP_017136220.1| PREDICTED: proton-coupled amino acid transporter 2 isoform
>gi|1036795305|ref|XP_017136221.1| PREDICTED: proton-coupled amino acid transporter 2 isoform
>gi|1036795322|ref|XP_017136223.1| PREDICTED: leucine-rich repeat-containing protein 59 [Drosop
>gi|1036795343|ref|XP_017136224.1| PREDICTED: 40S ribosomal protein S9 [Drosophila miranda]
>gi|1036768060|ref|XP_017134661.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036836076|ref|XP_017138488.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1036836095|ref|XP_017138489.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1036836059|ref|XP_017138487.1| PREDICTED: sodium-independent sulfate anion transporter iso
>gi|1036833285|ref|XP_017138336.1| PREDICTED: CCR4-NOT transcription complex subunit 7 isoform
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>gi|1036833303|ref|XP_017138337.1| PREDICTED: CCR4-NOT transcription complex subunit 7 isoform
>gi|1036805840|ref|XP_017136815.1| PREDICTED: PAX-interacting protein 1 [Drosophila miranda]
>gi|1036788020|ref|XP_017135809.1| PREDICTED: uncharacterized protein LOC108151621 [Drosophila
>gi|1036788039|ref|XP_017135810.1| PREDICTED: uncharacterized protein LOC108151621 [Drosophila
>gi|1036836954|ref|XP_017138539.1| PREDICTED: nitrogen permease regulator 3-like protein [Dros
>gi|1036835497|ref|XP_017138453.1| PREDICTED: zinc finger protein 391 isoform X1 [Drosophila mi
>gi|1036835516|ref|XP_017138454.1| PREDICTED: zinc finger protein 425 isoform X2 [Drosophila mi
>gi|1036760672|ref|XP_017156911.1| PREDICTED: uncharacterized zinc finger protein CG2678 [Dros
>gi|1036836540|ref|XP_017138515.1| PREDICTED: probable 28S ribosomal protein S6, mitochondrial
>gi|1036836502|ref|XP_017138513.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1036836521|ref|XP_017138514.1| PREDICTED: probable RNA methyltransferase CG11342 [Drosophil
>gi|1036843837|ref|XP_017138927.1| PREDICTED: beta-2 adrenergic receptor [Drosophila miranda]
>gi|1036766845|ref|XP_017134597.1| PREDICTED: uncharacterized protein LOC108150805 [Drosophila
>gi|1036844247|ref|XP_017138949.1| PREDICTED: uncharacterized protein LOC108153444 isoform X2
>gi|1036844228|ref|XP_017138948.1| PREDICTED: uncharacterized protein LOC108153444 isoform X1
>gi|1036835592|ref|XP_017138459.1| PREDICTED: syntaxin-17 [Drosophila miranda]
>gi|1036835611|ref|XP_017138460.1| PREDICTED: protein LLP homolog [Drosophila miranda]
>gi|1036824507|ref|XP_017137839.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824526|ref|XP_017137840.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824545|ref|XP_017137841.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824562|ref|XP_017137842.1| PREDICTED: ras guanine nucleotide exchange factor V isoform
>gi|1036824488|ref|XP_017137838.1| PREDICTED: uncharacterized protein LOC108152771 [Drosophila
>gi|1036842053|ref|XP_017138826.1| PREDICTED: acetylcholine receptor subunit beta-like 1 isofo
>gi|1036842034|ref|XP_017138825.1| PREDICTED: acetylcholine receptor subunit beta-like 1 isofo
>gi|1036851781|ref|XP_017139369.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036767905|ref|XP_017134652.1| PREDICTED: cyclin-dependent kinase inhibitor 1C [Drosophila
>gi|1036767944|ref|XP_017134654.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036760691|ref|XP_017156912.1| PREDICTED: early nodulin-75 [Drosophila miranda]
>gi|1036760709|ref|XP_017156913.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036760728|ref|XP_017156914.1| PREDICTED: uncharacterized protein LOC108165388 [Drosophila
>gi|1036851033|ref|XP_017139324.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036851069|ref|XP_017139326.1| PREDICTED: uncharacterized protein LOC108153710 isoform X1
>gi|1036851088|ref|XP_017139327.1| PREDICTED: uncharacterized protein LOC108153710 isoform X2
>gi|1036851050|ref|XP_017139325.1| PREDICTED: skin secretory protein xP2 [Drosophila miranda]
>gi|1036852490|ref|XP_017139413.1| PREDICTED: uncharacterized protein LOC108153782 isoform X1
>gi|1036852508|ref|XP_017139414.1| PREDICTED: uncharacterized protein LOC108153782 isoform X2
>gi|1036760744|ref|XP_017156915.1| PREDICTED: uncharacterized protein LOC108165389 [Drosophila
>gi|1036851164|ref|XP_017139331.1| PREDICTED: LOW QUALITY PROTEIN: extensin-2 [Drosophila mira
>gi|1036846273|ref|XP_017139047.1| PREDICTED: extensin-3 [Drosophila miranda]
>gi|1036769665|ref|XP_017134753.1| PREDICTED: glycine and tyrosine-rich protein [Drosophila mi
>gi|1036760763|ref|XP_017156916.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036850470|ref|XP_017139291.1| PREDICTED: uncharacterized protein LOC108153684 isoform X1
>gi|1036850487|ref|XP_017139292.1| PREDICTED: uncharacterized protein LOC108153684 isoform X1
>gi|1036850506|ref|XP_017139293.1| PREDICTED: uncharacterized protein LOC108153684 isoform X1
>gi|1036850542|ref|XP_017139295.1| PREDICTED: uncharacterized protein LOC108153684 isoform X3
>gi|1036850525|ref|XP_017139294.1| PREDICTED: uncharacterized protein LOC108153684 isoform X2
>gi|1036850559|ref|XP_017139296.1| PREDICTED: uncharacterized protein LOC108153684 isoform X4
>gi|1036844062|ref|XP_017138938.1| PREDICTED: LOW QUALITY PROTEIN: mucin-5AC, partial [Drosophi
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>gi|1036760782|ref|XP_017156918.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosop
>gi|1036819139|ref|XP_017137546.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036819160|ref|XP_017137547.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036819181|ref|XP_017137548.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1036819219|ref|XP_017137551.1| PREDICTED: uncharacterized protein LOC108152603 isoform X1
>gi|1036819240|ref|XP_017137552.1| PREDICTED: uncharacterized protein LOC108152603 isoform X2
>gi|1036819200|ref|XP_017137549.1| PREDICTED: beta-1,3-galactosyltransferase 5 [Drosophila mira
>gi|1036844097|ref|XP_017138940.1| PREDICTED: band 7 protein AGAP004871 isoform X2 [Drosophila
>gi|1036844081|ref|XP_017138939.1| PREDICTED: band 7 protein AGAP004871 isoform X1 [Drosophila
>gi|1036844171|ref|XP_017138945.1| PREDICTED: band 7 protein AGAP004871 isoform X6 [Drosophila
>gi|1036844153|ref|XP_017138944.1| PREDICTED: band 7 protein AGAP004871 isoform X5 [Drosophila
>gi|1036844190|ref|XP_017138946.1| PREDICTED: band 7 protein AGAP004871 isoform X7 [Drosophila
>gi|1036844115|ref|XP_017138941.1| PREDICTED: band 7 protein AGAP004871 isoform X3 [Drosophila
>gi|1036844134|ref|XP_017138942.1| PREDICTED: band 7 protein AGAP004871 isoform X4 [Drosophila
>gi|1036805165|ref|XP_017136776.1| PREDICTED: tyrosine-protein kinase Src64B isoform X2 [Drosop
>gi|1036805146|ref|XP_017136775.1| PREDICTED: tyrosine-protein kinase Src64B isoform X1 [Drosop
>gi|1036826750|ref|XP_017137963.1| PREDICTED: histone deacetylase Rpd3 isoform X1 [Drosophila r
>gi|1036826769|ref|XP_017137964.1| PREDICTED: histone deacetylase Rpd3 isoform X2 [Drosophila r
>gi|1036819572|ref|XP_017137568.1| PREDICTED: uncharacterized protein LOC108152614 isoform X1
>gi|1036819591|ref|XP_017137569.1| PREDICTED: uncharacterized protein LOC108152614 isoform X1
>gi|1036819666|ref|XP_017137574.1| PREDICTED: contactin-associated protein-like 2 isoform X5 [D
>gi|1036819742|ref|XP_017137578.1| PREDICTED: CASP-like protein 4U1 [Drosophila miranda]
>gi|1036819628|ref|XP_017137572.1| PREDICTED: uncharacterized protein LOC108152614 isoform X3
>gi|1036819610|ref|XP_017137570.1| PREDICTED: uncharacterized protein LOC108152614 isoform X2
>gi|1036819647|ref|XP_017137573.1| PREDICTED: contactin-associated protein-like 2 isoform X4 [D
>gi|1036819685|ref|XP_017137575.1| PREDICTED: probable tubulin polyglutamylase TTLL1 [Drosophil
>gi|1036819704|ref|XP_017137576.1| PREDICTED: uncharacterized protein LOC108152616 [Drosophila
>gi|1036819723|ref|XP_017137577.1| PREDICTED: uncharacterized protein LOC108152617 [Drosophila
>gi|1036810941|ref|XP_017137092.1| PREDICTED: uncharacterized protein LOC108152340 [Drosophila
>gi|1036810998|ref|XP_017137096.1| PREDICTED: uncharacterized protein LOC108152343 [Drosophila
>gi|1036810979|ref|XP_017137095.1| PREDICTED: uncharacterized protein LOC108152342 [Drosophila
>gi|1036810960|ref|XP_017137093.1| PREDICTED: uncharacterized protein LOC108152341 [Drosophila
>gi|1036810884|ref|XP_017137089.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036810903|ref|XP_017137090.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036810922|ref|XP_017137091.1| PREDICTED: mucin-19 [Drosophila miranda]
>gi|1036852017|ref|XP_017139385.1| PREDICTED: platelet glycoprotein V isoform X3 [Drosophila mi
>gi|1036851998|ref|XP_017139384.1| PREDICTED: platelet glycoprotein V isoform X2 [Drosophila mi
>gi|1036851981|ref|XP_017139383.1| PREDICTED: platelet glycoprotein V isoform X1 [Drosophila mi
>gi|1036852036|ref|XP_017139386.1| PREDICTED: leucine-rich repeat-containing protein 15 isoform
>gi|1036794285|ref|XP_017136163.1| PREDICTED: uncharacterized oxidoreductase YjmC [Drosophila r
>gi|1036794247|ref|XP_017136161.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X3 [Droso
>gi|1036794266|ref|XP_017136162.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X4 [Droso
>gi|1036794211|ref|XP_017136159.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X1 [Droso
>gi|1036794229|ref|XP_017136160.1| PREDICTED: dynein heavy chain, cytoplasmic isoform X2 [Droso
>gi|1036828986|ref|XP_017138091.1| PREDICTED: probable leucine--tRNA ligase, mitochondrial [Dro
>gi|1036828966|ref|XP_017138090.1| PREDICTED: uncharacterized protein LOC108152921 [Drosophila
>gi|1036848806|ref|XP_017139191.1| PREDICTED: uncharacterized protein LOC108153606 isoform X2
>gi|1036848787|ref|XP_017139190.1| PREDICTED: uncharacterized protein LOC108153606 isoform X1
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>gi|1036839865|ref|XP_017138709.1| PREDICTED: dynein heavy chain 3, axonemal [Drosophila miran
>gi|1036766102|ref|XP_017134558.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036849445|ref|XP_017139230.1| PREDICTED: uncharacterized protein LOC108153637 [Drosophila
>gi|1036848457|ref|XP_017139171.1| PREDICTED: ammonium transporter Rh type B [Drosophila miran
>gi|1036842480|ref|XP_017138848.1| PREDICTED: connectin [Drosophila miranda]
>gi|1036842499|ref|XP_017138849.1| PREDICTED: ubiquitin-conjugating enzyme E2 L3 [Drosophila m
>gi|1036842517|ref|XP_017138850.1| PREDICTED: uncharacterized protein LOC108153387 [Drosophila
>gi|1036760801|ref|XP_017156919.1| PREDICTED: uncharacterized protein LOC108165392 [Drosophila
>gi|1036845189|ref|XP_017138994.1| PREDICTED: uncharacterized protein LOC108153473 [Drosophila
>gi|1036845208|ref|XP_017138995.1| PREDICTED: uncharacterized protein LOC108153473 [Drosophila
>gi|1036845225|ref|XP_017138996.1| PREDICTED: uncharacterized protein LOC108153473 [Drosophila
>gi|1036845171|ref|XP_017138992.1| PREDICTED: glycine-rich cell wall structural protein 1.8 [D
>gi|1036760820|ref|XP_017156920.1| PREDICTED: uncharacterized protein LOC108165393 [Drosophila
>gi|1036815106|ref|XP_017137317.1| PREDICTED: uncharacterized protein LOC108152463 [Drosophila
>gi|1036815127|ref|XP_017137318.1| PREDICTED: uncharacterized protein LOC108152463 [Drosophila
>gi|1036815146|ref|XP_017137319.1| PREDICTED: uncharacterized protein LOC108152463 [Drosophila
>gi|1036815165|ref|XP_017137320.1| PREDICTED: uncharacterized protein LOC108152463 [Drosophila
>gi|1036815278|ref|XP_017137327.1| PREDICTED: lysozyme S [Drosophila miranda]
>gi|1036815259|ref|XP_017137325.1| PREDICTED: lysozyme E-like [Drosophila miranda]
>gi|1036815240|ref|XP_017137324.1| PREDICTED: lysozyme B-like [Drosophila miranda]
>gi|1036815296|ref|XP_017137328.1| PREDICTED: lysozyme D-like [Drosophila miranda]
>gi|1036815395|ref|XP_017137333.1| PREDICTED: lysozyme B-like [Drosophila miranda]
>gi|1036815374|ref|XP_017137332.1| PREDICTED: lysozyme D-like isoform X2 [Drosophila miranda]
>gi|1036815355|ref|XP_017137331.1| PREDICTED: lysozyme D-like isoform X1 [Drosophila miranda]
>gi|1036815202|ref|XP_017137322.1| PREDICTED: lysozyme A/C-like [Drosophila miranda]
>gi|1036815220|ref|XP_017137323.1| PREDICTED: lysozyme A/C-like [Drosophila miranda]
>gi|1036815334|ref|XP_017137330.1| PREDICTED: lysozyme A/C-like isoform X2 [Drosophila miranda]
>gi|1036815315|ref|XP_017137329.1| PREDICTED: lysozyme A/C-like isoform X1 [Drosophila miranda]
>gi|1036815435|ref|XP_017137336.1| PREDICTED: lysozyme A/C-like [Drosophila miranda]
>gi|1036815416|ref|XP_017137334.1| PREDICTED: lysozyme A/C [Drosophila miranda]
>gi|1036815183|ref|XP_017137321.1| PREDICTED: ester hydrolase C11orf54 homolog [Drosophila mira
>gi|1036797501|ref|XP_017136344.1| PREDICTED: mitochondrial carrier homolog 2 [Drosophila miran
>gi|1036797539|ref|XP_017136346.1| PREDICTED: tRNA pseudouridine synthase-like 1 [Drosophila m
>gi|1036797557|ref|XP_017136347.1| PREDICTED: dual specificity protein phosphatase 19 [Drosophi
>gi|1036797393|ref|XP_017136338.1| PREDICTED: uncharacterized protein LOC108151926 isoform X2
>gi|1036797335|ref|XP_017136335.1| PREDICTED: uncharacterized protein LOC108151926 isoform X1
>gi|1036797356|ref|XP_017136336.1| PREDICTED: uncharacterized protein LOC108151926 isoform X1
>gi|1036797376|ref|XP_017136337.1| PREDICTED: uncharacterized protein LOC108151926 isoform X1
>gi|1036797410|ref|XP_017136339.1| PREDICTED: uncharacterized protein LOC108151926 isoform X3
>gi|1036797429|ref|XP_017136340.1| PREDICTED: uncharacterized protein LOC108151926 isoform X3
>gi|1036797447|ref|XP_017136341.1| PREDICTED: uncharacterized protein LOC108151926 isoform X3
>gi|1036797464|ref|XP_017136342.1| PREDICTED: arrestin domain-containing protein 4 [Drosophila
>gi|1036797482|ref|XP_017136343.1| PREDICTED: sulfatase-modifying factor 1 [Drosophila miranda]
>gi|1036797520|ref|XP_017136345.1| PREDICTED: uncharacterized protein LOC108151931 [Drosophila
>gi|1036818759|ref|XP_017137524.1| PREDICTED: breast cancer anti-estrogen resistance protein 1
>gi|1036818778|ref|XP_017137525.1| PREDICTED: breast cancer anti-estrogen resistance protein 1
>gi|1036818797|ref|XP_017137526.1| PREDICTED: breast cancer anti-estrogen resistance protein 1
>gi|1036818816|ref|XP_017137527.1| PREDICTED: WD repeat-containing protein 78 [Drosophila miran
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>gi|1036807754|ref|XP_017136918.1| PREDICTED: 3-phosphoinositide-dependent protein kinase 1 iso
>gi|1036807770|ref|XP_017136919.1| PREDICTED: 3-phosphoinositide-dependent protein kinase 1 iso
>gi|1036852322|ref|XP_017139404.1| PREDICTED: glycine-rich protein 5 [Drosophila miranda]
>gi|1036760839|ref|XP_017156921.1| PREDICTED: kinesin-like protein KIF21A [Drosophila miranda]
>gi|1036760858|ref|XP_017156922.1| PREDICTED: lysozyme A/C-like, partial [Drosophila miranda]
>gi|1036782157|ref|XP_017135472.1| PREDICTED: uncharacterized protein LOC108151396 [Drosophila
>gi|1036782043|ref|XP_017135467.1| PREDICTED: probable G-protein coupled receptor Mth-like 8 [D
>gi|1036782024|ref|XP_017135466.1| PREDICTED: probable G-protein coupled receptor Mth-like 9 [D
>gi|1036781892|ref|XP_017135458.1| PREDICTED: G-protein coupled receptor Mth2 isoform X5 [Droso
>gi|1036781828|ref|XP_017135455.1| PREDICTED: G-protein coupled receptor Mth2 isoform X2 [Droso
>gi|1036781854|ref|XP_017135456.1| PREDICTED: G-protein coupled receptor Mth2 isoform X3 [Droso
>gi|1036781873|ref|XP_017135457.1| PREDICTED: G-protein coupled receptor Mth2 isoform X4 [Droso
>gi|1036781810|ref|XP_017135454.1| PREDICTED: G-protein coupled receptor Mth2 isoform X1 [Droso
>gi|1036781910|ref|XP_017135459.1| PREDICTED: G-protein coupled receptor Mth2 isoform X6 [Droso
>gi|1036781928|ref|XP_017135460.1| PREDICTED: G-protein coupled receptor Mth2 isoform X6 [Droso
>gi|1036781947|ref|XP_017135461.1| PREDICTED: G-protein coupled receptor Mth2 isoform X6 [Droso
>gi|1036782081|ref|XP_017135469.1| PREDICTED: odorant receptor 94b-like isoform X2 [Drosophila
>gi|1036782061|ref|XP_017135468.1| PREDICTED: odorant receptor 94b-like isoform X1 [Drosophila
>gi|1036781966|ref|XP_017135462.1| PREDICTED: G-protein coupled receptor Mth-like isoform X1 [
>gi|1036782005|ref|XP_017135464.1| PREDICTED: G-protein coupled receptor Mth-like isoform X3 [
>gi|1036781986|ref|XP_017135463.1| PREDICTED: G-protein coupled receptor Mth-like isoform X2 [
>gi|1036782136|ref|XP_017135471.1| PREDICTED: G-protein coupled receptor Mth-like isoform X2 [
>gi|1036782099|ref|XP_017135470.1| PREDICTED: G-protein coupled receptor Mth-like isoform X1 [
>gi|1036781791|ref|XP_017135453.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 4
>gi|1036775797|ref|XP_017135095.1| PREDICTED: uncharacterized protein DDB_G0283357 [Drosophila
>gi|1036775912|ref|XP_017135101.1| PREDICTED: probable phospholipid hydroperoxide glutathione p
>gi|1036775893|ref|XP_017135100.1| PREDICTED: probable phospholipid hydroperoxide glutathione p
>gi|1036775930|ref|XP_017135102.1| PREDICTED: probable phospholipid hydroperoxide glutathione p
>gi|1036775816|ref|XP_017135096.1| PREDICTED: striatin-interacting protein 1 homolog [Drosophil
>gi|1036775857|ref|XP_017135098.1| PREDICTED: nicotinamide riboside kinase 2 [Drosophila mirand
>gi|1036775948|ref|XP_017135103.1| PREDICTED: dolichol phosphate-mannose biosynthesis regulato
>gi|1036775874|ref|XP_017135099.1| PREDICTED: phosphatidylinositol N-acetylglucosaminyltransfe
>gi|1036775837|ref|XP_017135097.1| PREDICTED: LOW QUALITY PROTEIN: zinc/cadmium resistance prot
>gi|1036760877|ref|XP_017156923.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036774210|ref|XP_017135011.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774133|ref|XP_017135007.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774152|ref|XP_017135008.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774172|ref|XP_017135009.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774191|ref|XP_017135010.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774229|ref|XP_017135012.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774285|ref|XP_017135016.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774268|ref|XP_017135015.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036774250|ref|XP_017135014.1| PREDICTED: nucleosome-remodeling factor subunit NURF301 iso
>gi|1036802701|ref|XP_017136636.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036760894|ref|XP_017156924.1| PREDICTED: uncharacterized protein LOC108165397 [Drosophila
>gi|1036802737|ref|XP_017136639.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036802718|ref|XP_017136638.1| PREDICTED: ATP-dependent RNA helicase p62-like [Drosophila
>gi|1036802756|ref|XP_017136640.1| PREDICTED: uncharacterized protein LOC108152100 [Drosophila
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>gi|1036837292|ref|XP_017138560.1| PREDICTED: uncharacterized protein LOC108153234 [Drosophila
>gi|1036777416|ref|XP_017135190.1| PREDICTED: follistatin-related protein 1 [Drosophila miranda
>gi|1036777378|ref|XP_017135188.1| PREDICTED: 39S ribosomal protein L17, mitochondrial [Drosoph
>gi|1036777361|ref|XP_017135187.1| PREDICTED: DNA replication complex GINS protein PSF1 [Drosop
>gi|1036777342|ref|XP_017135186.1| PREDICTED: staphylococcal nuclease domain-containing protei
>gi|1036777397|ref|XP_017135189.1| PREDICTED: uncharacterized protein LOC108151219 [Drosophila
>gi|1036777305|ref|XP_017135184.1| PREDICTED: disco-interacting protein 2 isoform X1 [Drosophil
>gi|1036777323|ref|XP_017135185.1| PREDICTED: disco-interacting protein 2 isoform X2 [Drosophil
>gi|1036849839|ref|XP_017139254.1| PREDICTED: probable G-protein coupled receptor Mth-like 14
>gi|1036765116|ref|XP_017157153.1| PREDICTED: ankyrin-1 isoform X1 [Drosophila miranda]
>gi|1036765133|ref|XP_017157154.1| PREDICTED: ankyrin-1 isoform X2 [Drosophila miranda]
>gi|1036765099|ref|XP_017157152.1| PREDICTED: LOW QUALITY PROTEIN: ATP-dependent RNA helicase
>gi|1036765224|ref|XP_017157158.1| PREDICTED: uncharacterized protein LOC108165595 [Drosophila
>gi|1036765023|ref|XP_017157147.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dr
>gi|1036765041|ref|XP_017157148.1| PREDICTED: phosphatidylinositol 4-kinase beta isoform X2 [D
>gi|1036765166|ref|XP_017157156.1| PREDICTED: uncharacterized protein LOC108165592 [Drosophila
>gi|1036765242|ref|XP_017157159.1| PREDICTED: uncharacterized protein LOC108165596 [Drosophila
>gi|1036765190|ref|XP_017157157.1| PREDICTED: uncharacterized protein LOC108165593 [Drosophila
>gi|1036765079|ref|XP_017157151.1| PREDICTED: transient receptor potential channel pyrexia [Dr
>gi|1036765260|ref|XP_017157161.1| PREDICTED: uncharacterized protein LOC108165597 [Drosophila
>gi|1036765276|ref|XP_017157162.1| PREDICTED: uncharacterized protein LOC108165598 [Drosophila
>gi|1036765059|ref|XP_017157150.1| PREDICTED: DIS3-like exonuclease 2 [Drosophila miranda]
>gi|1036765150|ref|XP_017157155.1| PREDICTED: THO complex protein 7 [Drosophila miranda]
>gi|1036785500|ref|XP_017135667.1| PREDICTED: serine/threonine-protein kinase prp4 [Drosophila
>gi|1036785519|ref|XP_017135669.1| PREDICTED: serine/threonine-protein kinase prp4 [Drosophila
>gi|1036785611|ref|XP_017135674.1| PREDICTED: uncharacterized protein C3orf38 homolog [Drosoph
>gi|1036785536|ref|XP_017135670.1| PREDICTED: glycerol kinase [Drosophila miranda]
>gi|1036785573|ref|XP_017135672.1| PREDICTED: nitrilase and fragile histidine triad fusion pro
>gi|1036785554|ref|XP_017135671.1| PREDICTED: nitrilase and fragile histidine triad fusion pro
>gi|1036785592|ref|XP_017135673.1| PREDICTED: BTB/POZ domain-containing protein 10 [Drosophila
>gi|1036796958|ref|XP_017136313.1| PREDICTED: PHD finger protein rhinoceros [Drosophila miranda
>gi|1036796977|ref|XP_017136314.1| PREDICTED: PHD finger protein rhinoceros [Drosophila miranda
>gi|1036796996|ref|XP_017136315.1| PREDICTED: PHD finger protein rhinoceros [Drosophila miranda
>gi|1036810451|ref|XP_017137065.1| PREDICTED: serine/threonine-protein kinase NLK isoform X1 [
>gi|1036810470|ref|XP_017137066.1| PREDICTED: serine/threonine-protein kinase NLK isoform X2 [
>gi|1036833910|ref|XP_017138362.1| PREDICTED: ribonuclease P protein subunit p29 [Drosophila m
>gi|1036833893|ref|XP_017138361.1| PREDICTED: UBX domain-containing protein 1 [Drosophila mira
>gi|1036833929|ref|XP_017138363.1| PREDICTED: uncharacterized protein LOC108153093 [Drosophila
>gi|1036850321|ref|XP_017139282.1| PREDICTED: uncharacterized protein LOC108153677 [Drosophila
>gi|1036850340|ref|XP_017139283.1| PREDICTED: uncharacterized protein LOC108153677 [Drosophila
>gi|1036764753|ref|XP_017157134.1| PREDICTED: titin [Drosophila miranda]
>gi|1036760913|ref|XP_017156926.1| PREDICTED: uncharacterized protein LOC108165398 [Drosophila
>gi|1036767032|ref|XP_017134608.1| PREDICTED: uncharacterized protein LOC108150815 [Drosophila
>gi|1036767811|ref|XP_017134647.1| PREDICTED: uncharacterized protein LOC108150855 [Drosophila
>gi|1036849475|ref|XP_017139231.1| PREDICTED: uncharacterized protein LOC108153639 [Drosophila
>gi|1036791852|ref|XP_017136026.1| PREDICTED: probable protein phosphatase 2C T23F11.1 [Drosop
>gi|1036791793|ref|XP_017136023.1| PREDICTED: la-related protein CG11505 isoform X1 [Drosophila
>gi|1036791831|ref|XP_017136025.1| PREDICTED: la-related protein CG11505 isoform X3 [Drosophila
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>gi|1036791812|ref|XP_017136024.1| PREDICTED: la-related protein CG11505 isoform X2 [Drosophila
>gi|1036845781|ref|XP_017139020.1| PREDICTED: protein translation factor SUI1 homolog [Drosophi
>gi|1036820266|ref|XP_017137607.1| PREDICTED: 60S ribosomal protein L28 [Drosophila miranda]
>gi|1036820191|ref|XP_017137602.1| PREDICTED: putative neutral sphingomyelinase [Drosophila mi
>gi|1036820212|ref|XP_017137604.1| PREDICTED: uncharacterized protein LOC108152635 [Drosophila
>gi|1036820229|ref|XP_017137605.1| PREDICTED: uncharacterized protein LOC108152635 [Drosophila
>gi|1036820247|ref|XP_017137606.1| PREDICTED: uncharacterized protein LOC108152635 [Drosophila
>gi|1036773064|ref|XP_017134943.1| PREDICTED: BAI1-associated protein 3 [Drosophila miranda]
>gi|1036773083|ref|XP_017134944.1| PREDICTED: BAI1-associated protein 3 [Drosophila miranda]
>gi|1036773100|ref|XP_017134945.1| PREDICTED: BAI1-associated protein 3 [Drosophila miranda]
>gi|1036773119|ref|XP_017134946.1| PREDICTED: BAI1-associated protein 3 [Drosophila miranda]
>gi|1036773228|ref|XP_017134952.1| PREDICTED: uncharacterized protein F13E9.13, mitochondrial
>gi|1036773190|ref|XP_017134950.1| PREDICTED: uncharacterized protein LOC108151078 [Drosophila
>gi|1036773172|ref|XP_017134949.1| PREDICTED: RINT1-like protein [Drosophila miranda]
>gi|1036773209|ref|XP_017134951.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036773265|ref|XP_017134955.1| PREDICTED: uncharacterized protein LOC108151083 [Drosophila
>gi|1036773138|ref|XP_017134947.1| PREDICTED: structure-specific endonuclease subunit SLX4 [Dro
>gi|1036773282|ref|XP_017134956.1| PREDICTED: uncharacterized protein LOC108151084 [Drosophila
>gi|1036773247|ref|XP_017134953.1| PREDICTED: uncharacterized protein LOC108151082 [Drosophila
>gi|1036773154|ref|XP_017134948.1| PREDICTED: N-acetyltransferase eco [Drosophila miranda]
>gi|1036801562|ref|XP_017136575.1| PREDICTED: RNA-binding protein lark [Drosophila miranda]
>gi|1036801395|ref|XP_017136566.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801413|ref|XP_017136567.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801433|ref|XP_017136568.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801450|ref|XP_017136569.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801469|ref|XP_017136570.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801488|ref|XP_017136571.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801507|ref|XP_017136572.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801526|ref|XP_017136573.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1036801544|ref|XP_017136574.1| PREDICTED: uncharacterized protein LOC108152065 [Drosophila
>gi|1036801615|ref|XP_017136579.1| PREDICTED: signal recognition particle 19 kDa protein [Droso
>gi|1036801580|ref|XP_017136576.1| PREDICTED: geranylgeranyl pyrophosphate synthase isoform X1
>gi|1036801596|ref|XP_017136577.1| PREDICTED: geranylgeranyl pyrophosphate synthase isoform X2
>gi|1036829556|ref|XP_017138122.1| PREDICTED: transcriptional repressor CTCF [Drosophila miranc
>gi|1036810405|ref|XP_017137062.1| PREDICTED: regulator of chromosome condensation [Drosophila
>gi|1036810392|ref|XP_017137061.1| PREDICTED: LOW QUALITY PROTEIN: motile sperm domain-containi
>gi|1036810419|ref|XP_017137063.1| PREDICTED: motile sperm domain-containing protein 2 [Drosopl
>gi|1036810434|ref|XP_017137064.1| PREDICTED: motile sperm domain-containing protein 2 [Drosopl
>gi|1036849251|ref|XP_017139217.1| PREDICTED: xenotropic and polytropic retrovirus receptor 1
>gi|1036852436|ref|XP_017139411.1| PREDICTED: uncharacterized protein LOC108153780 [Drosophila
>gi|1036760931|ref|XP_017156927.1| PREDICTED: organic cation transporter protein [Drosophila m
>gi|1036833841|ref|XP_017138357.1| PREDICTED: glutaminyl-peptide cyclotransferase-like protein
>gi|1036833805|ref|XP_017138355.1| PREDICTED: glutaminyl-peptide cyclotransferase-like [Drosopl
>gi|1036833823|ref|XP_017138356.1| PREDICTED: ribosome biogenesis regulatory protein homolog [I
>gi|1036766082|ref|XP_017134557.1| PREDICTED: solute carrier family 22 member 2 [Drosophila mi
>gi|1036838372|ref|XP_017138625.1| PREDICTED: DNA polymerase epsilon subunit 2 [Drosophila mira
>gi|1036827576|ref|XP_017138012.1| PREDICTED: transcription factor btd [Drosophila miranda]
>gi|1036827538|ref|XP_017138010.1| PREDICTED: protein vein isoform X4 [Drosophila miranda]
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>gi|1036827520|ref|XP_017138008.1| PREDICTED: protein vein isoform X3 [Drosophila miranda]
>gi|1036827501|ref|XP_017138007.1| PREDICTED: protein vein isoform X2 [Drosophila miranda]
>gi|1036827557|ref|XP_017138011.1| PREDICTED: protein vein isoform X5 [Drosophila miranda]
>gi|1036827489|ref|XP_017138006.1| PREDICTED: protein vein isoform X1 [Drosophila miranda]
>gi|1036792862|ref|XP_017136084.1| PREDICTED: mitotic spindle assembly checkpoint protein MAD2A
>gi|1036792809|ref|XP_017136082.1| PREDICTED: uracil phosphoribosyltransferase homolog [Drosop]
>gi|1036792792|ref|XP_017136081.1| PREDICTED: regulator of chromosome condensation-like [Droso]
>gi|1036792773|ref|XP_017136080.1| PREDICTED: ribosomal protein S6 kinase beta-2 [Drosophila m]
>gi|1036792754|ref|XP_017136079.1| PREDICTED: uncharacterized protein LOC108151752 [Drosophila]
>gi|1036851800|ref|XP_017139370.1| PREDICTED: tektin-1 [Drosophila miranda]
>gi|1036821294|ref|XP_017137657.1| PREDICTED: E3 ubiquitin-protein ligase Bre1 [Drosophila mir]
>gi|1036813489|ref|XP_017137225.1| PREDICTED: odorant receptor 67b [Drosophila miranda]
>gi|1036813449|ref|XP_017137223.1| PREDICTED: laminin subunit gamma-1 [Drosophila miranda]
>gi|1036813466|ref|XP_017137224.1| PREDICTED: laminin subunit gamma-1 [Drosophila miranda]
>gi|1036760947|ref|XP_017156928.1| PREDICTED: uncharacterized protein LOC108165400 [Drosophila]
>gi|1036844022|ref|XP_017138936.1| PREDICTED: MORN repeat-containing protein 5 [Drosophila mir]
>gi|1036844041|ref|XP_017138937.1| PREDICTED: serine proteases 1/2 [Drosophila miranda]
>gi|1036760965|ref|XP_017156929.1| PREDICTED: uncharacterized protein LOC108165401 [Drosophila]
>gi|1036835955|ref|XP_017138481.1| PREDICTED: capon-like protein isoform X4 [Drosophila miranda]
>gi|1036835938|ref|XP_017138480.1| PREDICTED: uncharacterized protein LOC108153181 isoform X3
>gi|1036835904|ref|XP_017138477.1| PREDICTED: capon-like protein isoform X2 [Drosophila miranda]
>gi|1036835921|ref|XP_017138478.1| PREDICTED: capon-like protein isoform X2 [Drosophila miranda]
>gi|1036835887|ref|XP_017138476.1| PREDICTED: capon-like protein isoform X1 [Drosophila miranda]
>gi|1036760983|ref|XP_017156930.1| PREDICTED: uncharacterized protein LOC108165402 [Drosophila]
>gi|1036761002|ref|XP_017156931.1| PREDICTED: uncharacterized protein LOC108165403 [Drosophila]
>gi|1036825921|ref|XP_017137914.1| PREDICTED: DNA replication licensing factor Mcm7-like [Dros]
>gi|1036825807|ref|XP_017137907.1| PREDICTED: protein TANC1 isoform X1 [Drosophila miranda]
>gi|1036825825|ref|XP_017137908.1| PREDICTED: protein TANC1 isoform X1 [Drosophila miranda]
>gi|1036825863|ref|XP_017137910.1| PREDICTED: protein TANC2 isoform X3 [Drosophila miranda]
>gi|1036825844|ref|XP_017137909.1| PREDICTED: protein TANC1 isoform X2 [Drosophila miranda]
>gi|1036825882|ref|XP_017137912.1| PREDICTED: protein TANC2 isoform X4 [Drosophila miranda]
>gi|1036825903|ref|XP_017137913.1| PREDICTED: protein TANC2 isoform X5 [Drosophila miranda]
>gi|1036758102|ref|XP_017156757.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036842109|ref|XP_017138829.1| PREDICTED: isovaleryl-CoA dehydrogenase, mitochondrial [Dros]
>gi|1036815963|ref|XP_017137364.1| PREDICTED: uncharacterized protein LOC108152501 isoform X1
>gi|1036815981|ref|XP_017137365.1| PREDICTED: uncharacterized protein LOC108152501 isoform X1
>gi|1036816038|ref|XP_017137369.1| PREDICTED: uncharacterized protein LOC108152501 isoform X4
>gi|1036816002|ref|XP_017137367.1| PREDICTED: uncharacterized protein LOC108152501 isoform X2
>gi|1036816114|ref|XP_017137373.1| PREDICTED: uncharacterized protein LOC108152501 isoform X7
>gi|1036816077|ref|XP_017137371.1| PREDICTED: uncharacterized protein LOC108152501 isoform X6
>gi|1036816095|ref|XP_017137372.1| PREDICTED: uncharacterized protein LOC108152501 isoform X6
>gi|1036816019|ref|XP_017137368.1| PREDICTED: uncharacterized protein LOC108152501 isoform X3
>gi|1036816056|ref|XP_017137370.1| PREDICTED: uncharacterized protein LOC108152501 isoform X5
>gi|1036816133|ref|XP_017137374.1| PREDICTED: negative elongation factor E [Drosophila miranda]
>gi|1036829693|ref|XP_017138130.1| PREDICTED: 60S ribosomal protein L14 [Drosophila miranda]
>gi|1036829674|ref|XP_017138129.1| PREDICTED: LETM1 domain-containing protein 1 [Drosophila mi]
>gi|1036818740|ref|XP_017137523.1| PREDICTED: uncharacterized protein LOC108152587 [Drosophila]
>gi|1036818639|ref|XP_017137517.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
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>gi|1036818656|ref|XP_017137518.1| PREDICTED: cytosol aminopeptidase [Drosophila miranda]
>gi|1036818677|ref|XP_017137519.1| PREDICTED: TBC1 domain family member 13 isoform X1 [Drosoph
>gi|1036818719|ref|XP_017137522.1| PREDICTED: TBC1 domain family member 13 isoform X3 [Drosoph
>gi|1036818698|ref|XP_017137520.1| PREDICTED: TBC1 domain family member 13 isoform X2 [Drosoph
>gi|1036802318|ref|XP_017136619.1| PREDICTED: uncharacterized protein LOC108152090 isoform X1
>gi|1036802337|ref|XP_017136620.1| PREDICTED: uncharacterized protein LOC108152090 isoform X2
>gi|1036802391|ref|XP_017136623.1| PREDICTED: uncharacterized protein LOC108152090 isoform X5
>gi|1036802409|ref|XP_017136624.1| PREDICTED: uncharacterized protein LOC108152090 isoform X6
>gi|1036802355|ref|XP_017136621.1| PREDICTED: uncharacterized protein LOC108152090 isoform X3
>gi|1036802373|ref|XP_017136622.1| PREDICTED: PDZ and LIM domain protein 2 isoform X4 [Drosoph
>gi|1036802428|ref|XP_017136625.1| PREDICTED: PDZ and LIM domain protein 4 isoform X7 [Drosoph
>gi|1036802447|ref|XP_017136626.1| PREDICTED: uncharacterized protein LOC108152090 isoform X8
>gi|1036802466|ref|XP_017136627.1| PREDICTED: uncharacterized protein LOC108152090 isoform X9
>gi|1036802485|ref|XP_017136628.1| PREDICTED: uncharacterized protein LOC108152090 isoform X10
>gi|1036802299|ref|XP_017136618.1| PREDICTED: cell division control protein 6 homolog [Drosoph
>gi|1036802628|ref|XP_017136633.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036802680|ref|XP_017136635.1| PREDICTED: putative uncharacterized protein DDB_G0294196 iso
>gi|1036802661|ref|XP_017136634.1| PREDICTED: putative uncharacterized protein DDB_G0294196 iso
>gi|1036802572|ref|XP_017136630.1| PREDICTED: phosrestin-1 [Drosophila miranda]
>gi|1036802609|ref|XP_017136632.1| PREDICTED: peroxisomal targeting signal 2 receptor [Drosoph
>gi|1036802590|ref|XP_017136631.1| PREDICTED: protein hairy [Drosophila miranda]
>gi|1036761020|ref|XP_017156932.1| PREDICTED: selenide, water dikinase-like [Drosophila miranda
>gi|1036802110|ref|XP_017136607.1| PREDICTED: signal recognition particle receptor subunit beta
>gi|1036802129|ref|XP_017136608.1| PREDICTED: tail-anchored protein insertion receptor WRB [Dro
>gi|1036802057|ref|XP_017136606.1| PREDICTED: erythroid differentiation-related factor 1 [Droso
>gi|1036802147|ref|XP_017136609.1| PREDICTED: chorion protein S18 [Drosophila miranda]
>gi|1036766994|ref|XP_017134606.1| PREDICTED: chorion protein S15 [Drosophila miranda]
>gi|1036766884|ref|XP_017134600.1| PREDICTED: chorion protein S19 [Drosophila miranda]
>gi|1036766406|ref|XP_017134571.1| PREDICTED: chorion protein S16 [Drosophila miranda]
>gi|1036836335|ref|XP_017138503.1| PREDICTED: paramyosin, long form isoform X1 [Drosophila mira
>gi|1036836354|ref|XP_017138505.1| PREDICTED: paramyosin, short form isoform X2 [Drosophila mir
>gi|1036836373|ref|XP_017138506.1| PREDICTED: uncharacterized protein LOC108153198 [Drosophila
>gi|1036836392|ref|XP_017138507.1| PREDICTED: uncharacterized protein LOC108153198 [Drosophila
>gi|1036836411|ref|XP_017138508.1| PREDICTED: uncharacterized protein LOC108153198 [Drosophila
>gi|1036809164|ref|XP_017136995.1| PREDICTED: uncharacterized protein LOC108152288 isoform X1
>gi|1036809221|ref|XP_017136997.1| PREDICTED: uncharacterized protein LOC108152288 isoform X3
>gi|1036809200|ref|XP_017136996.1| PREDICTED: uncharacterized protein LOC108152288 isoform X2
>gi|1036809297|ref|XP_017137000.1| PREDICTED: uncharacterized protein LOC108152289 [Drosophila
>gi|1036809242|ref|XP_017136998.1| PREDICTED: FH1/FH2 domain-containing protein 1 isoform X4 [
>gi|1036809278|ref|XP_017136999.1| PREDICTED: FH1/FH2 domain-containing protein 3 isoform X5 [
>gi|1036809316|ref|XP_017137001.1| PREDICTED: acyl-CoA-binding protein [Drosophila miranda]
>gi|1036761038|ref|XP_017156933.1| PREDICTED: uncharacterized protein LOC108165406 [Drosophila
>gi|1036761052|ref|XP_017156934.1| PREDICTED: thioredoxin-related transmembrane protein 1-like
>gi|1036764149|ref|XP_017157103.1| PREDICTED: uncharacterized protein LOC108165540 [Drosophila
>gi|1036761068|ref|XP_017156935.1| PREDICTED: mucin-5AC-like [Drosophila miranda]
>gi|1036847066|ref|XP_017139094.1| PREDICTED: uncharacterized protein LOC108153539 [Drosophila
>gi|1036847047|ref|XP_017139092.1| PREDICTED: spore coat protein SP96 [Drosophila miranda]
>gi|1036850895|ref|XP_017139316.1| PREDICTED: uncharacterized protein LOC108153700 [Drosophila
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>gi|1036850913|ref|XP_017139317.1| PREDICTED: uncharacterized protein LOC108153701 [Drosophila
>gi|1036816630|ref|XP_017137405.1| PREDICTED: division abnormally delayed protein [Drosophila
>gi|1036816611|ref|XP_017137404.1| PREDICTED: nascent polypeptide-associated complex subunit al
>gi|1036826844|ref|XP_017137968.1| PREDICTED: DNA replication licensing factor Mcm7 [Drosophila
>gi|1036826825|ref|XP_017137967.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036826806|ref|XP_017137966.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036826788|ref|XP_017137965.1| PREDICTED: transient receptor potential cation channel subfa
>gi|1036849397|ref|XP_017139226.1| PREDICTED: fer-1-like protein 6 [Drosophila miranda]
>gi|1036836244|ref|XP_017138498.1| PREDICTED: tetraspanin-9 [Drosophila miranda]
>gi|1036836262|ref|XP_017138499.1| PREDICTED: tetraspanin-9 [Drosophila miranda]
>gi|1036783249|ref|XP_017135537.1| PREDICTED: peptidyl-prolyl cis-trans isomerase D [Drosophila
>gi|1036783174|ref|XP_017135533.1| PREDICTED: protein furry isoform X7 [Drosophila miranda]
>gi|1036783155|ref|XP_017135532.1| PREDICTED: protein furry isoform X6 [Drosophila miranda]
>gi|1036783117|ref|XP_017135530.1| PREDICTED: protein furry isoform X4 [Drosophila miranda]
>gi|1036783099|ref|XP_017135529.1| PREDICTED: protein furry isoform X3 [Drosophila miranda]
>gi|1036783083|ref|XP_017135527.1| PREDICTED: protein furry isoform X2 [Drosophila miranda]
>gi|1036783068|ref|XP_017135526.1| PREDICTED: protein furry isoform X1 [Drosophila miranda]
>gi|1036783212|ref|XP_017135535.1| PREDICTED: protein furry isoform X9 [Drosophila miranda]
>gi|1036783136|ref|XP_017135531.1| PREDICTED: protein furry isoform X5 [Drosophila miranda]
>gi|1036783193|ref|XP_017135534.1| PREDICTED: protein furry isoform X8 [Drosophila miranda]
>gi|1036783266|ref|XP_017135538.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036783286|ref|XP_017135539.1| PREDICTED: serine proteases 1/2-like [Drosophila miranda]
>gi|1036783230|ref|XP_017135536.1| PREDICTED: uncharacterized protein LOC108151437 [Drosophila
>gi|1036830828|ref|XP_017138198.1| PREDICTED: UPF0046 protein C25E10.12 [Drosophila miranda]
>gi|1036830809|ref|XP_017138197.1| PREDICTED: tubulin alpha-4 chain [Drosophila miranda]
>gi|1036830847|ref|XP_017138199.1| PREDICTED: uncharacterized protein LOC108152984 [Drosophila
>gi|1036817059|ref|XP_017137429.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X4
>gi|1036817040|ref|XP_017137428.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X3
>gi|1036817098|ref|XP_017137431.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X6
>gi|1036817079|ref|XP_017137430.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X5
>gi|1036817021|ref|XP_017137427.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X2
>gi|1036817003|ref|XP_017137426.1| PREDICTED: ribose-phosphate pyrophosphokinase 2 isoform X1
>gi|1036817135|ref|XP_017137434.1| PREDICTED: sperm flagellar protein 1-like [Drosophila mirand
>gi|1036817117|ref|XP_017137432.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa [Drosophil
>gi|1036816983|ref|XP_017137425.1| PREDICTED: uncharacterized protein KIAA1841 homolog [Drosoph
>gi|1036834860|ref|XP_017138416.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834823|ref|XP_017138414.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834768|ref|XP_017138410.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834841|ref|XP_017138415.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834787|ref|XP_017138411.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834879|ref|XP_017138417.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834806|ref|XP_017138413.1| PREDICTED: eukaryotic translation initiation factor 4E isof
>gi|1036834898|ref|XP_017138418.1| PREDICTED: sentrin-specific protease 1-like [Drosophila mir
>gi|1036828817|ref|XP_017138081.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1036803828|ref|XP_017136700.1| PREDICTED: tyrosine-protein kinase Abl isoform X1 [Drosophil
>gi|1036803901|ref|XP_017136704.1| PREDICTED: tyrosine-protein kinase Abl isoform X5 [Drosophil
>gi|1036803862|ref|XP_017136702.1| PREDICTED: tyrosine-protein kinase Abl isoform X3 [Drosophil
>gi|1036803844|ref|XP_017136701.1| PREDICTED: tyrosine-protein kinase Abl isoform X2 [Drosophil
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>gi|1036803881|ref|XP_017136703.1| PREDICTED: tyrosine-protein kinase Abl isoform X4 [Drosophil
>gi|1036819929|ref|XP_017137587.1| PREDICTED: protein Mo25 [Drosophila miranda]
>gi|1036819966|ref|XP_017137589.1| PREDICTED: protein anon-73B1 [Drosophila miranda]
>gi|1036819947|ref|XP_017137588.1| PREDICTED: prostaglandin E synthase 2 [Drosophila miranda]
>gi|1036820833|ref|XP_017137632.1| PREDICTED: ribosome biogenesis protein BRX1 homolog [Drosop]
>gi|1036820815|ref|XP_017137631.1| PREDICTED: CTL-like protein 1 [Drosophila miranda]
>gi|1036766826|ref|XP_017134596.1| PREDICTED: uncharacterized protein LOC108150804 [Drosophila
>gi|1036835079|ref|XP_017138429.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036835098|ref|XP_017138430.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036835116|ref|XP_017138431.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036835134|ref|XP_017138432.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036828909|ref|XP_017138087.1| PREDICTED: V-type proton ATPase subunit e 2 [Drosophila mira
>gi|1036828894|ref|XP_017138086.1| PREDICTED: zinc finger matrin-type protein 2 [Drosophila mi]
>gi|1036828876|ref|XP_017138084.1| PREDICTED: protein ABHD16A [Drosophila miranda]
>gi|1036763440|ref|XP_017157066.1| PREDICTED: uncharacterized protein LOC108165519 isoform X2
>gi|1036763425|ref|XP_017157065.1| PREDICTED: uncharacterized protein LOC108165519 isoform X1
>gi|1036761085|ref|XP_017156937.1| PREDICTED: uncharacterized protein LOC108165409 [Drosophila
>gi|1036780681|ref|XP_017135386.1| PREDICTED: NF-X1-type zinc finger protein NFXL1 [Drosophila
>gi|1036780700|ref|XP_017135387.1| PREDICTED: PQ-loop repeat-containing protein 3 [Drosophila r
>gi|1036780662|ref|XP_017135385.1| PREDICTED: F-box/WD repeat-containing protein 7 [Drosophila
>gi|1036780718|ref|XP_017135388.1| PREDICTED: kinesin-like protein KIF23 [Drosophila miranda]
>gi|1036843149|ref|XP_017138886.1| PREDICTED: sulfide:quinone oxidoreductase, mitochondrial [Dr
>gi|1036785666|ref|XP_017135677.1| PREDICTED: neural/ectodermal development factor IMP-L2 isofo
>gi|1036785685|ref|XP_017135678.1| PREDICTED: neural/ectodermal development factor IMP-L2 isofo
>gi|1036785647|ref|XP_017135676.1| PREDICTED: neurobeachin-like protein 1 isoform X2 [Drosophil
>gi|1036785628|ref|XP_017135675.1| PREDICTED: neurobeachin-like protein 1 isoform X1 [Drosophil
>gi|1036785704|ref|XP_017135679.1| PREDICTED: larval cuticle protein A3A [Drosophila miranda]
>gi|1036785742|ref|XP_017135682.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1036785759|ref|XP_017135683.1| PREDICTED: larval cuticle protein A3A [Drosophila miranda]
>gi|1036785723|ref|XP_017135681.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1036758120|ref|XP_017156758.1| PREDICTED: extensin [Drosophila miranda]
>gi|1036807493|ref|XP_017136904.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807512|ref|XP_017136905.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807531|ref|XP_017136906.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807552|ref|XP_017136907.1| PREDICTED: protein tipE [Drosophila miranda]
>gi|1036807644|ref|XP_017136912.1| PREDICTED: V-type proton ATPase subunit e [Drosophila mirand
>gi|1036807571|ref|XP_017136908.1| PREDICTED: uncharacterized protein LOC108152231 [Drosophila
>gi|1036807588|ref|XP_017136909.1| PREDICTED: uncharacterized protein LOC108152231 [Drosophila
>gi|1036807626|ref|XP_017136911.1| PREDICTED: protein tipE isoform X2 [Drosophila miranda]
>gi|1036807607|ref|XP_017136910.1| PREDICTED: protein tipE isoform X1 [Drosophila miranda]
>gi|1036807407|ref|XP_017136899.1| PREDICTED: uncharacterized protein LOC108152228 isoform X1
>gi|1036807426|ref|XP_017136900.1| PREDICTED: uncharacterized protein LOC108152228 isoform X2
>gi|1036761102|ref|XP_017156938.1| PREDICTED: UPF0769 protein CG18675 [Drosophila miranda]
>gi|1036807444|ref|XP_017136901.1| PREDICTED: ero1-like protein [Drosophila miranda]
>gi|1036807460|ref|XP_017136903.1| PREDICTED: ero1-like protein [Drosophila miranda]
>gi|1036838500|ref|XP_017138633.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1036848898|ref|XP_017139197.1| PREDICTED: NGFI-A-binding protein homolog isoform X1 [Drosop
>gi|1036848914|ref|XP_017139198.1| PREDICTED: NGFI-A-binding protein homolog isoform X2 [Drosop
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>gi|1036821619|ref|XP_017137674.1| PREDICTED: venom serine protease Bi-VSP isoform X2 [Drosophi
>gi|1036821656|ref|XP_017137677.1| PREDICTED: venom protease isoform X4 [Drosophila miranda]
>gi|1036821637|ref|XP_017137676.1| PREDICTED: venom protease isoform X3 [Drosophila miranda]
>gi|1036821600|ref|XP_017137673.1| PREDICTED: venom serine protease Bi-VSP isoform X1 [Drosophi
>gi|1036821757|ref|XP_017137682.1| PREDICTED: venom protease isoform X2 [Drosophila miranda]
>gi|1036821734|ref|XP_017137681.1| PREDICTED: proclotting enzyme isoform X1 [Drosophila miranda
>gi|1036821715|ref|XP_017137680.1| PREDICTED: adipokinetic hormone [Drosophila miranda]
>gi|1036821694|ref|XP_017137679.1| PREDICTED: ras-like protein 2 [Drosophila miranda]
>gi|1036821675|ref|XP_017137678.1| PREDICTED: replication factor C subunit 2 [Drosophila mirand
>gi|1036804292|ref|XP_017136727.1| PREDICTED: flocculation protein FLO11 isoform X1 [Drosophila
>gi|1036804368|ref|XP_017136731.1| PREDICTED: PHD finger protein rhinoceros isoform X5 [Drosoph
>gi|1036804424|ref|XP_017136734.1| PREDICTED: ensconsin isoform X8 [Drosophila miranda]
>gi|1036804460|ref|XP_017136737.1| PREDICTED: ensconsin isoform X10 [Drosophila miranda]
>gi|1036804349|ref|XP_017136730.1| PREDICTED: flocculation protein FLO11 isoform X4 [Drosophila
>gi|1036804405|ref|XP_017136733.1| PREDICTED: ensconsin isoform X7 [Drosophila miranda]
>gi|1036804386|ref|XP_017136732.1| PREDICTED: protein split ends isoform X6 [Drosophila miranda
>gi|1036804443|ref|XP_017136736.1| PREDICTED: ensconsin isoform X9 [Drosophila miranda]
>gi|1036804498|ref|XP_017136739.1| PREDICTED: ensconsin isoform X12 [Drosophila miranda]
>gi|1036804311|ref|XP_017136728.1| PREDICTED: flocculation protein FLO11 isoform X2 [Drosophila
>gi|1036804479|ref|XP_017136738.1| PREDICTED: ensconsin isoform X11 [Drosophila miranda]
>gi|1036804515|ref|XP_017136740.1| PREDICTED: MAP7 domain-containing protein 1 isoform X13 [Dro
>gi|1036804534|ref|XP_017136741.1| PREDICTED: ensconsin isoform X14 [Drosophila miranda]
>gi|1036804330|ref|XP_017136729.1| PREDICTED: flocculation protein FLO11 isoform X3 [Drosophila
>gi|1036804553|ref|XP_017136742.1| PREDICTED: ensconsin isoform X15 [Drosophila miranda]
>gi|1036835461|ref|XP_017138451.1| PREDICTED: myophilin isoform X1 [Drosophila miranda]
>gi|1036835480|ref|XP_017138452.1| PREDICTED: myophilin isoform X2 [Drosophila miranda]
>gi|1036796737|ref|XP_017136299.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC1
>gi|1036796718|ref|XP_017136298.1| PREDICTED: uncharacterized protein F23B12.7 [Drosophila mira
>gi|1036796062|ref|XP_017136263.1| PREDICTED: SPRY domain-containing SOCS box protein 3 [Drosop
>gi|1036796043|ref|XP_017136262.1| PREDICTED: ATP-dependent helicase brm isoform X2 [Drosophila
>gi|1036796025|ref|XP_017136261.1| PREDICTED: ATP-dependent helicase brm isoform X1 [Drosophila
>gi|1036796102|ref|XP_017136265.1| PREDICTED: inositol monophosphatase 1 [Drosophila miranda]
>gi|1036796142|ref|XP_017136268.1| PREDICTED: inositol monophosphatase 2 [Drosophila miranda]
>gi|1036796123|ref|XP_017136267.1| PREDICTED: inositol monophosphatase ttx-7 [Drosophila mirand
>gi|1036796081|ref|XP_017136264.1| PREDICTED: inositol monophosphatase 2 [Drosophila miranda]
>gi|1036819118|ref|XP_017137545.1| PREDICTED: ADP-ribosylation factor-like protein 1 [Drosophil
>gi|1036819099|ref|XP_017137544.1| PREDICTED: DNA polymerase delta catalytic subunit [Drosophil
>gi|1036821795|ref|XP_017137684.1| PREDICTED: palmitoyltransferase Hip14 [Drosophila miranda]
>gi|1036831960|ref|XP_017138261.1| PREDICTED: carboxy-terminal domain RNA polymerase II polypep
>gi|1036831979|ref|XP_017138262.1| PREDICTED: carboxy-terminal domain RNA polymerase II polypep
>gi|1036822269|ref|XP_017137711.1| PREDICTED: 28S ribosomal protein S31, mitochondrial [Drosoph
>gi|1036822250|ref|XP_017137710.1| PREDICTED: E3 ubiquitin-protein ligase mind-bomb [Drosophila
>gi|1036826203|ref|XP_017137930.1| PREDICTED: uncharacterized protein LOC108152836 isoform X2
>gi|1036826180|ref|XP_017137929.1| PREDICTED: uncharacterized protein LOC108152836 isoform X1
>gi|1036761120|ref|XP_017156939.1| PREDICTED: cytosol aminopeptidase-like, partial [Drosophila
>gi|1036761138|ref|XP_017156940.1| PREDICTED: isocitrate dehydrogenase [NAD] catalytic subunit
>gi|1036826259|ref|XP_017137934.1| PREDICTED: hydroxyacylglutathione hydrolase, mitochondrial
>gi|1036826239|ref|XP_017137933.1| PREDICTED: hydroxyacylglutathione hydrolase, mitochondrial
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>gi|1036826278|ref|XP_017137935.1| PREDICTED: hydroxyacylglutathione hydrolase, mitochondrial
>gi|1036826221|ref|XP_017137931.1| PREDICTED: uncharacterized protein LOC108152837 [Drosophila
>gi|1036796788|ref|XP_017136303.1| PREDICTED: homeobox protein SIX6 [Drosophila miranda]
>gi|1036796838|ref|XP_017136306.1| PREDICTED: uncharacterized protein LOC108151906 isoform X2
>gi|1036796821|ref|XP_017136305.1| PREDICTED: uncharacterized protein LOC108151906 isoform X1
>gi|1036796771|ref|XP_017136301.1| PREDICTED: trafficking protein particle complex subunit 12
>gi|1036796754|ref|XP_017136300.1| PREDICTED: phosphorylated CTD-interacting factor 1 [Drosoph
>gi|1036796804|ref|XP_017136304.1| PREDICTED: peroxisomal membrane protein PEX14 [Drosophila m
>gi|1036788984|ref|XP_017135863.1| PREDICTED: peroxisomal membrane protein PEX16 [Drosophila m
>gi|1036788931|ref|XP_017135860.1| PREDICTED: serine/threonine-protein kinase PITSLRE [Drosoph
>gi|1036789059|ref|XP_017135868.1| PREDICTED: ATP synthase subunit s-like protein [Drosophila
>gi|1036789097|ref|XP_017135870.1| PREDICTED: coiled-coil-helix-coiled-coil-helix domain-conta
>gi|1036789003|ref|XP_017135864.1| PREDICTED: Down syndrome critical region protein 3 homolog
>gi|1036788947|ref|XP_017135861.1| PREDICTED: uncharacterized protein LOC108151653 [Drosophila
>gi|1036788966|ref|XP_017135862.1| PREDICTED: uncharacterized protein LOC108151653 [Drosophila
>gi|1036789116|ref|XP_017135871.1| PREDICTED: putative gustatory receptor 77a [Drosophila mira
>gi|1036789021|ref|XP_017135866.1| PREDICTED: uncharacterized protein LOC108151657 [Drosophila
>gi|1036789040|ref|XP_017135867.1| PREDICTED: uncharacterized protein LOC108151657 [Drosophila
>gi|1036789078|ref|XP_017135869.1| PREDICTED: uncharacterized protein LOC108151659 [Drosophila
>gi|1036843582|ref|XP_017138913.1| PREDICTED: uncharacterized protein LOC108153421 [Drosophila
>gi|1036761158|ref|XP_017156941.1| PREDICTED: extracellular sulfatase SULF-1 homolog [Drosophi
>gi|1036843564|ref|XP_017138911.1| PREDICTED: zygotic gap protein knirps [Drosophila miranda]
>gi|1036761178|ref|XP_017156942.1| PREDICTED: uncharacterized protein LOC108165414 [Drosophila
>gi|1036832902|ref|XP_017138313.1| PREDICTED: knirps-related protein [Drosophila miranda]
>gi|1036829827|ref|XP_017138138.1| PREDICTED: cell wall protein IFF6 [Drosophila miranda]
>gi|1036829863|ref|XP_017138140.1| PREDICTED: uncharacterized protein LOC108152952 [Drosophila
>gi|1036829844|ref|XP_017138139.1| PREDICTED: cytosolic Fe-S cluster assembly factor NUBP2 hom
>gi|1036785395|ref|XP_017135661.1| PREDICTED: phosphatidylserine synthase 1 [Drosophila mirand
>gi|1036785483|ref|XP_017135666.1| PREDICTED: vesicle transport protein SFT2A [Drosophila mira
>gi|1036785433|ref|XP_017135663.1| PREDICTED: transcription termination factor 3, mitochondrial
>gi|1036785447|ref|XP_017135664.1| PREDICTED: uncharacterized protein LOC108151526 isoform X1
>gi|1036785464|ref|XP_017135665.1| PREDICTED: uncharacterized protein LOC108151526 isoform X2
>gi|1036785377|ref|XP_017135660.1| PREDICTED: uncharacterized protein LOC108151520 isoform X3
>gi|1036785340|ref|XP_017135658.1| PREDICTED: uncharacterized protein LOC108151520 isoform X1
>gi|1036785358|ref|XP_017135659.1| PREDICTED: C2 domain-containing protein 3 isoform X2 [Droso
>gi|1036785414|ref|XP_017135662.1| PREDICTED: hippocampus abundant transcript-like protein 1 [
>gi|1036839101|ref|XP_017138671.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcompl
>gi|1036800954|ref|XP_017136539.1| PREDICTED: zinc transporter 1 [Drosophila miranda]
>gi|1036800992|ref|XP_017136541.1| PREDICTED: glutaminyl-peptide cyclotransferase isoform X1 [
>gi|1036801010|ref|XP_017136543.1| PREDICTED: glutaminyl-peptide cyclotransferase isoform X2 [
>gi|1036801045|ref|XP_017136545.1| PREDICTED: vacuolar ATPase assembly integral membrane prote
>gi|1036801026|ref|XP_017136544.1| PREDICTED: uncharacterized protein LOC108152051 [Drosophila
>gi|1036800973|ref|XP_017136540.1| PREDICTED: L-threonine 3-dehydrogenase, mitochondrial [Dros
>gi|1036839049|ref|XP_017138669.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1036839009|ref|XP_017138667.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1036839028|ref|XP_017138668.1| PREDICTED: chaoptin [Drosophila miranda]
>gi|1036829250|ref|XP_017138107.1| PREDICTED: uncharacterized protein LOC108152931 [Drosophila
>gi|1036829273|ref|XP_017138108.1| PREDICTED: uncharacterized protein LOC108152931 [Drosophila
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>gi|1036829292|ref|XP_017138109.1| PREDICTED: uncharacterized protein LOC108152932 [Drosophila
>gi|1036829311|ref|XP_017138110.1| PREDICTED: uncharacterized protein LOC108152932 [Drosophila
>gi|1036829328|ref|XP_017138112.1| PREDICTED: uncharacterized protein LOC108152932 [Drosophila
>gi|1036793663|ref|XP_017136126.1| PREDICTED: 4-hydroxyphenylpyruvate dioxygenase [Drosophila r
>gi|1036793624|ref|XP_017136124.1| PREDICTED: uncharacterized protein LOC108151787 isoform X1
>gi|1036793643|ref|XP_017136125.1| PREDICTED: uncharacterized protein LOC108151787 isoform X2
>gi|1036793682|ref|XP_017136127.1| PREDICTED: 39S ribosomal protein L15, mitochondrial [Drosopl
>gi|1036793605|ref|XP_017136123.1| PREDICTED: presenilin homolog isoform X3 [Drosophila miranda
>gi|1036793587|ref|XP_017136122.1| PREDICTED: presenilin homolog isoform X2 [Drosophila miranda
>gi|1036793572|ref|XP_017136121.1| PREDICTED: presenilin homolog isoform X1 [Drosophila miranda
>gi|1036793501|ref|XP_017136117.1| PREDICTED: Golgi apparatus protein 1 isoform X1 [Drosophila
>gi|1036793519|ref|XP_017136118.1| PREDICTED: Golgi apparatus protein 1 isoform X2 [Drosophila
>gi|1036793536|ref|XP_017136119.1| PREDICTED: glutactin isoform X1 [Drosophila miranda]
>gi|1036793555|ref|XP_017136120.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1036831997|ref|XP_017138264.1| PREDICTED: nuclear hormone receptor HR78 isoform X1 [Drosopl
>gi|1036832015|ref|XP_017138265.1| PREDICTED: nuclear hormone receptor HR78 isoform X2 [Drosopl
>gi|1036832053|ref|XP_017138267.1| PREDICTED: protein PBDC1 [Drosophila miranda]
>gi|1036832034|ref|XP_017138266.1| PREDICTED: uncharacterized protein LOC108153023 [Drosophila
>gi|1036800160|ref|XP_017136494.1| PREDICTED: uncharacterized protein LOC108152024 [Drosophila
>gi|1036800068|ref|XP_017136489.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800085|ref|XP_017136490.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800104|ref|XP_017136491.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800123|ref|XP_017136492.1| PREDICTED: speract receptor-like isoform X1 [Drosophila mira
>gi|1036800142|ref|XP_017136493.1| PREDICTED: speract receptor-like isoform X2 [Drosophila mira
>gi|1036813202|ref|XP_017137208.1| PREDICTED: mucolipin-3 [Drosophila miranda]
>gi|1036813276|ref|XP_017137213.1| PREDICTED: uncharacterized protein LOC108152411 [Drosophila
>gi|1036813220|ref|XP_017137209.1| PREDICTED: uncharacterized protein LOC108152410 isoform X1
>gi|1036813238|ref|XP_017137211.1| PREDICTED: uncharacterized protein LOC108152410 isoform X2
>gi|1036813257|ref|XP_017137212.1| PREDICTED: uncharacterized protein LOC108152410 isoform X3
>gi|1036813295|ref|XP_017137214.1| PREDICTED: guanine nucleotide-binding protein subunit beta-2
>gi|1036835535|ref|XP_017138455.1| PREDICTED: uncharacterized protein LOC108153162 [Drosophila
>gi|1036848474|ref|XP_017139172.1| PREDICTED: protein Teyrha-meyrha isoform X1 [Drosophila mira
>gi|1036848551|ref|XP_017139176.1| PREDICTED: protein Teyrha-meyrha isoform X5 [Drosophila mira
>gi|1036848493|ref|XP_017139173.1| PREDICTED: protein Teyrha-meyrha isoform X2 [Drosophila mira
>gi|1036848567|ref|XP_017139177.1| PREDICTED: protein Teyrha-meyrha isoform X6 [Drosophila mira
>gi|1036848513|ref|XP_017139174.1| PREDICTED: protein Teyrha-meyrha isoform X3 [Drosophila mira
>gi|1036848532|ref|XP_017139175.1| PREDICTED: protein Teyrha-meyrha isoform X4 [Drosophila mira
>gi|1036848585|ref|XP_017139178.1| PREDICTED: protein Teyrha-meyrha isoform X7 [Drosophila mira
>gi|1036848603|ref|XP_017139179.1| PREDICTED: protein Teyrha-meyrha isoform X8 [Drosophila mira
>gi|1036848621|ref|XP_017139180.1| PREDICTED: protein Teyrha-meyrha isoform X9 [Drosophila mira
>gi|1036848145|ref|XP_017139153.1| PREDICTED: ornithine aminotransferase, mitochondrial [Drosop
>gi|1036766120|ref|XP_017134559.1| PREDICTED: uncharacterized protein LOC108150774 [Drosophila
>gi|1036817801|ref|XP_017137470.1| PREDICTED: E3 ubiquitin-protein ligase rnf146 isoform X1 [Dr
>gi|1036817821|ref|XP_017137471.1| PREDICTED: E3 ubiquitin-protein ligase RNF146 isoform X2 [Dr
>gi|1036761196|ref|XP_017156943.1| PREDICTED: E3 ubiquitin-protein ligase rnf146-like [Drosophi
>gi|1036783322|ref|XP_017135542.1| PREDICTED: mitogen-activated protein kinase kinase kinase 13
>gi|1036783341|ref|XP_017135543.1| PREDICTED: mitogen-activated protein kinase kinase kinase 13
>gi|1036783360|ref|XP_017135544.1| PREDICTED: mitogen-activated protein kinase kinase kinase 13
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>gi|1036783379|ref|XP_017135545.1| PREDICTED: uncharacterized protein LOC108151444 [Drosophila
>gi|1036783417|ref|XP_017135547.1| PREDICTED: histone chaperone asf1 [Drosophila miranda]
>gi|1036783304|ref|XP_017135540.1| PREDICTED: trafficking protein particle complex subunit 8 [
>gi|1036783398|ref|XP_017135546.1| PREDICTED: glycosylphosphatidylinositol anchor biosynthesis
>gi|1036777143|ref|XP_017135174.1| PREDICTED: LOW QUALITY PROTEIN: lon protease homolog, mitoch
>gi|1036777215|ref|XP_017135178.1| PREDICTED: venom protease [Drosophila miranda]
>gi|1036777268|ref|XP_017135181.1| PREDICTED: uncharacterized protein LOC108151213 isoform X2
>gi|1036777249|ref|XP_017135180.1| PREDICTED: uncharacterized protein LOC108151213 isoform X1
>gi|1036777286|ref|XP_017135183.1| PREDICTED: general odorant-binding protein lush [Drosophila
>gi|1036777180|ref|XP_017135176.1| PREDICTED: transcription initiation factor TFIID subunit 6
>gi|1036777198|ref|XP_017135177.1| PREDICTED: transcription initiation factor TFIID subunit 6
>gi|1036777162|ref|XP_017135175.1| PREDICTED: transcription initiation factor TFIID subunit 6
>gi|1036777104|ref|XP_017135172.1| PREDICTED: histone-lysine N-methyltransferase ash1 isoform
>gi|1036777123|ref|XP_017135173.1| PREDICTED: histone-lysine N-methyltransferase ash1 isoform
>gi|1036777230|ref|XP_017135179.1| PREDICTED: proteasome subunit beta type-1-like [Drosophila
>gi|1036810076|ref|XP_017137043.1| PREDICTED: rRNA methyltransferase 3, mitochondrial [Drosoph
>gi|1036809980|ref|XP_017137038.1| PREDICTED: ATP-binding cassette sub-family F member 3 [Dros
>gi|1036810094|ref|XP_017137044.1| PREDICTED: UPF0389 protein GA21628 [Drosophila miranda]
>gi|1036809999|ref|XP_017137039.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036810017|ref|XP_017137040.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036810038|ref|XP_017137041.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036810057|ref|XP_017137042.1| PREDICTED: potassium voltage-gated channel protein Shal iso
>gi|1036775510|ref|XP_017135080.1| PREDICTED: uncharacterized protein LOC108151161 [Drosophila
>gi|1036775416|ref|XP_017135075.1| PREDICTED: E3 ubiquitin-protein ligase listerin [Drosophila
>gi|1036775624|ref|XP_017135086.1| PREDICTED: uncharacterized protein LOC108151168 [Drosophila
>gi|1036775434|ref|XP_017135076.1| PREDICTED: dynactin subunit 1 [Drosophila miranda]
>gi|1036775453|ref|XP_017135077.1| PREDICTED: fibroin heavy chain [Drosophila miranda]
>gi|1036775472|ref|XP_017135078.1| PREDICTED: fibroin heavy chain [Drosophila miranda]
>gi|1036761214|ref|XP_017156944.1| PREDICTED: LOW QUALITY PROTEIN: MICOS complex subunit Mic10-
>gi|1036775548|ref|XP_017135082.1| PREDICTED: uncharacterized protein LOC108151163 [Drosophila
>gi|1036758138|ref|XP_017156759.1| PREDICTED: adult-specific cuticular protein ACP-20 [Drosophi
>gi|1036775645|ref|XP_017135087.1| PREDICTED: cuticle protein 19.8 [Drosophila miranda]
>gi|1036775567|ref|XP_017135083.1| PREDICTED: venom acid phosphatase Acph-1 [Drosophila miranda
>gi|1036775586|ref|XP_017135084.1| PREDICTED: venom acid phosphatase Acph-1 [Drosophila miranda
>gi|1036775605|ref|XP_017135085.1| PREDICTED: venom acid phosphatase Acph-1 [Drosophila miranda
>gi|1036775529|ref|XP_017135081.1| PREDICTED: sulfhydryl oxidase 1-like [Drosophila miranda]
>gi|1036775491|ref|XP_017135079.1| PREDICTED: polycystin-2 [Drosophila miranda]
>gi|1036761232|ref|XP_017156945.1| PREDICTED: glutaconyl-CoA decarboxylase subunit gamma-like
>gi|1036758155|ref|XP_017156760.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036761251|ref|XP_017156946.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036761270|ref|XP_017156948.1| PREDICTED: cyclin-dependent kinase inhibitor 1C-like [Drosop
>gi|1036761287|ref|XP_017156949.1| PREDICTED: cyclin-dependent kinase inhibitor 1C [Drosophila
>gi|1036849064|ref|XP_017139207.1| PREDICTED: uncharacterized protein LOC108153617 [Drosophila
>gi|1036820061|ref|XP_017137595.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036820003|ref|XP_017137592.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036820042|ref|XP_017137594.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036819985|ref|XP_017137590.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
>gi|1036820023|ref|XP_017137593.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like is
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>gi|1036758171|ref|XP_017156761.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [
>gi|1036820134|ref|XP_017137599.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [
>gi|1036820171|ref|XP_017137601.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [
>gi|1036820153|ref|XP_017137600.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like, 
>gi|1036820116|ref|XP_017137598.1| PREDICTED: heparan sulfate 2-O-sulfotransferase pipe-like [
>gi|1036820080|ref|XP_017137596.1| PREDICTED: deoxynucleoside triphosphate triphosphohydrolase
>gi|1036820098|ref|XP_017137597.1| PREDICTED: deoxynucleoside triphosphate triphosphohydrolase
>gi|1036833503|ref|XP_017138348.1| PREDICTED: glycogen-binding subunit 76A [Drosophila miranda]
>gi|1036833520|ref|XP_017138349.1| PREDICTED: glycogen-binding subunit 76A [Drosophila miranda]
>gi|1036833538|ref|XP_017138350.1| PREDICTED: peroxiredoxin-2 [Drosophila miranda]
>gi|1036761305|ref|XP_017156950.1| PREDICTED: collagen alpha-1(IX) chain [Drosophila miranda]
>gi|1036770105|ref|XP_017134775.1| PREDICTED: uncharacterized protein LOC108150952 [Drosophila
>gi|1036761324|ref|XP_017156951.1| PREDICTED: uncharacterized protein LOC108165423 [Drosophila
>gi|1036798272|ref|XP_017136387.1| PREDICTED: putative aldehyde dehydrogenase family 7 member A
>gi|1036798383|ref|XP_017136394.1| PREDICTED: methyltransferase-like protein 5 [Drosophila mira
>gi|1036798457|ref|XP_017136399.1| PREDICTED: SOSS complex subunit C homolog [Drosophila mirand
>gi|1036798402|ref|XP_017136395.1| PREDICTED: protein max isoform X1 [Drosophila miranda]
>gi|1036798420|ref|XP_017136396.1| PREDICTED: protein max isoform X2 [Drosophila miranda]
>gi|1036798438|ref|XP_017136398.1| PREDICTED: protein max isoform X3 [Drosophila miranda]
>gi|1036798475|ref|XP_017136400.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798494|ref|XP_017136401.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798512|ref|XP_017136402.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798529|ref|XP_017136403.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798547|ref|XP_017136404.1| PREDICTED: BET1 homolog [Drosophila miranda]
>gi|1036798311|ref|XP_017136389.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798328|ref|XP_017136390.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798345|ref|XP_017136391.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798364|ref|XP_017136393.1| PREDICTED: lysophospholipid acyltransferase 5 [Drosophila mi
>gi|1036798292|ref|XP_017136388.1| PREDICTED: ruvB-like helicase 2 [Drosophila miranda]
>gi|1036844935|ref|XP_017138979.1| PREDICTED: 39S ribosomal protein L21, mitochondrial [Drosoph
>gi|1036768023|ref|XP_017134659.1| PREDICTED: uncharacterized protein LOC108150867 [Drosophila
>gi|1036851745|ref|XP_017139367.1| PREDICTED: protein IWS1 homolog [Drosophila miranda]
>gi|1036851762|ref|XP_017139368.1| PREDICTED: uncharacterized protein LOC108153749 [Drosophila
>gi|1036761343|ref|XP_017156952.1| PREDICTED: uncharacterized protein LOC108165425 [Drosophila
>gi|1036815846|ref|XP_017137358.1| PREDICTED: frizzled-2 [Drosophila miranda]
>gi|1036815864|ref|XP_017137359.1| PREDICTED: putative glycine-rich cell wall structural prote
>gi|1036834233|ref|XP_017138382.1| PREDICTED: serpin B6-like isoform X1 [Drosophila miranda]
>gi|1036834252|ref|XP_017138383.1| PREDICTED: serpin B3-like isoform X2 [Drosophila miranda]
>gi|1036761362|ref|XP_017156953.1| PREDICTED: LOW QUALITY PROTEIN: accessory gland protein Acp
>gi|1036834214|ref|XP_017138381.1| PREDICTED: uncharacterized protein LOC108153106 [Drosophila
>gi|1036823262|ref|XP_017137768.1| PREDICTED: UPF0489 protein C5orf22 homolog [Drosophila mira
>gi|1036823318|ref|XP_017137772.1| PREDICTED: protein canopy 4 [Drosophila miranda]
>gi|1036823282|ref|XP_017137770.1| PREDICTED: uncharacterized protein LOC108152736 [Drosophila
>gi|1036823300|ref|XP_017137771.1| PREDICTED: protein Gemin2 [Drosophila miranda]
>gi|1036769533|ref|XP_017134745.1| PREDICTED: dual specificity protein phosphatase Mpk3 [Drosop
>gi|1036769551|ref|XP_017134746.1| PREDICTED: sideroflexin-2 [Drosophila miranda]
>gi|1036769474|ref|XP_017134741.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1036769359|ref|XP_017134735.1| PREDICTED: protein naked cuticle isoform X1 [Drosophila mira
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>gi|1036769380|ref|XP_017134736.1| PREDICTED: protein naked cuticle isoform X2 [Drosophila mira
>gi|1036769397|ref|XP_017134737.1| PREDICTED: protein naked cuticle isoform X3 [Drosophila mira
>gi|1036769515|ref|XP_017134743.1| PREDICTED: protein msta [Drosophila miranda]
>gi|1036769497|ref|XP_017134742.1| PREDICTED: tRNA (uracil-5-)-methyltransferase homolog A [Dro
>gi|1036769456|ref|XP_017134740.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 is
>gi|1036769435|ref|XP_017134739.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 is
>gi|1036769571|ref|XP_017134747.1| PREDICTED: uncharacterized protein LOC108150927 [Drosophila
>gi|1036769416|ref|XP_017134738.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [D
>gi|1036769590|ref|XP_017134748.1| PREDICTED: uncharacterized protein LOC108150928 [Drosophila
>gi|1036845930|ref|XP_017139029.1| PREDICTED: organic solute transporter alpha-like protein [D
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>gi|1036845965|ref|XP_017139031.1| PREDICTED: organic solute transporter alpha-like protein [D
>gi|1036845984|ref|XP_017139032.1| PREDICTED: organic solute transporter alpha-like protein [D
>gi|1036846003|ref|XP_017139033.1| PREDICTED: uncharacterized protein LOC108153494 [Drosophila
>gi|1036846020|ref|XP_017139034.1| PREDICTED: uncharacterized protein LOC108153494 [Drosophila
>gi|1036846037|ref|XP_017139035.1| PREDICTED: uncharacterized protein LOC108153494 [Drosophila
>gi|1036801139|ref|XP_017136551.1| PREDICTED: uncharacterized protein LOC108152057 [Drosophila
>gi|1036761383|ref|XP_017156954.1| PREDICTED: mite allergen Eur m 3 [Drosophila miranda]
>gi|1036801120|ref|XP_017136549.1| PREDICTED: uncharacterized protein LOC108152056 [Drosophila
>gi|1036761402|ref|XP_017156956.1| PREDICTED: uncharacterized protein LOC108165428 [Drosophila
>gi|1036801083|ref|XP_017136547.1| PREDICTED: pre-mRNA-processing factor 6 [Drosophila miranda]
>gi|1036801101|ref|XP_017136548.1| PREDICTED: COP9 signalosome complex subunit 1b [Drosophila m
>gi|1036801158|ref|XP_017136552.1| PREDICTED: corepressor interacting with RBPJ 1 [Drosophila m
>gi|1036833963|ref|XP_017138365.1| PREDICTED: gametocyte-specific factor 1 homolog [Drosophila
>gi|1036833981|ref|XP_017138367.1| PREDICTED: 60S ribosomal protein L26 [Drosophila miranda]
>gi|1036833998|ref|XP_017138368.1| PREDICTED: 60S ribosomal protein L26 [Drosophila miranda]
>gi|1036829091|ref|XP_017138098.1| PREDICTED: uncharacterized protein LOC108152926 [Drosophila
>gi|1036829111|ref|XP_017138099.1| PREDICTED: cell death-inducing p53-target protein 1 [Drosoph
>gi|1036829132|ref|XP_017138100.1| PREDICTED: cell death-inducing p53-target protein 1 [Drosoph
>gi|1036829153|ref|XP_017138101.1| PREDICTED: cell death-inducing p53-target protein 1 [Drosoph
>gi|1036831378|ref|XP_017138230.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1036831397|ref|XP_017138231.1| PREDICTED: extensin isoform X2 [Drosophila miranda]
>gi|1036831357|ref|XP_017138229.1| PREDICTED: integrator complex subunit 3 homolog isoform X1
>gi|1036803424|ref|XP_017136679.1| PREDICTED: uncharacterized protein LOC108152115 [Drosophila
>gi|1036803406|ref|XP_017136678.1| PREDICTED: vacuolar protein-sorting-associated protein 36 [
>gi|1036826090|ref|XP_017137924.1| PREDICTED: liprin-beta-1 [Drosophila miranda]
>gi|1036825997|ref|XP_017137918.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826015|ref|XP_017137919.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826034|ref|XP_017137921.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826052|ref|XP_017137922.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036826071|ref|XP_017137923.1| PREDICTED: uncharacterized protein LOC108152829 [Drosophila
>gi|1036839502|ref|XP_017138688.1| PREDICTED: probable WRKY transcription factor protein 1 [Dro
>gi|1036839483|ref|XP_017138687.1| PREDICTED: CUGBP Elav-like family member 4 isoform X9 [Droso
>gi|1036839428|ref|XP_017138684.1| PREDICTED: CUGBP Elav-like family member 4 isoform X6 [Droso
>gi|1036839392|ref|XP_017138681.1| PREDICTED: CUGBP Elav-like family member 4 isoform X4 [Droso
>gi|1036839356|ref|XP_017138679.1| PREDICTED: CUGBP Elav-like family member 4 isoform X2 [Droso
>gi|1036839464|ref|XP_017138686.1| PREDICTED: CUGBP Elav-like family member 4 isoform X8 [Droso
>gi|1036839409|ref|XP_017138683.1| PREDICTED: CUGBP Elav-like family member 4 isoform X5 [Droso
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>gi|1036839375|ref|XP_017138680.1| PREDICTED: CUGBP Elav-like family member 4 isoform X3 [Dros
>gi|1036839335|ref|XP_017138678.1| PREDICTED: CUGBP Elav-like family member 4 isoform X1 [Dros
>gi|1036839447|ref|XP_017138685.1| PREDICTED: CUGBP Elav-like family member 4 isoform X7 [Dros
>gi|1036852108|ref|XP_017139391.1| PREDICTED: uncharacterized protein LOC108153763 [Drosophila
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>gi|1036767540|ref|XP_017134635.1| PREDICTED: mastermind-like domain-containing protein 1 isof
>gi|1036818023|ref|XP_017137483.1| PREDICTED: ERAD-associated E3 ubiquitin-protein ligase HRD1A
>gi|1036818004|ref|XP_017137482.1| PREDICTED: protein panoramix [Drosophila miranda]
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>gi|1036768739|ref|XP_017134698.1| PREDICTED: protein new-glue 2-like [Drosophila miranda]
>gi|1036768721|ref|XP_017134697.1| PREDICTED: protein new-glue 1-like [Drosophila miranda]
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>gi|1036834733|ref|XP_017138408.1| PREDICTED: hemocytin [Drosophila miranda]
>gi|1036840869|ref|XP_017138766.1| PREDICTED: GPN-loop GTPase 2 [Drosophila miranda]
>gi|1036830865|ref|XP_017138201.1| PREDICTED: actin-interacting protein 1 [Drosophila miranda]
>gi|1036843908|ref|XP_017138929.1| PREDICTED: uncharacterized protein LOC108153435 [Drosophila
>gi|1036843927|ref|XP_017138930.1| PREDICTED: uncharacterized protein LOC108153435 [Drosophila
>gi|1036844003|ref|XP_017138935.1| PREDICTED: uncharacterized protein LOC108153438 [Drosophila
>gi|1036843948|ref|XP_017138931.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036843966|ref|XP_017138932.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036843984|ref|XP_017138933.1| PREDICTED: centrosomal and chromosomal factor [Drosophila mi
>gi|1036765677|ref|XP_017134533.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
>gi|1036765487|ref|XP_017157174.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
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>gi|1036844463|ref|XP_017138960.1| PREDICTED: uncharacterized protein LOC108153451 [Drosophila
>gi|1036781054|ref|XP_017135410.1| PREDICTED: uncharacterized protein LOC108151362 [Drosophila
>gi|1036781147|ref|XP_017135415.1| PREDICTED: jmjC domain-containing protein 7 [Drosophila mira
>gi|1036781165|ref|XP_017135416.1| PREDICTED: GSK3-beta interaction protein [Drosophila miranda
>gi|1036781129|ref|XP_017135414.1| PREDICTED: ATP-dependent RNA helicase Ddx1 [Drosophila miran
>gi|1036781073|ref|XP_017135411.1| PREDICTED: guanine nucleotide exchange factor subunit Rich
>gi|1036781091|ref|XP_017135412.1| PREDICTED: guanine nucleotide exchange factor subunit Rich
>gi|1036781110|ref|XP_017135413.1| PREDICTED: guanine nucleotide exchange factor subunit Rich
>gi|1036820962|ref|XP_017137639.1| PREDICTED: MOXD1 homolog 2 [Drosophila miranda]
>gi|1036771162|ref|XP_017134835.1| PREDICTED: 60S acidic ribosomal protein P0 [Drosophila miran
>gi|1036771032|ref|XP_017134828.1| PREDICTED: NEDD4-binding protein 2 [Drosophila miranda]
>gi|1036771051|ref|XP_017134829.1| PREDICTED: NEDD4-binding protein 2 [Drosophila miranda]
>gi|1036771195|ref|XP_017134838.1| PREDICTED: uncharacterized protein LOC108150997 isoform X2
>gi|1036771177|ref|XP_017134837.1| PREDICTED: uncharacterized protein LOC108150997 isoform X1
>gi|1036771214|ref|XP_017134839.1| PREDICTED: tyramine/octopamine receptor [Drosophila miranda]
>gi|1036771125|ref|XP_017134833.1| PREDICTED: glucose-6-phosphate 1-dehydrogenase [Drosophila m
>gi|1036771144|ref|XP_017134834.1| PREDICTED: actin, larval muscle [Drosophila miranda]
>gi|1036771070|ref|XP_017134830.1| PREDICTED: delta-1-pyrroline-5-carboxylate synthase [Drosoph
>gi|1036771013|ref|XP_017134827.1| PREDICTED: uncharacterized protein LOC108150988 isoform X2
>gi|1036770993|ref|XP_017134826.1| PREDICTED: uncharacterized protein LOC108150988 isoform X1
>gi|1036771088|ref|XP_017134831.1| PREDICTED: organic cation transporter protein [Drosophila m
>gi|1036771106|ref|XP_017134832.1| PREDICTED: solute carrier family 22 member 3-like [Drosophil
>gi|1036770941|ref|XP_017134822.1| PREDICTED: uncharacterized protein LOC108150987 [Drosophila
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>gi|1036770959|ref|XP_017134823.1| PREDICTED: uncharacterized protein LOC108150987 [Drosophila
>gi|1036770976|ref|XP_017134825.1| PREDICTED: uncharacterized protein LOC108150987 [Drosophila
>gi|1036761438|ref|XP_017156958.1| PREDICTED: vesicle-associated membrane protein-associated pi
>gi|1036794440|ref|XP_017136171.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794516|ref|XP_017136175.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794535|ref|XP_017136176.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
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>gi|1036794478|ref|XP_017136173.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794584|ref|XP_017136179.1| PREDICTED: sterile alpha and TIR motif-containing protein 1
>gi|1036794600|ref|XP_017136180.1| PREDICTED: dyslexia-associated protein KIAA0319 [Drosophila
>gi|1036794675|ref|XP_017136185.1| PREDICTED: uncharacterized protein LOC108151827 [Drosophila
>gi|1036794618|ref|XP_017136181.1| PREDICTED: uncharacterized protein LOC108151826 isoform X1
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>gi|1036794656|ref|XP_017136183.1| PREDICTED: uncharacterized protein LOC108151826 isoform X2
>gi|1036765908|ref|XP_017134546.1| PREDICTED: borealin-like [Drosophila miranda]
>gi|1036848316|ref|XP_017139163.1| PREDICTED: JNK-interacting protein 1 isoform X2 [Drosophila
>gi|1036848297|ref|XP_017139162.1| PREDICTED: JNK-interacting protein 1 isoform X1 [Drosophila
>gi|1036847365|ref|XP_017139111.1| PREDICTED: tryptophan 5-hydroxylase 1 [Drosophila miranda]
>gi|1036847383|ref|XP_017139112.1| PREDICTED: uncharacterized protein LOC108153553 [Drosophila
>gi|1036826865|ref|XP_017137969.1| PREDICTED: protein bric-a-brac 2 isoform X1 [Drosophila mira
>gi|1036826884|ref|XP_017137970.1| PREDICTED: protein bric-a-brac 2 isoform X1 [Drosophila mira
>gi|1036826904|ref|XP_017137972.1| PREDICTED: protein bric-a-brac 2 isoform X2 [Drosophila mira
>gi|1036761456|ref|XP_017156959.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosopl
>gi|1036847142|ref|XP_017139098.1| PREDICTED: protein bric-a-brac 1 isoform X1 [Drosophila mira
>gi|1036847161|ref|XP_017139099.1| PREDICTED: protein bric-a-brac 1 isoform X2 [Drosophila mira
>gi|1036845594|ref|XP_017139009.1| PREDICTED: oxysterol-binding protein-related protein 11 [Dro
>gi|1036788185|ref|XP_017135817.1| PREDICTED: triple functional domain protein isoform X1 [Dros
>gi|1036788204|ref|XP_017135818.1| PREDICTED: triple functional domain protein isoform X2 [Dros
>gi|1036788223|ref|XP_017135819.1| PREDICTED: triple functional domain protein isoform X3 [Dros
>gi|1036788242|ref|XP_017135821.1| PREDICTED: rho guanine nucleotide exchange factor 25 isoforn
>gi|1036788300|ref|XP_017135824.1| PREDICTED: lipase member I [Drosophila miranda]
>gi|1036788280|ref|XP_017135823.1| PREDICTED: microfibril-associated glycoprotein 4 isoform X2
>gi|1036788261|ref|XP_017135822.1| PREDICTED: fibrinogen C domain-containing protein 1 isoform
>gi|1036796658|ref|XP_017136295.1| PREDICTED: cyclin-dependent kinase 14 isoform X3 [Drosophila
>gi|1036796640|ref|XP_017136294.1| PREDICTED: cyclin-dependent kinase 14 isoform X2 [Drosophila
>gi|1036796581|ref|XP_017136291.1| PREDICTED: cyclin-dependent kinase 14 isoform X1 [Drosophila
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>gi|1036796693|ref|XP_017136297.1| PREDICTED: cyclin-dependent kinase 14 isoform X5 [Drosophila
>gi|1036796676|ref|XP_017136296.1| PREDICTED: cyclin-dependent kinase 14 isoform X4 [Drosophila
>gi|1036843168|ref|XP_017138888.1| PREDICTED: 20-hydroxyecdysone protein [Drosophila miranda]
>gi|1036837087|ref|XP_017138547.1| PREDICTED: uncharacterized protein LOC108153227 isoform X2
>gi|1036837068|ref|XP_017138546.1| PREDICTED: uncharacterized protein LOC108153227 isoform X1
>gi|1036837104|ref|XP_017138548.1| PREDICTED: uncharacterized protein LOC108153227 isoform X3
>gi|1036837136|ref|XP_017138550.1| PREDICTED: uncharacterized protein LOC108153228 [Drosophila
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>gi|1036796159|ref|XP_017136269.1| PREDICTED: solute carrier family 35 member C2 [Drosophila m
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>gi|1036796215|ref|XP_017136273.1| PREDICTED: cyclin-J-like isoform X2 [Drosophila miranda]
>gi|1036796196|ref|XP_017136272.1| PREDICTED: cyclin-J-like isoform X1 [Drosophila miranda]
>gi|1036796272|ref|XP_017136276.1| PREDICTED: uncharacterized protein LOC108151887 [Drosophila
>gi|1036796253|ref|XP_017136275.1| PREDICTED: cyclin-J-like isoform X2 [Drosophila miranda]
>gi|1036796234|ref|XP_017136274.1| PREDICTED: cyclin-J-like isoform X1 [Drosophila miranda]
>gi|1036796290|ref|XP_017136277.1| PREDICTED: probable RNA helicase armi [Drosophila miranda]
>gi|1036820796|ref|XP_017137629.1| PREDICTED: eukaryotic translation initiation factor 5B [Dros
>gi|1036827452|ref|XP_017138004.1| PREDICTED: protein sprouty [Drosophila miranda]
>gi|1036827470|ref|XP_017138005.1| PREDICTED: protein sprouty [Drosophila miranda]
>gi|1036826626|ref|XP_017137955.1| PREDICTED: iduronate 2-sulfatase [Drosophila miranda]
>gi|1036826643|ref|XP_017137956.1| PREDICTED: brain protein I3 [Drosophila miranda]
>gi|1036826607|ref|XP_017137954.1| PREDICTED: cell division cycle protein 48 homolog [Drosophi
>gi|1036824580|ref|XP_017137843.1| PREDICTED: YTH domain-containing protein 1 [Drosophila mira
>gi|1036824597|ref|XP_017137844.1| PREDICTED: osteopetrosis-associated transmembrane protein 1
>gi|1036782975|ref|XP_017135520.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036782994|ref|XP_017135521.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036783031|ref|XP_017135523.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036783050|ref|XP_017135525.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036783012|ref|XP_017135522.1| PREDICTED: spectrin beta chain, non-erythrocytic 5 isoform
>gi|1036768461|ref|XP_017134683.1| PREDICTED: uncharacterized protein LOC108150885 [Drosophila
>gi|1036790307|ref|XP_017135939.1| PREDICTED: calcium uniporter protein, mitochondrial isoform
>gi|1036790326|ref|XP_017135940.1| PREDICTED: calcium uniporter protein, mitochondrial isoform
>gi|1036790345|ref|XP_017135941.1| PREDICTED: calcium uniporter protein, mitochondrial isoform
>gi|1036790288|ref|XP_017135938.1| PREDICTED: uncharacterized protein LOC108151696 [Drosophila
>gi|1036790364|ref|XP_017135942.1| PREDICTED: serine protease nudel [Drosophila miranda]
>gi|1036790382|ref|XP_017135943.1| PREDICTED: small RNA degrading nuclease 5 [Drosophila miran
>gi|1036790400|ref|XP_017135944.1| PREDICTED: innexin inx4 [Drosophila miranda]
>gi|1036845115|ref|XP_017138989.1| PREDICTED: uncharacterized protein LOC108153468 [Drosophila
>gi|1036851110|ref|XP_017139328.1| PREDICTED: glutamate receptor 1 [Drosophila miranda]
>gi|1036766014|ref|XP_017134552.1| PREDICTED: eukaryotic translation initiation factor 4E [Dros
>gi|1036761472|ref|XP_017156960.1| PREDICTED: uncharacterized protein CG32395 [Drosophila mira
>gi|1036841317|ref|XP_017138793.1| PREDICTED: uncharacterized protein LOC108153347 [Drosophila
>gi|1036841336|ref|XP_017138794.1| PREDICTED: uncharacterized protein LOC108153347 [Drosophila
>gi|1036850875|ref|XP_017139315.1| PREDICTED: ras-related and estrogen-regulated growth inhibit
>gi|1036847779|ref|XP_017139134.1| PREDICTED: tyrosine 3-monooxygenase [Drosophila miranda]
>gi|1036788370|ref|XP_017135826.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036788318|ref|XP_017135825.1| PREDICTED: serine/arginine repetitive matrix protein 2 isof
>gi|1036797211|ref|XP_017136328.1| PREDICTED: uncharacterized protein LOC108151921 isoform X1
>gi|1036797226|ref|XP_017136329.1| PREDICTED: uncharacterized protein LOC108151921 isoform X2
>gi|1036797245|ref|XP_017136330.1| PREDICTED: uncharacterized protein LOC108151921 isoform X3
>gi|1036797282|ref|XP_017136332.1| PREDICTED: ribosome maturation protein SBDS [Drosophila mira
>gi|1036797263|ref|XP_017136331.1| PREDICTED: LOW QUALITY PROTEIN: IST1 homolog [Drosophila mir
>gi|1036827359|ref|XP_017137998.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1036827321|ref|XP_017137996.1| PREDICTED: LOW QUALITY PROTEIN: POU domain protein CF1A [Dro
>gi|1036827340|ref|XP_017137997.1| PREDICTED: amidophosphoribosyltransferase [Drosophila miran
>gi|1036807133|ref|XP_017136884.1| PREDICTED: transcription factor SPT20 homolog isoform X3 [Dr
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>gi|1036807152|ref|XP_017136885.1| PREDICTED: zinc carboxypeptidase [Drosophila miranda]
>gi|1036807099|ref|XP_017136882.1| PREDICTED: transcription factor SPT20 homolog isoform X1 [Di
>gi|1036807118|ref|XP_017136883.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036807083|ref|XP_017136881.1| PREDICTED: ATP-dependent RNA helicase dbp2 [Drosophila miral
>gi|1036850635|ref|XP_017139301.1| PREDICTED: uncharacterized protein LOC108153688 [Drosophila
>gi|1036846639|ref|XP_017139069.1| PREDICTED: guanine nucleotide-binding protein G(f) subunit a
>gi|1036842128|ref|XP_017138830.1| PREDICTED: coatomer subunit zeta-1 [Drosophila miranda]
>gi|1036850128|ref|XP_017139272.1| PREDICTED: coiled-coil domain-containing protein 13 isoform
>gi|1036850146|ref|XP_017139273.1| PREDICTED: coiled-coil domain-containing protein 13 isoform
>gi|1036814831|ref|XP_017137302.1| PREDICTED: POU domain, class 3, transcription factor 2 [Dros
>gi|1036814810|ref|XP_017137301.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X4 [Droso
>gi|1036814792|ref|XP_017137300.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X3 [Droso
>gi|1036814773|ref|XP_017137299.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X2 [Droso
>gi|1036814754|ref|XP_017137298.1| PREDICTED: LIM and SH3 domain protein Lasp isoform X1 [Droso
>gi|1036810847|ref|XP_017137087.1| PREDICTED: protein disabled [Drosophila miranda]
>gi|1036810866|ref|XP_017137088.1| PREDICTED: uncharacterized protein LOC108152337 [Drosophila
>gi|1036768756|ref|XP_017134699.1| PREDICTED: cuticlin-2 [Drosophila miranda]
>gi|1036827652|ref|XP_017138016.1| PREDICTED: uncharacterized protein LOC108152888 [Drosophila
>gi|1036827633|ref|XP_017138015.1| PREDICTED: peptidoglycan-recognition protein SB2 [Drosophila
>gi|1036827614|ref|XP_017138014.1| PREDICTED: peptidoglycan-recognition protein SB1 [Drosophila
>gi|1036827595|ref|XP_017138013.1| PREDICTED: probable ATP-dependent RNA helicase Dbp73D [Dros
>gi|1036844534|ref|XP_017138964.1| PREDICTED: myrosinase 1 [Drosophila miranda]
>gi|1036789625|ref|XP_017135901.1| PREDICTED: neurotactin isoform X1 [Drosophila miranda]
>gi|1036789644|ref|XP_017135902.1| PREDICTED: neurotactin isoform X1 [Drosophila miranda]
>gi|1036789662|ref|XP_017135903.1| PREDICTED: neurotactin isoform X2 [Drosophila miranda]
>gi|1036789680|ref|XP_017135904.1| PREDICTED: E3 ubiquitin-protein ligase RFWD3 [Drosophila mi
>gi|1036789698|ref|XP_017135905.1| PREDICTED: cold shock domain-containing protein CG9705 [Dros
>gi|1036789716|ref|XP_017135906.1| PREDICTED: cold shock domain-containing protein CG9705 [Dros
>gi|1036789735|ref|XP_017135907.1| PREDICTED: cold shock domain-containing protein CG9705 [Dros
>gi|1036817427|ref|XP_017137448.1| PREDICTED: uncharacterized protein LOC108152547 [Drosophila
>gi|1036826463|ref|XP_017137946.1| PREDICTED: uncharacterized protein LOC108152846 isoform X1
>gi|1036826481|ref|XP_017137947.1| PREDICTED: uncharacterized protein LOC108152846 isoform X1
>gi|1036826552|ref|XP_017137951.1| PREDICTED: ephexin-1 isoform X3 [Drosophila miranda]
>gi|1036826570|ref|XP_017137952.1| PREDICTED: ephexin-1 isoform X4 [Drosophila miranda]
>gi|1036826497|ref|XP_017137948.1| PREDICTED: uncharacterized protein LOC108152846 isoform X2
>gi|1036826515|ref|XP_017137949.1| PREDICTED: uncharacterized protein LOC108152846 isoform X2
>gi|1036826534|ref|XP_017137950.1| PREDICTED: uncharacterized protein LOC108152846 isoform X2
>gi|1036807866|ref|XP_017136924.1| PREDICTED: 2-oxoglutarate dehydrogenase, mitochondrial isof
>gi|1036807788|ref|XP_017136920.1| PREDICTED: 2-oxoglutarate dehydrogenase-like, mitochondrial
>gi|1036807847|ref|XP_017136923.1| PREDICTED: 2-oxoglutarate dehydrogenase, mitochondrial isof
>gi|1036807807|ref|XP_017136921.1| PREDICTED: 2-oxoglutarate dehydrogenase-like, mitochondrial
>gi|1036807828|ref|XP_017136922.1| PREDICTED: 2-oxoglutarate dehydrogenase-like, mitochondrial
>gi|1036761493|ref|XP_017156962.1| PREDICTED: general odorant-binding protein 70 [Drosophila m
>gi|1036807885|ref|XP_017136925.1| PREDICTED: fibroin heavy chain isoform X1 [Drosophila miran
>gi|1036807907|ref|XP_017136926.1| PREDICTED: keratin, type I cytoskeletal 9 isoform X2 [Drosop
>gi|1036761512|ref|XP_017156963.1| PREDICTED: uncharacterized protein LOC108165435 [Drosophila
>gi|1036848014|ref|XP_017139148.1| PREDICTED: uncharacterized protein LOC108153578 isoform X2
>gi|1036848033|ref|XP_017139149.1| PREDICTED: uncharacterized protein LOC108153578 isoform X3
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>gi|1036847997|ref|XP_017139147.1| PREDICTED: uncharacterized protein LOC108153578 isoform X1
>gi|1036764820|ref|XP_017157138.1| PREDICTED: protein app1 [Drosophila miranda]
>gi|1036767699|ref|XP_017134641.1| PREDICTED: uncharacterized protein LOC108150848 [Drosophila
>gi|1036804065|ref|XP_017136714.1| PREDICTED: prickle-like protein 3 isoform X5 [Drosophila mi
>gi|1036804046|ref|XP_017136713.1| PREDICTED: four and a half LIM domains protein 2 isoform X4
>gi|1036804083|ref|XP_017136715.1| PREDICTED: prickle-like protein 3 isoform X6 [Drosophila mi
>gi|1036804011|ref|XP_017136710.1| PREDICTED: trichohyalin isoform X2 [Drosophila miranda]
>gi|1036804105|ref|XP_017136716.1| PREDICTED: four and a half LIM domains protein 2 isoform X7
>gi|1036804122|ref|XP_017136717.1| PREDICTED: four and a half LIM domains protein 2 isoform X8
>gi|1036804029|ref|XP_017136711.1| PREDICTED: trichohyalin isoform X3 [Drosophila miranda]
>gi|1036803992|ref|XP_017136709.1| PREDICTED: trichohyalin isoform X1 [Drosophila miranda]
>gi|1036804141|ref|XP_017136718.1| PREDICTED: peptidyl-prolyl cis-trans isomerase CYP95 [Drosop
>gi|1036840242|ref|XP_017138731.1| PREDICTED: coiled-coil domain-containing protein 22 homolog
>gi|1036787262|ref|XP_017135764.1| PREDICTED: opsin Rh4 [Drosophila miranda]
>gi|1036787297|ref|XP_017135765.1| PREDICTED: probable E3 ubiquitin-protein ligase sinah [Dros
>gi|1036787318|ref|XP_017135767.1| PREDICTED: E3 ubiquitin-protein ligase sina [Drosophila mir
>gi|1036787337|ref|XP_017135768.1| PREDICTED: E3 ubiquitin-protein ligase sina [Drosophila mir
>gi|1036761531|ref|XP_017156964.1| PREDICTED: uncharacterized protein LOC108165436 [Drosophila
>gi|1036787353|ref|XP_017135769.1| PREDICTED: uncharacterized protein LOC108151600 [Drosophila
>gi|1036802830|ref|XP_017136643.1| PREDICTED: transmembrane protein 258 homolog [Drosophila mi
>gi|1036802792|ref|XP_017136641.1| PREDICTED: tumor susceptibility gene 101 protein isoform X1
>gi|1036802811|ref|XP_017136642.1| PREDICTED: tumor susceptibility gene 101 protein isoform X2
>gi|1036802866|ref|XP_017136646.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 6
>gi|1036802848|ref|XP_017136645.1| PREDICTED: tyrosine-protein phosphatase non-receptor type 6
>gi|1036838660|ref|XP_017138646.1| PREDICTED: uncharacterized protein LOC108153285 [Drosophila
>gi|1036838641|ref|XP_017138645.1| PREDICTED: rhythmically expressed gene 2 protein [Drosophila
>gi|1036800197|ref|XP_017136496.1| PREDICTED: UDP-glucose 4-epimerase [Drosophila miranda]
>gi|1036800216|ref|XP_017136497.1| PREDICTED: UDP-glucose 4-epimerase [Drosophila miranda]
>gi|1036800253|ref|XP_017136499.1| PREDICTED: uncharacterized protein C45G9.7 [Drosophila mira
>gi|1036800235|ref|XP_017136498.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036800178|ref|XP_017136495.1| PREDICTED: DNA repair protein REV1 [Drosophila miranda]
>gi|1036834067|ref|XP_017138372.1| PREDICTED: uncharacterized protein LOC108153099 [Drosophila
>gi|1036842235|ref|XP_017138837.1| PREDICTED: uncharacterized protein LOC108153377 [Drosophila
>gi|1036842254|ref|XP_017138838.1| PREDICTED: uncharacterized protein LOC108153377 [Drosophila
>gi|1036836483|ref|XP_017138512.1| PREDICTED: uncharacterized protein LOC108153202 [Drosophila
>gi|1036809010|ref|XP_017136986.1| PREDICTED: retinoid-inducible serine carboxypeptidase [Dros
>gi|1036809028|ref|XP_017136987.1| PREDICTED: retinoid-inducible serine carboxypeptidase [Dros
>gi|1036809064|ref|XP_017136989.1| PREDICTED: ras-related protein Rab-14 [Drosophila miranda]
>gi|1036809046|ref|XP_017136988.1| PREDICTED: vesicle transport through interaction with t-SNA
>gi|1036808993|ref|XP_017136985.1| PREDICTED: LOW QUALITY PROTEIN: mediator of RNA polymerase
>gi|1036832798|ref|XP_017138308.1| PREDICTED: uncharacterized protein LOC108153047 isoform X2
>gi|1036832720|ref|XP_017138304.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036832741|ref|XP_017138305.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036832760|ref|XP_017138306.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036832779|ref|XP_017138307.1| PREDICTED: uncharacterized protein LOC108153047 isoform X1
>gi|1036851636|ref|XP_017139360.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036851656|ref|XP_017139361.1| PREDICTED: cuticle protein 38-like [Drosophila miranda]
>gi|1036851420|ref|XP_017139347.1| PREDICTED: cuticle protein 38-like [Drosophila miranda]
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>gi|1036851402|ref|XP_017139345.1| PREDICTED: cuticle protein 16.5-like [Drosophila miranda]
>gi|1036816806|ref|XP_017137415.1| PREDICTED: histone-lysine N-methyltransferase 2D [Drosophila
>gi|1036816825|ref|XP_017137416.1| PREDICTED: protein extra-macrochaetae [Drosophila miranda]
>gi|1036798180|ref|XP_017136382.1| PREDICTED: splicing factor 3B subunit 3 isoform X1 [Drosoph
>gi|1036798199|ref|XP_017136383.1| PREDICTED: splicing factor 3B subunit 3 isoform X2 [Drosoph
>gi|1036798218|ref|XP_017136384.1| PREDICTED: splicing factor 3B subunit 3 isoform X3 [Drosoph
>gi|1036798253|ref|XP_017136386.1| PREDICTED: uncharacterized protein LOC108151958 [Drosophila
>gi|1036798235|ref|XP_017136385.1| PREDICTED: kidney mitochondrial carrier protein 1 [Drosophil
>gi|1036765762|ref|XP_017134538.1| PREDICTED: vacuole membrane protein 1 [Drosophila miranda]
>gi|1036826695|ref|XP_017137960.1| PREDICTED: uncharacterized protein LOC108152853 [Drosophila
>gi|1036826711|ref|XP_017137961.1| PREDICTED: uncharacterized protein LOC108152853 [Drosophila
>gi|1036826729|ref|XP_017137962.1| PREDICTED: ubiquitin-conjugating enzyme E2-22 kDa-like [Dros
>gi|1036826660|ref|XP_017137957.1| PREDICTED: gustatory receptor 68a [Drosophila miranda]
>gi|1036826677|ref|XP_017137959.1| PREDICTED: gustatory receptor 68a [Drosophila miranda]
>gi|1036761550|ref|XP_017156965.1| PREDICTED: LOW QUALITY PROTEIN: 2-oxoglutarate dehydrogenase
>gi|1036767830|ref|XP_017134648.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1036807170|ref|XP_017136886.1| PREDICTED: phosphatidylinositol 4-phosphate 3-kinase C2 doma
>gi|1036807187|ref|XP_017136887.1| PREDICTED: phosphatidylinositol 4-phosphate 3-kinase C2 doma
>gi|1036807206|ref|XP_017136888.1| PREDICTED: phosphatidylinositol 4-phosphate 3-kinase C2 doma
>gi|1036807244|ref|XP_017136891.1| PREDICTED: uncharacterized protein LOC108152221 [Drosophila
>gi|1036807225|ref|XP_017136889.1| PREDICTED: uncharacterized protein LOC108152220 isoform X2
>gi|1036782214|ref|XP_017135475.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1036782252|ref|XP_017135478.1| PREDICTED: peptidyl-prolyl cis-trans isomerase CWC27 homolog
>gi|1036782232|ref|XP_017135476.1| PREDICTED: kinesin-like protein Klp68D [Drosophila miranda]
>gi|1036782288|ref|XP_017135480.1| PREDICTED: 2-amino-3-ketobutyrate coenzyme A ligase, mitocho
>gi|1036782175|ref|XP_017135473.1| PREDICTED: cubilin homolog isoform X1 [Drosophila miranda]
>gi|1036782196|ref|XP_017135474.1| PREDICTED: cubilin homolog isoform X2 [Drosophila miranda]
>gi|1036782305|ref|XP_017135481.1| PREDICTED: uncharacterized protein LOC108151404 [Drosophila
>gi|1036782270|ref|XP_017135479.1| PREDICTED: uncharacterized protein LOC108151402 [Drosophila
>gi|1036782324|ref|XP_017135482.1| PREDICTED: 60S ribosomal protein L10a-2 [Drosophila miranda]
>gi|1036793404|ref|XP_017136111.1| PREDICTED: uncharacterized protein LOC108151779 [Drosophila
>gi|1036793465|ref|XP_017136115.1| PREDICTED: NADH-cytochrome b5 reductase 3 isoform X1 [Drosop
>gi|1036793484|ref|XP_017136116.1| PREDICTED: NADH-cytochrome b5 reductase 3 isoform X2 [Drosop
>gi|1036793423|ref|XP_017136112.1| PREDICTED: G2/mitotic-specific cyclin-A [Drosophila miranda]
>gi|1036793446|ref|XP_017136114.1| PREDICTED: RIB43A-like with coiled-coils protein 2 [Drosophi
>gi|1036843187|ref|XP_017138889.1| PREDICTED: alpha-ketoglutarate-dependent dioxygenase alkB ho
>gi|1036843262|ref|XP_017138893.1| PREDICTED: BPTI/Kunitz domain-containing protein-like [Droso
>gi|1036843223|ref|XP_017138891.1| PREDICTED: uncharacterized protein LOC108153408 isoform X1
>gi|1036843241|ref|XP_017138892.1| PREDICTED: BPTI/Kunitz domain-containing protein-like isofor
>gi|1036843206|ref|XP_017138890.1| PREDICTED: kunitz-type serine protease inhibitor NACI [Droso
>gi|1036761570|ref|XP_017156966.1| PREDICTED: palmitoleoyl-protein carboxylesterase NOTUM [Dros
>gi|1036808506|ref|XP_017136961.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808524|ref|XP_017136962.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808542|ref|XP_017136963.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808560|ref|XP_017136964.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808579|ref|XP_017136965.1| PREDICTED: uncharacterized protein LOC108152263 isoform X1
>gi|1036808618|ref|XP_017136967.1| PREDICTED: uncharacterized protein LOC108152263 isoform X3
>gi|1036808599|ref|XP_017136966.1| PREDICTED: probable RNA-binding protein orb2 isoform X2 [Dro
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>gi|1036808636|ref|XP_017136968.1| PREDICTED: gram-negative bacteria-binding protein 3 [Drosop]
>gi|1036793983|ref|XP_017136142.1| PREDICTED: 50S ribosomal protein L7/L12 [Drosophila miranda]
>gi|1036793945|ref|XP_017136140.1| PREDICTED: uncharacterized protein LOC108151800 [Drosophila
>gi|1036793925|ref|XP_017136139.1| PREDICTED: valine--tRNA ligase, mitochondrial [Drosophila m:
>gi|1036793887|ref|XP_017136137.1| PREDICTED: coatomer subunit alpha [Drosophila miranda]
>gi|1036793906|ref|XP_017136138.1| PREDICTED: coatomer subunit alpha [Drosophila miranda]
>gi|1036793964|ref|XP_017136141.1| PREDICTED: protein big brother [Drosophila miranda]
>gi|1036831812|ref|XP_017138253.1| PREDICTED: methyltransferase-like protein isoform X2 [Drosop]
>gi|1036831793|ref|XP_017138252.1| PREDICTED: methyltransferase-like protein isoform X1 [Drosop]
>gi|1036831846|ref|XP_017138256.1| PREDICTED: vesicle-associated membrane protein 2 isoform X2
>gi|1036831865|ref|XP_017138257.1| PREDICTED: vesicle-associated membrane protein 2 isoform X3
>gi|1036831829|ref|XP_017138255.1| PREDICTED: vesicle-associated membrane protein 2 isoform X1
>gi|1036831901|ref|XP_017138258.1| PREDICTED: vesicle-associated membrane protein 2 isoform X4
>gi|1036793173|ref|XP_017136101.1| PREDICTED: uncharacterized protein LOC108151768 [Drosophila
>gi|1036793062|ref|XP_017136095.1| PREDICTED: vitamin K-dependent gamma-carboxylase isoform X1
>gi|1036793079|ref|XP_017136096.1| PREDICTED: vitamin K-dependent gamma-carboxylase isoform X2
>gi|1036793043|ref|XP_017136094.1| PREDICTED: ATP-binding cassette sub-family B member 7, mitc
>gi|1036793025|ref|XP_017136093.1| PREDICTED: ATP-binding cassette sub-family B member 7, mitc
>gi|1036793154|ref|XP_017136100.1| PREDICTED: SET domain-containing protein 4 [Drosophila mira]
>gi|1036793191|ref|XP_017136102.1| PREDICTED: protein OPI10 homolog [Drosophila miranda]
>gi|1036793006|ref|XP_017136092.1| PREDICTED: 2-oxoglutarate dehydrogenase, mitochondrial [Dros
>gi|1036792988|ref|XP_017136091.1| PREDICTED: uncharacterized protein LOC108151760 [Drosophila
>gi|1036793118|ref|XP_017136098.1| PREDICTED: uncharacterized protein LOC108151765 isoform X2
>gi|1036793099|ref|XP_017136097.1| PREDICTED: uncharacterized protein LOC108151765 isoform X1
>gi|1036793135|ref|XP_017136099.1| PREDICTED: uncharacterized protein LOC108151765 isoform X3
>gi|1036795601|ref|XP_017136237.1| PREDICTED: TOX high mobility group box family member 4-like
>gi|1036761589|ref|XP_017156967.1| PREDICTED: TOX high mobility group box family member 4-like
>gi|1036795513|ref|XP_017136232.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795531|ref|XP_017136233.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795550|ref|XP_017136234.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795566|ref|XP_017136235.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036795584|ref|XP_017136236.1| PREDICTED: uncharacterized protein LOC108151859 [Drosophila
>gi|1036851274|ref|XP_017139338.1| PREDICTED: uncharacterized protein LOC108153721 isoform X1
>gi|1036851292|ref|XP_017139339.1| PREDICTED: ninjurin-2 isoform X2 [Drosophila miranda]
>gi|1036837030|ref|XP_017138544.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 alpha subcomple
>gi|1036837049|ref|XP_017138545.1| PREDICTED: uncharacterized protein LOC108153226 [Drosophila
>gi|1036837011|ref|XP_017138543.1| PREDICTED: guanylate kinase [Drosophila miranda]
>gi|1036761608|ref|XP_017156968.1| PREDICTED: LOW QUALITY PROTEIN: DNA polymerase delta small s
>gi|1036842162|ref|XP_017138832.1| PREDICTED: WD repeat-containing protein 35 isoform X2 [Drosc
>gi|1036842144|ref|XP_017138831.1| PREDICTED: WD repeat-containing protein 35 isoform X1 [Drosc
>gi|1036851364|ref|XP_017139343.1| PREDICTED: uncharacterized protein At4g17910 isoform X1 [Dro
>gi|1036851383|ref|XP_017139344.1| PREDICTED: uncharacterized protein At4g17910 isoform X2 [Dro
>gi|1036835025|ref|XP_017138425.1| PREDICTED: glycerol kinase [Drosophila miranda]
>gi|1036835043|ref|XP_017138427.1| PREDICTED: glycerol kinase [Drosophila miranda]
>gi|1036817538|ref|XP_017137455.1| PREDICTED: uncharacterized protein LOC108152552 isoform X1
>gi|1036817559|ref|XP_017137456.1| PREDICTED: uncharacterized protein LOC108152552 isoform X2
>gi|1036817578|ref|XP_017137457.1| PREDICTED: uncharacterized protein LOC108152552 isoform X3
>gi|1036817597|ref|XP_017137458.1| PREDICTED: uncharacterized protein LOC108152552 isoform X4
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>gi|1036817615|ref|XP_017137459.1| PREDICTED: probable chitinase 2 isoform X1 [Drosophila miran
>gi|1036817635|ref|XP_017137460.1| PREDICTED: probable chitinase 2 isoform X2 [Drosophila miran
>gi|1036817655|ref|XP_017137462.1| PREDICTED: probable chitinase 2 isoform X2 [Drosophila miran
>gi|1036813090|ref|XP_017137202.1| PREDICTED: membrane-associated protein Hem [Drosophila miran
>gi|1036813184|ref|XP_017137207.1| PREDICTED: SRSF protein kinase 3 isoform X5 [Drosophila mira
>gi|1036813145|ref|XP_017137205.1| PREDICTED: SRSF protein kinase 1 isoform X3 [Drosophila mira
>gi|1036813166|ref|XP_017137206.1| PREDICTED: SRSF protein kinase 3 isoform X4 [Drosophila mira
>gi|1036813126|ref|XP_017137204.1| PREDICTED: SRSF protein kinase 1 isoform X2 [Drosophila mira
>gi|1036813108|ref|XP_017137203.1| PREDICTED: SRSF protein kinase 1 isoform X1 [Drosophila mira
>gi|1036823186|ref|XP_017137764.1| PREDICTED: dnaJ homolog subfamily C member 5 homolog isoform
>gi|1036823168|ref|XP_017137763.1| PREDICTED: dnaJ homolog subfamily C member 5 homolog isoform
>gi|1036837740|ref|XP_017138588.1| PREDICTED: pancreatic lipase-related protein 3 [Drosophila m
>gi|1036837646|ref|XP_017138582.1| PREDICTED: speract receptor isoform X1 [Drosophila miranda]
>gi|1036837683|ref|XP_017138584.1| PREDICTED: speract receptor isoform X3 [Drosophila miranda]
>gi|1036837664|ref|XP_017138583.1| PREDICTED: speract receptor isoform X2 [Drosophila miranda]
>gi|1036837702|ref|XP_017138585.1| PREDICTED: speract receptor isoform X4 [Drosophila miranda]
>gi|1036837721|ref|XP_017138587.1| PREDICTED: dual specificity protein phosphatase 22-like [Dro
>gi|1036845743|ref|XP_017139018.1| PREDICTED: uncharacterized protein LOC108153488 isoform X2
>gi|1036845724|ref|XP_017139017.1| PREDICTED: uncharacterized protein LOC108153488 isoform X1
>gi|1036818834|ref|XP_017137530.1| PREDICTED: BAG domain-containing protein Samui isoform X1 [
>gi|1036818873|ref|XP_017137532.1| PREDICTED: BAG domain-containing protein Samui isoform X3 [
>gi|1036818854|ref|XP_017137531.1| PREDICTED: BAG domain-containing protein Samui isoform X2 [
>gi|1036818892|ref|XP_017137533.1| PREDICTED: BAG domain-containing protein Samui isoform X4 [
>gi|1036818913|ref|XP_017137534.1| PREDICTED: BAG domain-containing protein Samui isoform X5 [
>gi|1036818931|ref|XP_017137535.1| PREDICTED: BAG domain-containing protein Samui isoform X5 [
>gi|1036818951|ref|XP_017137536.1| PREDICTED: drebrin-like protein [Drosophila miranda]
>gi|1036785030|ref|XP_017135640.1| PREDICTED: protein suppressor of hairy wing [Drosophila mira
>gi|1036785012|ref|XP_017135639.1| PREDICTED: protein-cysteine N-palmitoyltransferase Rasp [Dro
>gi|1036784993|ref|XP_017135638.1| PREDICTED: uncharacterized protein LOC108151504 [Drosophila
>gi|1036785049|ref|XP_017135641.1| PREDICTED: NADH dehydrogenase [ubiquinone] iron-sulfur prote
>gi|1036784938|ref|XP_017135634.1| PREDICTED: histone-lysine N-methyltransferase 2D isoform X1
>gi|1036784974|ref|XP_017135636.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036784956|ref|XP_017135635.1| PREDICTED: longitudinals lacking protein, isoforms A/B/D/L
>gi|1036792881|ref|XP_017136085.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 5 [Drosophi
>gi|1036792899|ref|XP_017136086.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 5 [Drosophi
>gi|1036792917|ref|XP_017136087.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 5 [Drosophi
>gi|1036792971|ref|XP_017136090.1| PREDICTED: alpha-1,3/1,6-mannosyltransferase ALG2 [Drosophil
>gi|1036792953|ref|XP_017136089.1| PREDICTED: hippocampus abundant transcript 1 protein isoform
>gi|1036792934|ref|XP_017136088.1| PREDICTED: hippocampus abundant transcript 1 protein isoform
>gi|1036847642|ref|XP_017139126.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1036805614|ref|XP_017136802.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036805633|ref|XP_017136803.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036805652|ref|XP_017136804.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036805689|ref|XP_017136806.1| PREDICTED: prominin-like protein isoform X3 [Drosophila mira
>gi|1036805670|ref|XP_017136805.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1036805708|ref|XP_017136807.1| PREDICTED: prominin-like protein isoform X4 [Drosophila mira
>gi|1036805727|ref|XP_017136809.1| PREDICTED: prominin-like protein isoform X5 [Drosophila mira
>gi|1036845685|ref|XP_017139015.1| PREDICTED: probable chitinase 3 [Drosophila miranda]
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>gi|1036840034|ref|XP_017138719.1| PREDICTED: prominin-like protein isoform X1 [Drosophila mira
>gi|1036840051|ref|XP_017138720.1| PREDICTED: prominin-like protein isoform X2 [Drosophila mira
>gi|1036811818|ref|XP_017137139.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036811799|ref|XP_017137138.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036811780|ref|XP_017137137.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036811837|ref|XP_017137140.1| PREDICTED: PAB-dependent poly(A)-specific ribonuclease subu
>gi|1036780403|ref|XP_017135371.1| PREDICTED: cysteine and histidine-rich protein 1 homolog [D
>gi|1036780442|ref|XP_017135373.1| PREDICTED: protein FAM207A [Drosophila miranda]
>gi|1036780365|ref|XP_017135368.1| PREDICTED: cyclin-dependent kinase 8 [Drosophila miranda]
>gi|1036780330|ref|XP_017135367.1| PREDICTED: putative glycerol kinase 5 [Drosophila miranda]
>gi|1036780422|ref|XP_017135372.1| PREDICTED: CRAL-TRIO domain-containing protein C3H8.02 [Dros
>gi|1036780296|ref|XP_017135365.1| PREDICTED: sphingosine kinase 2 [Drosophila miranda]
>gi|1036780311|ref|XP_017135366.1| PREDICTED: dnaJ homolog subfamily C member 2 [Drosophila mi
>gi|1036780384|ref|XP_017135369.1| PREDICTED: islet cell autoantigen 1 [Drosophila miranda]
>gi|1036823112|ref|XP_017137759.1| PREDICTED: uncharacterized protein LOC108152730 [Drosophila
>gi|1036823129|ref|XP_017137760.1| PREDICTED: uncharacterized protein LOC108152730 [Drosophila
>gi|1036823147|ref|XP_017137761.1| PREDICTED: uncharacterized protein LOC108152731 [Drosophila
>gi|1036822457|ref|XP_017137721.1| PREDICTED: dnaJ protein homolog 1-like isoform X1 [Drosophil
>gi|1036822476|ref|XP_017137722.1| PREDICTED: dnaJ protein homolog 1-like isoform X2 [Drosophil
>gi|1036822497|ref|XP_017137724.1| PREDICTED: LOW QUALITY PROTEIN: dnaJ protein homolog 1-like
>gi|1036761627|ref|XP_017156969.1| PREDICTED: enhancer of rudimentary homolog [Drosophila mira
>gi|1036845556|ref|XP_017139007.1| PREDICTED: uncharacterized protein LOC108153479 [Drosophila
>gi|1036845575|ref|XP_017139008.1| PREDICTED: uncharacterized protein LOC108153479 [Drosophila
>gi|1036764858|ref|XP_017157140.1| PREDICTED: phosphoglycolate phosphatase 2 [Drosophila miran
>gi|1036848219|ref|XP_017139158.1| PREDICTED: glycerol-3-phosphate phosphatase [Drosophila mira
>gi|1036848200|ref|XP_017139157.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1036850267|ref|XP_017139279.1| PREDICTED: FMRFamide receptor [Drosophila miranda]
>gi|1036822696|ref|XP_017137735.1| PREDICTED: carnitine O-palmitoyltransferase 2, mitochondria
>gi|1036822773|ref|XP_017137739.1| PREDICTED: odorant receptor 63a [Drosophila miranda]
>gi|1036822538|ref|XP_017137726.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822519|ref|XP_017137725.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822673|ref|XP_017137734.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822654|ref|XP_017137732.1| PREDICTED: uncharacterized protein LOC108152711 isoform X8
>gi|1036822635|ref|XP_017137731.1| PREDICTED: uncharacterized protein LOC108152711 isoform X7
>gi|1036822595|ref|XP_017137729.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822614|ref|XP_017137730.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822576|ref|XP_017137728.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822557|ref|XP_017137727.1| PREDICTED: potassium voltage-gated channel protein Shab iso
>gi|1036822715|ref|XP_017137736.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036761646|ref|XP_017156970.1| PREDICTED: uncharacterized protein LOC108165443 [Drosophila
>gi|1036822734|ref|XP_017137737.1| PREDICTED: uncharacterized protein LOC108152714 [Drosophila
>gi|1036822754|ref|XP_017137738.1| PREDICTED: uncharacterized protein LOC108152714 [Drosophila
>gi|1036786583|ref|XP_017135729.1| PREDICTED: LOW QUALITY PROTEIN: DNA N6-methyl adenine demet
>gi|1036818382|ref|XP_017137503.1| PREDICTED: N-acetylgalactosaminyltransferase 6 [Drosophila
>gi|1036818400|ref|XP_017137504.1| PREDICTED: 28S ribosomal protein S35, mitochondrial [Drosop
>gi|1036809104|ref|XP_017136992.1| PREDICTED: pyridoxal kinase [Drosophila miranda]
>gi|1036809143|ref|XP_017136994.1| PREDICTED: uncharacterized protein LOC108152287 [Drosophila
>gi|1036809083|ref|XP_017136990.1| PREDICTED: kinesin-like protein KIF18A [Drosophila miranda]
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>gi|1036809123|ref|XP_017136993.1| PREDICTED: ribosome-recycling factor, mitochondrial [Drosoph
>gi|1036831095|ref|XP_017138214.1| PREDICTED: protein FAM76A isoform X1 [Drosophila miranda]
>gi|1036831115|ref|XP_017138215.1| PREDICTED: uncharacterized protein LOC108152991 isoform X2
>gi|1036831151|ref|XP_017138217.1| PREDICTED: adrenodoxin-like protein, mitochondrial [Drosoph
>gi|1036831133|ref|XP_017138216.1| PREDICTED: heat shock protein 67B3 [Drosophila miranda]
>gi|1036845134|ref|XP_017138990.1| PREDICTED: heat shock protein 67B2 [Drosophila miranda]
>gi|1036845153|ref|XP_017138991.1| PREDICTED: heat shock protein 22 [Drosophila miranda]
>gi|1036832955|ref|XP_017138317.1| PREDICTED: heat shock protein 27 [Drosophila miranda]
>gi|1036832938|ref|XP_017138316.1| PREDICTED: heat shock protein 26 [Drosophila miranda]
>gi|1036832919|ref|XP_017138314.1| PREDICTED: uncharacterized protein LOC108153054 [Drosophila
>gi|1036845613|ref|XP_017139010.1| PREDICTED: heat shock protein 23 [Drosophila miranda]
>gi|1036816787|ref|XP_017137414.1| PREDICTED: LOW QUALITY PROTEIN: heat shock protein 27 [Dros
>gi|1036816764|ref|XP_017137413.1| PREDICTED: uncharacterized protein LOC108152523 [Drosophila
>gi|1036831436|ref|XP_017138233.1| PREDICTED: 40S ribosomal protein S17 [Drosophila miranda]
>gi|1036831417|ref|XP_017138232.1| PREDICTED: SHC-transforming protein 2 [Drosophila miranda]
>gi|1036798693|ref|XP_017136412.1| PREDICTED: phosphoserine phosphatase [Drosophila miranda]
>gi|1036798565|ref|XP_017136405.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [
>gi|1036798600|ref|XP_017136406.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [
>gi|1036798619|ref|XP_017136407.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [
>gi|1036798637|ref|XP_017136408.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X1 [
>gi|1036798655|ref|XP_017136409.1| PREDICTED: alpha-tubulin N-acetyltransferase 1 isoform X2 [
>gi|1036798674|ref|XP_017136410.1| PREDICTED: ataxin-2 homolog [Drosophila miranda]
>gi|1036798711|ref|XP_017136413.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036798730|ref|XP_017136414.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036818252|ref|XP_017137495.1| PREDICTED: uncharacterized protein LOC108152569 [Drosophila
>gi|1036818271|ref|XP_017137497.1| PREDICTED: uncharacterized protein LOC108152570 [Drosophila
>gi|1036764038|ref|XP_017157101.1| PREDICTED: sentrin-specific protease 1-like [Drosophila mir
>gi|1036761663|ref|XP_017156971.1| PREDICTED: NADH-quinone oxidoreductase subunit C-like, part
>gi|1036790531|ref|XP_017135951.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X
>gi|1036790550|ref|XP_017135952.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X
>gi|1036790569|ref|XP_017135953.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X2
>gi|1036790640|ref|XP_017135957.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X6
>gi|1036790657|ref|XP_017135958.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X7
>gi|1036790676|ref|XP_017135959.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036790588|ref|XP_017135954.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X3
>gi|1036790769|ref|XP_017135964.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X
>gi|1036790695|ref|XP_017135960.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036790751|ref|XP_017135963.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X
>gi|1036790604|ref|XP_017135955.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X4
>gi|1036790622|ref|XP_017135956.1| PREDICTED: serine/threonine-protein kinase mig-15 isoform X5
>gi|1036790714|ref|XP_017135961.1| PREDICTED: mitogen-activated protein kinase kinase kinase ki
>gi|1036790733|ref|XP_017135962.1| PREDICTED: misshapen-like kinase 1 isoform X11 [Drosophila m
>gi|1036841577|ref|XP_017138808.1| PREDICTED: 60S ribosomal protein L8 [Drosophila miranda]
>gi|1036794021|ref|XP_017136147.1| PREDICTED: protein daughter of sevenless [Drosophila miranda
>gi|1036794054|ref|XP_017136149.1| PREDICTED: enkurin [Drosophila miranda]
>gi|1036794038|ref|XP_017136148.1| PREDICTED: uncharacterized protein LOC108151806 [Drosophila
>gi|1036794088|ref|XP_017136151.1| PREDICTED: acyl-coenzyme A thioesterase 13-like [Drosophila
>gi|1036794071|ref|XP_017136150.1| PREDICTED: acyl-coenzyme A thioesterase 13-like [Drosophila
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>gi|1036794105|ref|XP_017136152.1| PREDICTED: acyl-coenzyme A thioesterase 13 [Drosophila miran
>gi|1036794123|ref|XP_017136153.1| PREDICTED: acyl-coenzyme A thioesterase 13 [Drosophila miran
>gi|1036794002|ref|XP_017136143.1| PREDICTED: peroxidasin [Drosophila miranda]
>gi|1036804921|ref|XP_017136763.1| PREDICTED: uncharacterized protein KIAA1143 homolog [Drosoph
>gi|1036804845|ref|XP_017136758.1| PREDICTED: uncharacterized protein LOC108152153 isoform X1
>gi|1036804864|ref|XP_017136759.1| PREDICTED: uncharacterized protein LOC108152153 isoform X1
>gi|1036804883|ref|XP_017136760.1| PREDICTED: uncharacterized protein LOC108152153 isoform X2
>gi|1036804902|ref|XP_017136762.1| PREDICTED: uncharacterized protein LOC108152154 [Drosophila
>gi|1036851963|ref|XP_017139382.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1036840015|ref|XP_017138718.1| PREDICTED: uncharacterized protein LOC108153317 [Drosophila
>gi|1036839981|ref|XP_017138715.1| PREDICTED: probable proline--tRNA ligase, mitochondrial iso
>gi|1036840000|ref|XP_017138716.1| PREDICTED: probable proline--tRNA ligase, mitochondrial iso
>gi|1036831265|ref|XP_017138224.1| PREDICTED: snurportin-1 [Drosophila miranda]
>gi|1036831244|ref|XP_017138223.1| PREDICTED: uncharacterized protein LOC108152996 [Drosophila
>gi|1036831284|ref|XP_017138225.1| PREDICTED: uncharacterized protein LOC108152998 [Drosophila
>gi|1036833786|ref|XP_017138354.1| PREDICTED: signal recognition particle 54 kDa protein [Dros
>gi|1036772751|ref|XP_017134926.1| PREDICTED: flap endonuclease GEN [Drosophila miranda]
>gi|1036772677|ref|XP_017134922.1| PREDICTED: alanine--tRNA ligase, mitochondrial [Drosophila m
>gi|1036772789|ref|XP_017134928.1| PREDICTED: FIT family protein CG10671 [Drosophila miranda]
>gi|1036772849|ref|XP_017134930.1| PREDICTED: ganglioside-induced differentiation-associated pr
>gi|1036772868|ref|XP_017134932.1| PREDICTED: ganglioside-induced differentiation-associated pr
>gi|1036772828|ref|XP_017134929.1| PREDICTED: ganglioside-induced differentiation-associated pr
>gi|1036772887|ref|XP_017134933.1| PREDICTED: dehydrogenase/reductase SDR family member 4 [Dros
>gi|1036772937|ref|XP_017134936.1| PREDICTED: charged multivesicular body protein 2b-B [Drosoph
>gi|1036772920|ref|XP_017134935.1| PREDICTED: TP53-regulating kinase [Drosophila miranda]
>gi|1036772770|ref|XP_017134927.1| PREDICTED: pentatricopeptide repeat-containing protein 1, mi
>gi|1036772954|ref|XP_017134937.1| PREDICTED: protein Asterix [Drosophila miranda]
>gi|1036772903|ref|XP_017134934.1| PREDICTED: ubiquitin thioesterase OTU1 [Drosophila miranda]
>gi|1036773045|ref|XP_017134941.1| PREDICTED: uncharacterized protein LOC108151073 [Drosophila
>gi|1036773020|ref|XP_017134940.1| PREDICTED: uncharacterized protein LOC108151072 [Drosophila
>gi|1036761681|ref|XP_017156972.1| PREDICTED: uncharacterized protein LOC108165445 [Drosophila
>gi|1036761700|ref|XP_017156973.1| PREDICTED: LOW QUALITY PROTEIN: myb-like protein AA [Drosoph
>gi|1036772969|ref|XP_017134938.1| PREDICTED: uncharacterized protein LOC108151070 [Drosophila
>gi|1036772986|ref|XP_017134939.1| PREDICTED: uncharacterized protein LOC108151070 [Drosophila
>gi|1036772696|ref|XP_017134923.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036772715|ref|XP_017134924.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1036772733|ref|XP_017134925.1| PREDICTED: proteoglycan 4 isoform X2 [Drosophila miranda]
>gi|1036772658|ref|XP_017134921.1| PREDICTED: helicase POLQ-like [Drosophila miranda]
>gi|1036792714|ref|XP_017136077.1| PREDICTED: protein IWS1 homolog A-like [Drosophila miranda]
>gi|1036792700|ref|XP_017136075.1| PREDICTED: uncharacterized ATP-dependent helicase C29A10.10
>gi|1036792733|ref|XP_017136078.1| PREDICTED: phosphatidate cytidylyltransferase, photorecepto
>gi|1036827995|ref|XP_017138035.1| PREDICTED: SUMO-activating enzyme subunit 2 [Drosophila mira
>gi|1036799198|ref|XP_017136441.1| PREDICTED: uncharacterized protein LOC108151987 [Drosophila
>gi|1036799179|ref|XP_017136440.1| PREDICTED: 2-aminoethanethiol dioxygenase [Drosophila mirand
>gi|1036799161|ref|XP_017136439.1| PREDICTED: lariat debranching enzyme [Drosophila miranda]
>gi|1036799143|ref|XP_017136438.1| PREDICTED: importin-7 [Drosophila miranda]
>gi|1036812528|ref|XP_017137177.1| PREDICTED: actin-related protein 3 [Drosophila miranda]
>gi|1036812510|ref|XP_017137176.1| PREDICTED: transcription factor 25 [Drosophila miranda]
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>gi|1036818098|ref|XP_017137487.1| PREDICTED: DENN domain-containing protein 5B isoform X1 [Dro
>gi|1036818115|ref|XP_017137488.1| PREDICTED: DENN domain-containing protein 5B isoform X1 [Dro
>gi|1036818134|ref|XP_017137489.1| PREDICTED: DENN domain-containing protein 5B isoform X1 [Dro
>gi|1036818155|ref|XP_017137490.1| PREDICTED: DENN domain-containing protein 5B isoform X2 [Dro
>gi|1036837150|ref|XP_017138551.1| PREDICTED: protein phosphatase PTC7 homolog [Drosophila mira
>gi|1036779983|ref|XP_017135347.1| PREDICTED: DEP domain-containing protein 5 isoform X1 [Dros
>gi|1036780004|ref|XP_017135348.1| PREDICTED: DEP domain-containing protein 5 isoform X2 [Dros
>gi|1036780023|ref|XP_017135349.1| PREDICTED: DEP domain-containing protein 5 isoform X3 [Dros
>gi|1036780042|ref|XP_017135350.1| PREDICTED: DEP domain-containing protein 5 isoform X4 [Dros
>gi|1036780061|ref|XP_017135351.1| PREDICTED: DEP domain-containing protein 5 isoform X5 [Dros
>gi|1036780079|ref|XP_017135352.1| PREDICTED: DEP domain-containing protein 5 isoform X6 [Dros
>gi|1036780096|ref|XP_017135353.1| PREDICTED: DEP domain-containing protein 5 isoform X7 [Dros
>gi|1036780114|ref|XP_017135354.1| PREDICTED: DEP domain-containing protein 5 isoform X8 [Dros
>gi|1036780129|ref|XP_017135355.1| PREDICTED: DEP domain-containing protein 5 isoform X9 [Dros
>gi|1036780148|ref|XP_017135356.1| PREDICTED: DEP domain-containing protein 5 isoform X10 [Dros
>gi|1036780277|ref|XP_017135364.1| PREDICTED: uncharacterized protein LOC108151323 [Drosophila
>gi|1036780167|ref|XP_017135357.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X1 [Dr
>gi|1036780239|ref|XP_017135362.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X3 [Dr
>gi|1036780185|ref|XP_017135359.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X2 [Dr
>gi|1036780201|ref|XP_017135360.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X2 [Dr
>gi|1036780220|ref|XP_017135361.1| PREDICTED: puromycin-sensitive aminopeptidase isoform X2 [Dr
>gi|1036780258|ref|XP_017135363.1| PREDICTED: protein cueball [Drosophila miranda]
>gi|1036836205|ref|XP_017138496.1| PREDICTED: uncharacterized protein LOC108153190 [Drosophila
>gi|1036761717|ref|XP_017156974.1| PREDICTED: protein TonB-like [Drosophila miranda]
>gi|1036838351|ref|XP_017138624.1| PREDICTED: protein tracheanless-like [Drosophila miranda]
>gi|1036838239|ref|XP_017138618.1| PREDICTED: protein tracheanless isoform X2 [Drosophila mirand
>gi|1036838258|ref|XP_017138619.1| PREDICTED: protein tracheanless isoform X3 [Drosophila mirand
>gi|1036838220|ref|XP_017138617.1| PREDICTED: protein tracheanless isoform X1 [Drosophila mirand
>gi|1036838332|ref|XP_017138623.1| PREDICTED: protein tracheanless isoform X7 [Drosophila mirand
>gi|1036838296|ref|XP_017138621.1| PREDICTED: protein tracheanless isoform X5 [Drosophila mirand
>gi|1036838313|ref|XP_017138622.1| PREDICTED: protein tracheanless isoform X6 [Drosophila mirand
>gi|1036838277|ref|XP_017138620.1| PREDICTED: protein tracheanless isoform X4 [Drosophila mirand
>gi|1036774020|ref|XP_017135001.1| PREDICTED: uncharacterized protein LOC108151110 isoform X2
>gi|1036774039|ref|XP_017135002.1| PREDICTED: uncharacterized protein LOC108151110 isoform X3
>gi|1036774001|ref|XP_017135000.1| PREDICTED: uncharacterized protein LOC108151110 isoform X1
>gi|1036774058|ref|XP_017135003.1| PREDICTED: uncharacterized protein LOC108151110 isoform X4
>gi|1036774077|ref|XP_017135004.1| PREDICTED: nuclear anchorage protein 1 isoform X5 [Drosophil
>gi|1036774114|ref|XP_017135006.1| PREDICTED: uncharacterized protein LOC108151112 [Drosophila
>gi|1036758187|ref|XP_017156762.1| PREDICTED: uncharacterized protein LOC108165234, partial [Dr
>gi|1036774095|ref|XP_017135005.1| PREDICTED: uncharacterized protein LOC108151111 [Drosophila
>gi|1036842071|ref|XP_017138827.1| PREDICTED: uncharacterized protein C17orf59 homolog [Drosoph
>gi|1036842090|ref|XP_017138828.1| PREDICTED: peptidyl-prolyl cis-trans isomerase-like 1 [Dros
>gi|1036847703|ref|XP_017139129.1| PREDICTED: BTB/POZ domain-containing adapter for CUL3-mediat
>gi|1036811761|ref|XP_017137136.1| PREDICTED: LOW QUALITY PROTEIN: homeodomain-interacting prot
>gi|1036823205|ref|XP_017137765.1| PREDICTED: probable protein phosphatase 2C T23F11.1 [Drosoph
>gi|1036838537|ref|XP_017138635.1| PREDICTED: probable protein phosphatase 2C T23F11.1 [Drosoph
>gi|1036838518|ref|XP_017138634.1| PREDICTED: zinc finger protein 333 [Drosophila miranda]
>gi|1036807662|ref|XP_017136913.1| PREDICTED: mediator of RNA polymerase II transcription subun
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>gi|1036807681|ref|XP_017136914.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1036807718|ref|XP_017136917.1| PREDICTED: SEC14-like protein 2 isoform X2 [Drosophila miran
>gi|1036807699|ref|XP_017136916.1| PREDICTED: SEC14-like protein 2 isoform X1 [Drosophila miran
>gi|1036852268|ref|XP_017139401.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036852284|ref|XP_017139402.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036852303|ref|XP_017139403.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036761736|ref|XP_017156975.1| PREDICTED: DC-STAMP domain-containing protein 2 [Drosophila
>gi|1036851707|ref|XP_017139365.1| PREDICTED: uncharacterized protein LOC108153746 [Drosophila
>gi|1036851599|ref|XP_017139358.1| PREDICTED: alpha-(1,3)-fucosyltransferase C [Drosophila mira
>gi|1036850984|ref|XP_017139320.1| PREDICTED: cohesin subunit SA-1 [Drosophila miranda]
>gi|1036851001|ref|XP_017139321.1| PREDICTED: cohesin subunit SA-1 [Drosophila miranda]
>gi|1036841669|ref|XP_017138813.1| PREDICTED: protein rhomboid [Drosophila miranda]
>gi|1036767373|ref|XP_017134626.1| PREDICTED: N-alpha-acetyltransferase 30 [Drosophila miranda]
>gi|1036829347|ref|XP_017138113.1| PREDICTED: protein rhomboid isoform X1 [Drosophila miranda]
>gi|1036829423|ref|XP_017138117.1| PREDICTED: protein rhomboid isoform X4 [Drosophila miranda]
>gi|1036829366|ref|XP_017138114.1| PREDICTED: protein rhomboid isoform X2 [Drosophila miranda]
>gi|1036829385|ref|XP_017138115.1| PREDICTED: protein rhomboid isoform X3 [Drosophila miranda]
>gi|1036829404|ref|XP_017138116.1| PREDICTED: protein rhomboid isoform X3 [Drosophila miranda]
>gi|1036831592|ref|XP_017138242.1| PREDICTED: protein zer-1 homolog isoform X2 [Drosophila mira
>gi|1036831533|ref|XP_017138239.1| PREDICTED: protein zer-1 homolog isoform X1 [Drosophila mira
>gi|1036831554|ref|XP_017138240.1| PREDICTED: protein zer-1 homolog isoform X1 [Drosophila mira
>gi|1036831573|ref|XP_017138241.1| PREDICTED: protein zer-1 homolog isoform X1 [Drosophila mira
>gi|1036825280|ref|XP_017137881.1| PREDICTED: poly(U)-binding-splicing factor half pint [Drosop
>gi|1036791210|ref|XP_017135990.1| PREDICTED: cell division cycle 5-like protein [Drosophila mi
>gi|1036791247|ref|XP_017135992.1| PREDICTED: RING-box protein 1B [Drosophila miranda]
>gi|1036791229|ref|XP_017135991.1| PREDICTED: uncharacterized protein LOC108151725 [Drosophila
>gi|1036791191|ref|XP_017135988.1| PREDICTED: bifunctional heparan sulfate N-deacetylase/N-sul
>gi|1036799780|ref|XP_017136473.1| PREDICTED: vacuolar protein sorting-associated protein 8 hom
>gi|1036799888|ref|XP_017136478.1| PREDICTED: BTB/POZ domain-containing protein 9-like [Drosoph
>gi|1036799817|ref|XP_017136475.1| PREDICTED: uncharacterized protein C18orf8 isoform X2 [Drosc
>gi|1036799835|ref|XP_017136476.1| PREDICTED: uncharacterized protein C18orf8 isoform X3 [Drosc
>gi|1036799798|ref|XP_017136474.1| PREDICTED: uncharacterized protein C18orf8 isoform X1 [Drosc
>gi|1036799853|ref|XP_017136477.1| PREDICTED: zinc finger protein 93 [Drosophila miranda]
>gi|1036805182|ref|XP_017136778.1| PREDICTED: zinc finger protein 253-like isoform X1 [Drosophi
>gi|1036805202|ref|XP_017136779.1| PREDICTED: zinc finger protein 675-like isoform X2 [Drosophi
>gi|1036805220|ref|XP_017136780.1| PREDICTED: uncharacterized protein LOC108152169 [Drosophila
>gi|1036805239|ref|XP_017136782.1| PREDICTED: ubiquitin-conjugating enzyme E2 D4-like [Drosophi
>gi|1036841716|ref|XP_017138817.1| PREDICTED: putative cyclin-dependent serine/threonine-prote
>gi|1036841731|ref|XP_017138818.1| PREDICTED: uncharacterized protein LOC108153362 [Drosophila
>gi|1036761754|ref|XP_017156976.1| PREDICTED: odorant receptor 65a-like [Drosophila miranda]
>gi|1036761770|ref|XP_017156977.1| PREDICTED: ubiquitin-conjugating enzyme E2 D4-like, partial
>gi|1036766030|ref|XP_017134553.1| PREDICTED: collagenase-like [Drosophila miranda]
>gi|1036761788|ref|XP_017156978.1| PREDICTED: uncharacterized protein LOC108165452 [Drosophila
>gi|1036761809|ref|XP_017156980.1| PREDICTED: protein phosphatase inhibitor 2-like [Drosophila
>gi|1036761827|ref|XP_017156981.1| PREDICTED: protein phosphatase inhibitor 2-like [Drosophila
>gi|1036847347|ref|XP_017139110.1| PREDICTED: protein phosphatase inhibitor 2-like isoform X3
>gi|1036847309|ref|XP_017139108.1| PREDICTED: protein phosphatase inhibitor 2-like isoform X1
>gi|1036847328|ref|XP_017139109.1| PREDICTED: protein phosphatase inhibitor 2-like isoform X2
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>gi|1036762929|ref|XP_017157042.1| PREDICTED: L-lactate dehydrogenase [Drosophila miranda]
>gi|1036762908|ref|XP_017157041.1| PREDICTED: phospholipase A1 member A [Drosophila miranda]
>gi|1036761847|ref|XP_017156982.1| PREDICTED: uncharacterized protein LOC108165455 [Drosophila
>gi|1036761865|ref|XP_017156983.1| PREDICTED: uncharacterized protein LOC108165456 [Drosophila
>gi|1036762659|ref|XP_017157026.1| PREDICTED: multidrug resistance protein homolog 65 [Drosophi
>gi|1036762640|ref|XP_017157025.1| PREDICTED: multidrug resistance protein homolog 65 [Drosophi
>gi|1036762995|ref|XP_017157046.1| PREDICTED: uncharacterized protein LOC108165499 [Drosophila
>gi|1036763012|ref|XP_017157047.1| PREDICTED: uncharacterized protein LOC108165500 [Drosophila
>gi|1036762956|ref|XP_017157043.1| PREDICTED: uncharacterized protein LOC108165497 [Drosophila
>gi|1036763031|ref|XP_017157048.1| PREDICTED: uncharacterized protein LOC108165501 [Drosophila
>gi|1036762574|ref|XP_017157021.1| PREDICTED: laminin subunit alpha [Drosophila miranda]
>gi|1036762676|ref|XP_017157027.1| PREDICTED: transportin-1 [Drosophila miranda]
>gi|1036762975|ref|XP_017157045.1| PREDICTED: splicing factor 3B subunit 6-like protein [Drosop
>gi|1036762731|ref|XP_017157030.1| PREDICTED: zinc finger protein 879 isoform X2 [Drosophila mi
>gi|1036762712|ref|XP_017157029.1| PREDICTED: zinc finger protein 708 isoform X1 [Drosophila mi
>gi|1036762694|ref|XP_017157028.1| PREDICTED: zinc finger protein 668 [Drosophila miranda]
>gi|1036762750|ref|XP_017157031.1| PREDICTED: protein ERGIC-53 [Drosophila miranda]
>gi|1036762593|ref|XP_017157022.1| PREDICTED: talin-2 isoform X1 [Drosophila miranda]
>gi|1036762603|ref|XP_017157023.1| PREDICTED: talin-2 isoform X2 [Drosophila miranda]
>gi|1036762620|ref|XP_017157024.1| PREDICTED: ATP-dependent DNA helicase Q4 [Drosophila miranda
>gi|1036762789|ref|XP_017157034.1| PREDICTED: glycylpeptide N-tetradecanoyltransferase [Drosoph
>gi|1036762804|ref|XP_017157035.1| PREDICTED: UDP-glucuronic acid decarboxylase 1 [Drosophila m
>gi|1036762773|ref|XP_017157033.1| PREDICTED: steroid hormone receptor ERR2 [Drosophila miranda
>gi|1036762827|ref|XP_017157036.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762841|ref|XP_017157037.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762875|ref|XP_017157039.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762859|ref|XP_017157038.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036762890|ref|XP_017157040.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1036761883|ref|XP_017156984.1| PREDICTED: chaperone protein DnaJ, partial [Drosophila miran
>gi|1036815069|ref|XP_017137315.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1036815088|ref|XP_017137316.1| PREDICTED: nedd8-conjugating enzyme Ubc12 [Drosophila mirand
>gi|1036839119|ref|XP_017138673.1| PREDICTED: probable deoxyhypusine synthase isoform X1 [Droso
>gi|1036839138|ref|XP_017138674.1| PREDICTED: probable deoxyhypusine synthase isoform X2 [Droso
>gi|1036849321|ref|XP_017139221.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1036763387|ref|XP_017157063.1| PREDICTED: uncharacterized protein LOC108165517 [Drosophila
>gi|1036850837|ref|XP_017139313.1| PREDICTED: uncharacterized protein LOC108153698 isoform X1
>gi|1036850856|ref|XP_017139314.1| PREDICTED: uncharacterized protein LOC108153698 isoform X2
>gi|1036766424|ref|XP_017134573.1| PREDICTED: uncharacterized protein LOC108150784 [Drosophila
>gi|1036850763|ref|XP_017139308.1| PREDICTED: pupal cuticle protein C1B [Drosophila miranda]
>gi|1036850782|ref|XP_017139309.1| PREDICTED: vitelline membrane protein Vm26Ab [Drosophila mir
>gi|1036850746|ref|XP_017139307.1| PREDICTED: eukaryotic translation initiation factor 4E [Dros
>gi|1036765435|ref|XP_017157170.1| PREDICTED: uncharacterized protein LOC108165606 [Drosophila
>gi|1036765453|ref|XP_017157171.1| PREDICTED: uncharacterized protein LOC108165606 [Drosophila
>gi|1036847439|ref|XP_017139116.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036847458|ref|XP_017139117.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036847476|ref|XP_017139118.1| PREDICTED: Down syndrome cell adhesion molecule-like protei
>gi|1036784789|ref|XP_017135625.1| PREDICTED: collagen type IV alpha-3-binding protein [Drosoph
>gi|1036784805|ref|XP_017135626.1| PREDICTED: uncharacterized protein LOC108151498 [Drosophila
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>gi|1036784879|ref|XP_017135631.1| PREDICTED: ras-related protein Rab-43 isoform X1 [Drosophila
>gi|1036784898|ref|XP_017135632.1| PREDICTED: ras-related protein Rab-43 isoform X2 [Drosophila
>gi|1036784919|ref|XP_017135633.1| PREDICTED: ADP-ribosylation factor-like protein 5A [Drosophi
>gi|1036784772|ref|XP_017135624.1| PREDICTED: uncharacterized protein LOC108151495 [Drosophila
>gi|1036784823|ref|XP_017135627.1| PREDICTED: selenocysteine insertion sequence-binding protei
>gi|1036761902|ref|XP_017156985.1| PREDICTED: gustatory receptor for bitter taste 66a [Drosophi
>gi|1036784842|ref|XP_017135629.1| PREDICTED: bax inhibitor 1 isoform X1 [Drosophila miranda]
>gi|1036784861|ref|XP_017135630.1| PREDICTED: bax inhibitor 1 isoform X2 [Drosophila miranda]
>gi|1036823396|ref|XP_017137776.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036823377|ref|XP_017137775.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036823358|ref|XP_017137774.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036823337|ref|XP_017137773.1| PREDICTED: cleavage and polyadenylation specificity factor s
>gi|1036848334|ref|XP_017139164.1| PREDICTED: alkaline phosphatase [Drosophila miranda]
>gi|1036838391|ref|XP_017138626.1| PREDICTED: syntaxin-7 [Drosophila miranda]
>gi|1036846454|ref|XP_017139058.1| PREDICTED: fork head domain-containing protein crocodile [Di
>gi|1036847123|ref|XP_017139097.1| PREDICTED: forkhead box protein biniou [Drosophila miranda]
>gi|1036761920|ref|XP_017156986.1| PREDICTED: keratin-associated protein 5-1-like [Drosophila r
>gi|1036764298|ref|XP_017157113.1| PREDICTED: sperm-specific protein Don juan [Drosophila mirar
>gi|1036764318|ref|XP_017157114.1| PREDICTED: sperm-specific protein Don juan [Drosophila mirar
>gi|1036809371|ref|XP_017137004.1| PREDICTED: G protein alpha i subunit [Drosophila miranda]
>gi|1036809390|ref|XP_017137005.1| PREDICTED: uncharacterized protein LOC108152294 [Drosophila
>gi|1036809352|ref|XP_017137003.1| PREDICTED: putative serine protease F56F10.1 [Drosophila mi
>gi|1036809335|ref|XP_017137002.1| PREDICTED: cilia- and flagella-associated protein 52 [Drosop
>gi|1036784699|ref|XP_017135620.1| PREDICTED: UDP-glucose 6-dehydrogenase [Drosophila miranda]
>gi|1036784717|ref|XP_017135621.1| PREDICTED: SURF1-like protein [Drosophila miranda]
>gi|1036784736|ref|XP_017135622.1| PREDICTED: ubiquinol-cytochrome-c reductase complex assembly
>gi|1036784753|ref|XP_017135623.1| PREDICTED: transcription factor Adf-1 [Drosophila miranda]
>gi|1036784680|ref|XP_017135619.1| PREDICTED: CLIP-associating protein [Drosophila miranda]
>gi|1036825221|ref|XP_017137877.1| PREDICTED: LOW QUALITY PROTEIN: asparagine synthetase B [glu
>gi|1036825240|ref|XP_017137878.1| PREDICTED: asparagine synthetase [glutamine-hydrolyzing] 2-1
>gi|1036829537|ref|XP_017138121.1| PREDICTED: uncharacterized protein LOC108152938 [Drosophila
>gi|1036829476|ref|XP_017138118.1| PREDICTED: adenylate cyclase type 8 isoform X1 [Drosophila r
>gi|1036829497|ref|XP_017138119.1| PREDICTED: adenylate cyclase type 8 isoform X2 [Drosophila r
>gi|1036829516|ref|XP_017138120.1| PREDICTED: adenylate cyclase type 8 isoform X3 [Drosophila r
>gi|1036805107|ref|XP_017136773.1| PREDICTED: GTPase-activating protein isoform X1 [Drosophila
>gi|1036805127|ref|XP_017136774.1| PREDICTED: GTPase-activating protein isoform X2 [Drosophila
>gi|1036786635|ref|XP_017135730.1| PREDICTED: uncharacterized protein LOC108151570 isoform X1
>gi|1036786656|ref|XP_017135731.1| PREDICTED: myotubularin-related protein 9-like isoform X2 [
>gi|1036786675|ref|XP_017135732.1| PREDICTED: myotubularin-related protein 9-like isoform X3 [
>gi|1036786694|ref|XP_017135733.1| PREDICTED: ATP-binding cassette sub-family E member 1 [Droso
>gi|1036786713|ref|XP_017135734.1| PREDICTED: uncharacterized protein LOC108151573 [Drosophila
>gi|1036786791|ref|XP_017135739.1| PREDICTED: uncharacterized protein LOC108151575 [Drosophila
>gi|1036786810|ref|XP_017135740.1| PREDICTED: uncharacterized protein LOC108151575 [Drosophila
>gi|1036786734|ref|XP_017135735.1| PREDICTED: N-acetylgalactosamine kinase [Drosophila miranda]
>gi|1036786753|ref|XP_017135736.1| PREDICTED: N-acetylgalactosamine kinase [Drosophila miranda]
>gi|1036786772|ref|XP_017135738.1| PREDICTED: N-acetylgalactosamine kinase [Drosophila miranda]
>gi|1036800456|ref|XP_017136510.1| PREDICTED: vesicle transport protein USE1 isoform X1 [Drosop
>gi|1036800474|ref|XP_017136511.1| PREDICTED: vesicle transport protein USE1 isoform X2 [Drosop
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>gi|1036800492|ref|XP_017136512.1| PREDICTED: dihydropteridine reductase [Drosophila miranda]
>gi|1036800271|ref|XP_017136500.1| PREDICTED: protein boule isoform X1 [Drosophila miranda]
>gi|1036800290|ref|XP_017136501.1| PREDICTED: protein boule isoform X1 [Drosophila miranda]
>gi|1036800309|ref|XP_017136502.1| PREDICTED: protein boule isoform X2 [Drosophila miranda]
>gi|1036800327|ref|XP_017136503.1| PREDICTED: protein boule isoform X3 [Drosophila miranda]
>gi|1036800344|ref|XP_017136504.1| PREDICTED: protein boule isoform X4 [Drosophila miranda]
>gi|1036800362|ref|XP_017136505.1| PREDICTED: protein boule isoform X5 [Drosophila miranda]
>gi|1036800382|ref|XP_017136506.1| PREDICTED: protein boule isoform X6 [Drosophila miranda]
>gi|1036800401|ref|XP_017136507.1| PREDICTED: protein boule isoform X7 [Drosophila miranda]
>gi|1036800418|ref|XP_017136508.1| PREDICTED: protein boule isoform X8 [Drosophila miranda]
>gi|1036800437|ref|XP_017136509.1| PREDICTED: protein boule isoform X9 [Drosophila miranda]
>gi|1036848370|ref|XP_017139166.1| PREDICTED: uncharacterized protein LOC108153596 [Drosophila
>gi|1036850361|ref|XP_017139284.1| PREDICTED: uncharacterized protein LOC108153678 [Drosophila
>gi|1036818582|ref|XP_017137514.1| PREDICTED: uncharacterized protein LOC108152581 isoform X4
>gi|1036818563|ref|XP_017137513.1| PREDICTED: uncharacterized protein LOC108152581 isoform X3
>gi|1036818525|ref|XP_017137511.1| PREDICTED: uncharacterized protein LOC108152581 isoform X1
>gi|1036818544|ref|XP_017137512.1| PREDICTED: uncharacterized protein LOC108152581 isoform X2
>gi|1036818603|ref|XP_017137515.1| PREDICTED: uncharacterized protein LOC108152581 isoform X5
>gi|1036818622|ref|XP_017137516.1| PREDICTED: mitochondrial inner membrane protein COX18 [Droso
>gi|1036786319|ref|XP_017135714.1| PREDICTED: F-box only protein 39 [Drosophila miranda]
>gi|1036786266|ref|XP_017135710.1| PREDICTED: arginine kinase isoform X1 [Drosophila miranda]
>gi|1036786285|ref|XP_017135711.1| PREDICTED: arginine kinase isoform X2 [Drosophila miranda]
>gi|1036786302|ref|XP_017135712.1| PREDICTED: arginine kinase isoform X3 [Drosophila miranda]
>gi|1036786338|ref|XP_017135715.1| PREDICTED: arginine kinase-like [Drosophila miranda]
>gi|1036841632|ref|XP_017138811.1| PREDICTED: T-box transcription factor TBX6 [Drosophila miran
>gi|1036838591|ref|XP_017138638.1| PREDICTED: T-box protein 2 [Drosophila miranda]
>gi|1036845704|ref|XP_017139016.1| PREDICTED: T-box transcription factor TBX6 [Drosophila miran
>gi|1036828014|ref|XP_017138036.1| PREDICTED: ubiquitin-protein ligase E3B [Drosophila miranda]
>gi|1036810809|ref|XP_017137085.1| PREDICTED: protein Smaug isoform X2 [Drosophila miranda]
>gi|1036810828|ref|XP_017137086.1| PREDICTED: protein Smaug isoform X3 [Drosophila miranda]
>gi|1036810771|ref|XP_017137083.1| PREDICTED: protein Smaug isoform X1 [Drosophila miranda]
>gi|1036810789|ref|XP_017137084.1| PREDICTED: protein Smaug isoform X1 [Drosophila miranda]
>gi|1036835814|ref|XP_017138472.1| PREDICTED: UPF0193 protein EVG1 homolog [Drosophila miranda]
>gi|1036835795|ref|XP_017138471.1| PREDICTED: protein phosphatase methylesterase 1 [Drosophila
>gi|1036843002|ref|XP_017138879.1| PREDICTED: uncharacterized protein LOC108153400 isoform X1
>gi|1036843021|ref|XP_017138880.1| PREDICTED: uncharacterized protein LOC108153400 isoform X1
>gi|1036843041|ref|XP_017138881.1| PREDICTED: uncharacterized protein LOC108153400 isoform X2
>gi|1036822080|ref|XP_017137700.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X1
>gi|1036822098|ref|XP_017137702.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X1
>gi|1036822116|ref|XP_017137703.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X1
>gi|1036822134|ref|XP_017137704.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X2
>gi|1036822174|ref|XP_017137706.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X4
>gi|1036822193|ref|XP_017137707.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X5
>gi|1036822153|ref|XP_017137705.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X3
>gi|1036822212|ref|XP_017137708.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X6
>gi|1036822231|ref|XP_017137709.1| PREDICTED: F-BAR and double SH3 domains protein 2 isoform X7
>gi|1036828491|ref|XP_017138064.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828414|ref|XP_017138059.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
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>gi|1036828726|ref|XP_017138076.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828510|ref|XP_017138065.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828432|ref|XP_017138060.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828609|ref|XP_017138070.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828569|ref|XP_017138068.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828550|ref|XP_017138067.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828451|ref|XP_017138062.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828357|ref|XP_017138056.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828376|ref|XP_017138057.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828688|ref|XP_017138074.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828590|ref|XP_017138069.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828470|ref|XP_017138063.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828395|ref|XP_017138058.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828531|ref|XP_017138066.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828630|ref|XP_017138071.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828648|ref|XP_017138072.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828669|ref|XP_017138073.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036828707|ref|XP_017138075.1| PREDICTED: gamma-aminobutyric acid receptor subunit beta iso
>gi|1036834176|ref|XP_017138379.1| PREDICTED: proton-associated sugar transporter A [Drosophila
>gi|1036834195|ref|XP_017138380.1| PREDICTED: proton-associated sugar transporter A [Drosophila
>gi|1036835275|ref|XP_017138440.1| PREDICTED: sodium-dependent nutrient amino acid transporter
>gi|1036847722|ref|XP_017139130.1| PREDICTED: glutamate receptor 1 [Drosophila miranda]
>gi|1036850691|ref|XP_017139304.1| PREDICTED: peptidoglycan-recognition protein LA isoform X1
>gi|1036850729|ref|XP_017139306.1| PREDICTED: peptidoglycan-recognition protein LA isoform X3
>gi|1036850710|ref|XP_017139305.1| PREDICTED: peptidoglycan-recognition protein LA isoform X2
>gi|1036803078|ref|XP_017136658.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803097|ref|XP_017136660.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803116|ref|XP_017136661.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803134|ref|XP_017136662.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803152|ref|XP_017136663.1| PREDICTED: peptidoglycan-recognition protein LC isoform X1
>gi|1036803190|ref|XP_017136665.1| PREDICTED: peptidoglycan-recognition protein LC isoform X3
>gi|1036803228|ref|XP_017136667.1| PREDICTED: peptidoglycan-recognition protein LC isoform X5
>gi|1036803209|ref|XP_017136666.1| PREDICTED: peptidoglycan-recognition protein LC isoform X4
>gi|1036803171|ref|XP_017136664.1| PREDICTED: peptidoglycan-recognition protein LC isoform X2
>gi|1036803246|ref|XP_017136668.1| PREDICTED: peptidoglycan-recognition protein LC isoform X6
>gi|1036803265|ref|XP_017136669.1| PREDICTED: peptidoglycan-recognition protein LC isoform X7
>gi|1036803355|ref|XP_017136675.1| PREDICTED: peptidoglycan-recognition protein LF isoform X1
>gi|1036803371|ref|XP_017136676.1| PREDICTED: peptidoglycan-recognition protein LF isoform X2
>gi|1036803338|ref|XP_017136674.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803322|ref|XP_017136673.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803303|ref|XP_017136672.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803284|ref|XP_017136671.1| PREDICTED: UTP--glucose-1-phosphate uridylyltransferase iso
>gi|1036803388|ref|XP_017136677.1| PREDICTED: uncharacterized protein LOC108152112 [Drosophila
>gi|1036804590|ref|XP_017136744.1| PREDICTED: small VCP/p97-interacting protein [Drosophila mir
>gi|1036804572|ref|XP_017136743.1| PREDICTED: autophagy-related protein 2 homolog A [Drosophila
>gi|1036821275|ref|XP_017137656.1| PREDICTED: uncharacterized protein LOC108152670 [Drosophila
>gi|1036821256|ref|XP_017137655.1| PREDICTED: putative adenosylhomocysteinase 2 [Drosophila mir
>gi|1036836040|ref|XP_017138486.1| PREDICTED: upstream activation factor subunit spp27 [Drosoph
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>gi|1036802282|ref|XP_017136617.1| PREDICTED: histone-lysine N-methyltransferase 2D [Drosophila
>gi|1036837979|ref|XP_017138602.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1036837998|ref|XP_017138603.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1036838017|ref|XP_017138604.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1036838069|ref|XP_017138608.1| PREDICTED: synaptic vesicle glycoprotein 2B-like [Drosophila
>gi|1036838051|ref|XP_017138607.1| PREDICTED: putative transporter svop-1 [Drosophila miranda]
>gi|1036838034|ref|XP_017138605.1| PREDICTED: putative transporter SVOPL [Drosophila miranda]
>gi|1036764224|ref|XP_017157109.1| PREDICTED: uncharacterized protein LOC108165545 [Drosophila
>gi|1037099942|ref|XP_017144749.1| PREDICTED: extended synaptotagmin-2 isoform X2 [Drosophila n
>gi|1037099932|ref|XP_017144747.1| PREDICTED: extended synaptotagmin-2 isoform X1 [Drosophila n
>gi|1037099960|ref|XP_017144751.1| PREDICTED: extended synaptotagmin-1 isoform X4 [Drosophila n
>gi|1037099950|ref|XP_017144750.1| PREDICTED: extended synaptotagmin-2 isoform X3 [Drosophila n
>gi|1037079607|ref|XP_017143172.1| PREDICTED: protein CREBRF homolog [Drosophila miranda]
>gi|1037079621|ref|XP_017143173.1| PREDICTED: protein CREBRF homolog [Drosophila miranda]
>gi|1037079635|ref|XP_017143174.1| PREDICTED: BUD13 homolog [Drosophila miranda]
>gi|1037105933|ref|XP_017145427.1| PREDICTED: uncharacterized protein LOC108157757 [Drosophila
>gi|1037034935|ref|XP_017140187.1| PREDICTED: intraflagellar transport protein 56 [Drosophila n
>gi|1037085350|ref|XP_017143610.1| PREDICTED: uncharacterized protein LOC108156584 [Drosophila
>gi|1037033899|ref|XP_017140125.1| PREDICTED: mitochondrial import receptor subunit TOM40 homol
>gi|1037034305|ref|XP_017140151.1| PREDICTED: uncharacterized protein LOC108154396 [Drosophila
>gi|1037034760|ref|XP_017140178.1| PREDICTED: tetratricopeptide repeat protein 12 [Drosophila n
>gi|1037022592|ref|XP_017139431.1| PREDICTED: uncharacterized protein LOC108153843 [Drosophila
>gi|1037037994|ref|XP_017140356.1| PREDICTED: tetratricopeptide repeat protein 12 [Drosophila n
>gi|1037105894|ref|XP_017145421.1| PREDICTED: sodium-independent sulfate anion transporter [Dro
>gi|1037103816|ref|XP_017145131.1| PREDICTED: uncharacterized protein LOC108157539 [Drosophila
>gi|1037034339|ref|XP_017140153.1| PREDICTED: uncharacterized protein LOC108154398 [Drosophila
>gi|1037090337|ref|XP_017143965.1| PREDICTED: uncharacterized protein LOC108156797 [Drosophila
>gi|1037093993|ref|XP_017144226.1| PREDICTED: LOW QUALITY PROTEIN: xanthine dehydrogenase [Dros
>gi|1037022607|ref|XP_017139432.1| PREDICTED: xanthine dehydrogenase/oxidase-like [Drosophila n
>gi|1037093979|ref|XP_017144225.1| PREDICTED: indole-3-acetaldehyde oxidase [Drosophila miranda
>gi|1037092203|ref|XP_017144105.1| PREDICTED: LOW QUALITY PROTEIN: indole-3-acetaldehyde oxidas
>gi|1037092217|ref|XP_017144107.1| PREDICTED: xanthine dehydrogenase-like [Drosophila miranda]
>gi|1037037448|ref|XP_017140328.1| PREDICTED: uncharacterized protein LOC108154536 [Drosophila
>gi|1037033882|ref|XP_017140124.1| PREDICTED: uncharacterized protein LOC108154370 [Drosophila
>gi|1037105540|ref|XP_017145373.1| PREDICTED: uncharacterized protein LOC108157713 [Drosophila
>gi|1037105114|ref|XP_017145311.1| PREDICTED: uncharacterized protein LOC108157663 [Drosophila
>gi|1037103621|ref|XP_017145105.1| PREDICTED: dynein beta chain, ciliary isoform X2 [Drosophila
>gi|1037103613|ref|XP_017145104.1| PREDICTED: dynein beta chain, ciliary isoform X1 [Drosophila
>gi|1037022621|ref|XP_017139433.1| PREDICTED: uncharacterized protein LOC108153845 [Drosophila
>gi|1037035151|ref|XP_017140200.1| PREDICTED: uncharacterized protein LOC108154438 [Drosophila
>gi|1037034409|ref|XP_017140157.1| PREDICTED: uncharacterized protein LOC108154401 isoform X1
>gi|1037034427|ref|XP_017140159.1| PREDICTED: uncharacterized protein LOC108154401 isoform X2
>gi|1037034447|ref|XP_017140160.1| PREDICTED: uncharacterized protein LOC108154401 isoform X3
>gi|1037032758|ref|XP_017140057.1| PREDICTED: circadian clock-controlled protein [Drosophila mi
>gi|1037034813|ref|XP_017140182.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037091349|ref|XP_017144042.1| PREDICTED: protein Malvolio isoform X2 [Drosophila miranda]
>gi|1037091335|ref|XP_017144041.1| PREDICTED: protein Malvolio isoform X1 [Drosophila miranda]
>gi|1037093883|ref|XP_017144218.1| PREDICTED: hematopoietic lineage cell-specific protein [Dros
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>gi|1037093897|ref|XP_017144219.1| PREDICTED: hematopoietic lineage cell-specific protein [Dros
>gi|1037082949|ref|XP_017143427.1| PREDICTED: annexin B9 isoform X3 [Drosophila miranda]
>gi|1037082937|ref|XP_017143426.1| PREDICTED: annexin B9 isoform X2 [Drosophila miranda]
>gi|1037082910|ref|XP_017143424.1| PREDICTED: annexin B9 isoform X1 [Drosophila miranda]
>gi|1037082924|ref|XP_017143425.1| PREDICTED: annexin B9 isoform X1 [Drosophila miranda]
>gi|1037082896|ref|XP_017143422.1| PREDICTED: uridine 5'-monophosphate synthase [Drosophila mi
>gi|1037034221|ref|XP_017140144.1| PREDICTED: doublesex- and mab-3-related transcription facto
>gi|1037033178|ref|XP_017140083.1| PREDICTED: hematopoietically-expressed homeobox protein Hhe
>gi|1037067328|ref|XP_017142329.1| PREDICTED: hornerin [Drosophila miranda]
>gi|1037067344|ref|XP_017142330.1| PREDICTED: uncharacterized protein LOC108155781 [Drosophila
>gi|1037085504|ref|XP_017143620.1| PREDICTED: hematological and neurological expressed 1-like p
>gi|1037085490|ref|XP_017143619.1| PREDICTED: THO complex subunit 4 [Drosophila miranda]
>gi|1037092245|ref|XP_017144109.1| PREDICTED: dephospho-CoA kinase [Drosophila miranda]
>gi|1037092231|ref|XP_017144108.1| PREDICTED: UTP:RNA uridylyltransferase 1 [Drosophila miranda
>gi|1037085365|ref|XP_017143611.1| PREDICTED: serine/threonine-protein kinase/endoribonuclease
>gi|1037085989|ref|XP_017143655.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085933|ref|XP_017143651.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037085921|ref|XP_017143650.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037086000|ref|XP_017143656.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085961|ref|XP_017143653.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085947|ref|XP_017143652.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085975|ref|XP_017143654.1| PREDICTED: calcium uptake protein 3, mitochondrial isoform
>gi|1037085907|ref|XP_017143649.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037085893|ref|XP_017143647.1| PREDICTED: calcium uptake protein 2, mitochondrial isoform
>gi|1037086014|ref|XP_017143657.1| PREDICTED: uncharacterized protein LOC108156613 [Drosophila
>gi|1037085865|ref|XP_017143645.1| PREDICTED: serine-rich adhesin for platelets isoform X1 [Dr
>gi|1037085879|ref|XP_017143646.1| PREDICTED: serine-rich adhesin for platelets isoform X2 [Dr
>gi|1037106464|ref|XP_017145500.1| PREDICTED: 4-coumarate--CoA ligase-like 7 [Drosophila miran
>gi|1037106181|ref|XP_017145461.1| PREDICTED: 4-coumarate--CoA ligase-like [Drosophila miranda]
>gi|1037033750|ref|XP_017140115.1| PREDICTED: luciferin 4-monooxygenase [Drosophila miranda]
>gi|1037105779|ref|XP_017145406.1| PREDICTED: luciferin 4-monooxygenase-like [Drosophila miran
>gi|1037096331|ref|XP_017144415.1| PREDICTED: glutamate-gated chloride channel isoform X8 [Dros
>gi|1037096303|ref|XP_017144413.1| PREDICTED: glutamate-gated chloride channel isoform X6 [Dros
>gi|1037096252|ref|XP_017144409.1| PREDICTED: glutamate-gated chloride channel isoform X2 [Dros
>gi|1037096343|ref|XP_017144416.1| PREDICTED: glutamate-gated chloride channel isoform X9 [Dros
>gi|1037096291|ref|XP_017144412.1| PREDICTED: glutamate-gated chloride channel isoform X5 [Dros
>gi|1037096281|ref|XP_017144411.1| PREDICTED: glutamate-gated chloride channel isoform X4 [Dros
>gi|1037096226|ref|XP_017144406.1| PREDICTED: glutamate-gated chloride channel isoform X1 [Dros
>gi|1037096238|ref|XP_017144407.1| PREDICTED: glutamate-gated chloride channel isoform X1 [Dros
>gi|1037096315|ref|XP_017144414.1| PREDICTED: glutamate-gated chloride channel isoform X7 [Dros
>gi|1037096355|ref|XP_017144417.1| PREDICTED: glutamate-gated chloride channel isoform X10 [Dro
>gi|1037096270|ref|XP_017144410.1| PREDICTED: glutamate-gated chloride channel isoform X3 [Dros
>gi|1037038914|ref|XP_017140416.1| PREDICTED: uncharacterized protein LOC108154611 [Drosophila
>gi|1037096990|ref|XP_017144474.1| PREDICTED: pyrazinamidase/nicotinamidase [Drosophila miranda
>gi|1037097002|ref|XP_017144475.1| PREDICTED: pyrazinamidase/nicotinamidase [Drosophila miranda
>gi|1037096978|ref|XP_017144473.1| PREDICTED: serine-rich adhesin for platelets [Drosophila mi
>gi|1037077638|ref|XP_017143040.1| PREDICTED: glycine receptor subunit alpha-4 [Drosophila mira
>gi|1037077612|ref|XP_017143037.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
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>gi|1037077649|ref|XP_017143041.1| PREDICTED: developmentally-regulated GTP-binding protein 2
>gi|1037077625|ref|XP_017143039.1| PREDICTED: probable nucleoporin Nup58 [Drosophila miranda]
>gi|1037102983|ref|XP_017145032.1| PREDICTED: V-type proton ATPase subunit G [Drosophila miran
>gi|1037077531|ref|XP_017143032.1| PREDICTED: anoctamin-1 isoform X1 [Drosophila miranda]
>gi|1037077547|ref|XP_017143033.1| PREDICTED: anoctamin-1 isoform X2 [Drosophila miranda]
>gi|1037077559|ref|XP_017143034.1| PREDICTED: anoctamin-1 isoform X3 [Drosophila miranda]
>gi|1037077585|ref|XP_017143036.1| PREDICTED: uncharacterized protein LOC108156209 [Drosophila
>gi|1037077571|ref|XP_017143035.1| PREDICTED: organic cation transporter protein [Drosophila mi
>gi|1037022641|ref|XP_017139434.1| PREDICTED: uncharacterized protein LOC108153847 [Drosophila
>gi|1037100890|ref|XP_017144829.1| PREDICTED: organic cation transporter-like protein [Drosophi
>gi|1037106073|ref|XP_017145446.1| PREDICTED: organic cation transporter-like protein isoform )
>gi|1037106065|ref|XP_017145445.1| PREDICTED: organic cation transporter-like protein isoform )
>gi|1037106079|ref|XP_017145447.1| PREDICTED: uncharacterized protein LOC108157773 [Drosophila
>gi|1037047793|ref|XP_017141010.1| PREDICTED: cuticle protein 7 [Drosophila miranda]
>gi|1037047731|ref|XP_017141005.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X16 [Droso
>gi|1037047715|ref|XP_017141004.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X15 [Droso
>gi|1037047624|ref|XP_017140997.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X9 [Drosop
>gi|1037047608|ref|XP_017140996.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X8 [Drosop
>gi|1037047592|ref|XP_017140995.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X7 [Drosop
>gi|1037047682|ref|XP_017141002.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X13 [Dros
>gi|1037047576|ref|XP_017140994.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X6 [Drosop
>gi|1037047667|ref|XP_017141001.1| PREDICTED: dystrophin isoform X12 [Drosophila miranda]
>gi|1037047637|ref|XP_017140999.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X10 [Dros
>gi|1037047493|ref|XP_017140990.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X2 [Drosop
>gi|1037047697|ref|XP_017141003.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X14 [Dros
>gi|1037047652|ref|XP_017141000.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X11 [Dros
>gi|1037047540|ref|XP_017140992.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X4 [Drosop
>gi|1037047508|ref|XP_017140991.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X3 [Drosop
>gi|1037047479|ref|XP_017140989.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X1 [Drosop
>gi|1037047556|ref|XP_017140993.1| PREDICTED: dystrophin, isoforms A/C/F/G/H isoform X5 [Drosop
>gi|1037047821|ref|XP_017141012.1| PREDICTED: uncharacterized protein LOC108154984 [Drosophila
>gi|1037047747|ref|XP_017141006.1| PREDICTED: dystrophin-like [Drosophila miranda]
>gi|1037047763|ref|XP_017141007.1| PREDICTED: uncharacterized protein LOC108154980 [Drosophila
>gi|1037047807|ref|XP_017141011.1| PREDICTED: IDLSRF-like peptide [Drosophila miranda]
>gi|1037047779|ref|XP_017141009.1| PREDICTED: uncharacterized protein LOC108154981 [Drosophila
>gi|1037055229|ref|XP_017141518.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037055247|ref|XP_017141519.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037022659|ref|XP_017139435.1| PREDICTED: LOW QUALITY PROTEIN: aprataxin-like protein [Dros
>gi|1037055351|ref|XP_017141527.1| PREDICTED: LOW QUALITY PROTEIN: putative DNA helicase Ino80
>gi|1037055295|ref|XP_017141523.1| PREDICTED: uncharacterized protein LOC108155315 [Drosophila
>gi|1037055215|ref|XP_017141517.1| PREDICTED: LOW QUALITY PROTEIN: thymus-specific serine prote
>gi|1037055263|ref|XP_017141520.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037055277|ref|XP_017141521.1| PREDICTED: putative serine protease K12H4.7 [Drosophila mira
>gi|1037055336|ref|XP_017141526.1| PREDICTED: pollen-specific leucine-rich repeat extensin-like
>gi|1037055309|ref|XP_017141524.1| PREDICTED: uncharacterized protein LOC108155316 [Drosophila
>gi|1037055321|ref|XP_017141525.1| PREDICTED: uncharacterized protein LOC108155317 [Drosophila
>gi|1037033126|ref|XP_017140080.1| PREDICTED: uncharacterized protein LOC108154329 [Drosophila
>gi|1037079796|ref|XP_017143186.1| PREDICTED: neurogenic locus protein delta [Drosophila miran
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>gi|1037103661|ref|XP_017145111.1| PREDICTED: zinc finger CCCH domain-containing protein 18-li
>gi|1037103653|ref|XP_017145110.1| PREDICTED: uncharacterized protein LOC108157518 [Drosophila
>gi|1037085560|ref|XP_017143625.1| PREDICTED: uncharacterized protein LOC108156595 [Drosophila
>gi|1037085533|ref|XP_017143623.1| PREDICTED: protein unc-79 homolog isoform X2 [Drosophila mi
>gi|1037085520|ref|XP_017143622.1| PREDICTED: protein unc-79 homolog isoform X1 [Drosophila mi
>gi|1037085547|ref|XP_017143624.1| PREDICTED: uncharacterized protein LOC108156594 [Drosophila
>gi|1037105868|ref|XP_017145418.1| PREDICTED: uncharacterized protein LOC108157748 [Drosophila
>gi|1037105860|ref|XP_017145417.1| PREDICTED: neurofilament medium polypeptide [Drosophila mira
>gi|1037096899|ref|XP_017144465.1| PREDICTED: phosphatidylinositol transfer protein alpha isofo
>gi|1037096873|ref|XP_017144463.1| PREDICTED: phosphatidylinositol transfer protein alpha isofo
>gi|1037096887|ref|XP_017144464.1| PREDICTED: phosphatidylinositol transfer protein alpha isofo
>gi|1037105243|ref|XP_017145329.1| PREDICTED: cryptochrome-1 [Drosophila miranda]
>gi|1037094670|ref|XP_017144278.1| PREDICTED: 45 kDa calcium-binding protein [Drosophila miran
>gi|1037094634|ref|XP_017144276.1| PREDICTED: BRCA1-associated protein [Drosophila miranda]
>gi|1037094680|ref|XP_017144279.1| PREDICTED: uncharacterized protein LOC108156996 [Drosophila
>gi|1037094746|ref|XP_017144283.1| PREDICTED: zinc finger protein squeeze [Drosophila miranda]
>gi|1037094760|ref|XP_017144284.1| PREDICTED: zinc finger protein squeeze [Drosophila miranda]
>gi|1037094771|ref|XP_017144285.1| PREDICTED: zinc finger protein squeeze [Drosophila miranda]
>gi|1037077224|ref|XP_017143009.1| PREDICTED: C-1-tetrahydrofolate synthase, cytoplasmic [Dros
>gi|1037077178|ref|XP_017143005.1| PREDICTED: probable methyltransferase-like protein 15 homolo
>gi|1037077208|ref|XP_017143008.1| PREDICTED: uncharacterized protein LOC108156188 [Drosophila
>gi|1037077194|ref|XP_017143006.1| PREDICTED: peptide deformylase, mitochondrial [Drosophila mi
>gi|1037101196|ref|XP_017144859.1| PREDICTED: peptide deformylase, mitochondrial [Drosophila mi
>gi|1037101134|ref|XP_017144853.1| PREDICTED: uncharacterized protein LOC108157344 [Drosophila
>gi|1037101156|ref|XP_017144855.1| PREDICTED: uncharacterized protein LOC108157346 [Drosophila
>gi|1037101144|ref|XP_017144854.1| PREDICTED: uncharacterized protein LOC108157345 [Drosophila
>gi|1037101176|ref|XP_017144857.1| PREDICTED: uncharacterized protein LOC108157348 isoform X1
>gi|1037101186|ref|XP_017144858.1| PREDICTED: uncharacterized protein LOC108157348 isoform X2
>gi|1037101166|ref|XP_017144856.1| PREDICTED: very low-density lipoprotein receptor-like [Dros
>gi|1037022676|ref|XP_017139436.1| PREDICTED: bifunctional protein FolD-like [Drosophila miran
>gi|1037103567|ref|XP_017145098.1| PREDICTED: mitochondrial import receptor subunit TOM20 homol
>gi|1037103559|ref|XP_017145097.1| PREDICTED: tachykinin-like peptides receptor 86C isoform X2
>gi|1037103551|ref|XP_017145096.1| PREDICTED: tachykinin-like peptides receptor 86C isoform X1
>gi|1037106448|ref|XP_017145498.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X1 [Dros
>gi|1037106456|ref|XP_017145499.1| PREDICTED: synaptic vesicle glycoprotein 2B isoform X2 [Dros
>gi|1037093004|ref|XP_017144158.1| PREDICTED: synaptic vesicle glycoprotein 2B [Drosophila mira
>gi|1037093018|ref|XP_017144159.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1037077903|ref|XP_017143060.1| PREDICTED: uncharacterized protein LOC108156231 isoform X3
>gi|1037077875|ref|XP_017143058.1| PREDICTED: uncharacterized protein LOC108156231 isoform X1
>gi|1037077919|ref|XP_017143061.1| PREDICTED: uncharacterized protein LOC108156231 isoform X4
>gi|1037077889|ref|XP_017143059.1| PREDICTED: uncharacterized protein LOC108156231 isoform X2
>gi|1037077933|ref|XP_017143063.1| PREDICTED: general transcription factor IIF subunit 2 [Dros
>gi|1037078583|ref|XP_017143106.1| PREDICTED: alpha-1,2-mannosyltransferase ALG9 [Drosophila mi
>gi|1037078569|ref|XP_017143105.1| PREDICTED: protocadherin Fat 1 isoform X4 [Drosophila miran
>gi|1037078539|ref|XP_017143103.1| PREDICTED: protocadherin Fat 1 isoform X2 [Drosophila miran
>gi|1037078552|ref|XP_017143104.1| PREDICTED: protocadherin Fat 1 isoform X3 [Drosophila miran
>gi|1037078512|ref|XP_017143102.1| PREDICTED: protocadherin Fat 1 isoform X1 [Drosophila miran
>gi|1037078597|ref|XP_017143107.1| PREDICTED: nucleotide exchange factor SIL1 isoform X1 [Dros
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>gi|1037078611|ref|XP_017143108.1| PREDICTED: nucleotide exchange factor SIL1 isoform X2 [Drosc
>gi|1037078625|ref|XP_017143109.1| PREDICTED: nucleotide exchange factor SIL1 isoform X3 [Drosc
>gi|1037103249|ref|XP_017145063.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037103239|ref|XP_017145061.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037101047|ref|XP_017144844.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037101037|ref|XP_017144843.1| PREDICTED: protein bag-of-marbles [Drosophila miranda]
>gi|1037093952|ref|XP_017144224.1| PREDICTED: 40S ribosomal protein S27 [Drosophila miranda]
>gi|1037093938|ref|XP_017144223.1| PREDICTED: nucleoporin Nup37 [Drosophila miranda]
>gi|1037047148|ref|XP_017140966.1| PREDICTED: E3 SUMO-protein ligase RanBP2 [Drosophila miranda
>gi|1037047194|ref|XP_017140969.1| PREDICTED: LOW QUALITY PROTEIN: transcription factor Ouib-li
>gi|1037047180|ref|XP_017140968.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037047210|ref|XP_017140970.1| PREDICTED: HIG1 domain family member 1A, mitochondrial [Dros
>gi|1037047164|ref|XP_017140967.1| PREDICTED: dihydrolipoyllysine-residue succinyltransferase
>gi|1037067809|ref|XP_017142362.1| PREDICTED: serine/threonine-protein kinase MARK2 [Drosophila
>gi|1037067824|ref|XP_017142363.1| PREDICTED: serine/threonine-protein kinase MARK2 [Drosophila
>gi|1037067707|ref|XP_017142355.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1037067723|ref|XP_017142356.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1037067738|ref|XP_017142357.1| PREDICTED: homeobox protein prospero isoform X1 [Drosophila
>gi|1037067753|ref|XP_017142358.1| PREDICTED: homeobox protein prospero isoform X2 [Drosophila
>gi|1037067768|ref|XP_017142359.1| PREDICTED: homeobox protein prospero isoform X3 [Drosophila
>gi|1037067783|ref|XP_017142360.1| PREDICTED: homeobox protein prospero isoform X4 [Drosophila
>gi|1037067795|ref|XP_017142361.1| PREDICTED: homeobox protein prospero isoform X5 [Drosophila
>gi|1037057958|ref|XP_017141697.1| PREDICTED: 39S ribosomal protein L40, mitochondrial [Drosoph
>gi|1037057942|ref|XP_017141696.1| PREDICTED: prefoldin subunit 3 [Drosophila miranda]
>gi|1037057892|ref|XP_017141692.1| PREDICTED: ATP-dependent DNA helicase 2 subunit 1 [Drosophil
>gi|1037057876|ref|XP_017141691.1| PREDICTED: importin-9 [Drosophila miranda]
>gi|1037057926|ref|XP_017141694.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 [Dro
>gi|1037057910|ref|XP_017141693.1| PREDICTED: neither inactivation nor afterpotential protein (
>gi|1037058003|ref|XP_017141700.1| PREDICTED: D-tyrosyl-tRNA(Tyr) deacylase 1 isoform X2 [Drosc
>gi|1037057987|ref|XP_017141699.1| PREDICTED: D-tyrosyl-tRNA(Tyr) deacylase 1 isoform X1 [Drosc
>gi|1037057972|ref|XP_017141698.1| PREDICTED: probable ribosome biogenesis protein RLP24 [Drosc
>gi|1037058789|ref|XP_017141755.1| PREDICTED: ubiquitin carboxyl-terminal hydrolase 8 [Drosophi
>gi|1037058839|ref|XP_017141758.1| PREDICTED: putative tRNA (cytidine(32)/guanosine(34)-2'-O)-r
>gi|1037058823|ref|XP_017141757.1| PREDICTED: uncharacterized protein LOC108155452 [Drosophila
>gi|1037058853|ref|XP_017141759.1| PREDICTED: ribosome production factor 2 homolog [Drosophila
>gi|1037058884|ref|XP_017141761.1| PREDICTED: uncharacterized protein LOC108155456 isoform X2
>gi|1037058869|ref|XP_017141760.1| PREDICTED: uncharacterized protein LOC108155456 isoform X1
>gi|1037058900|ref|XP_017141762.1| PREDICTED: uncharacterized protein LOC108155456 isoform X3
>gi|1037058807|ref|XP_017141756.1| PREDICTED: ralA-binding protein 1 [Drosophila miranda]
>gi|1037022693|ref|XP_017139438.1| PREDICTED: LOW QUALITY PROTEIN: malate dehydrogenase, mitoch
>gi|1037068859|ref|XP_017142436.1| PREDICTED: retinal homeobox protein Rx [Drosophila miranda]
>gi|1037068829|ref|XP_017142434.1| PREDICTED: ribosomal protein S6 kinase delta-1 [Drosophila n
>gi|1037068845|ref|XP_017142435.1| PREDICTED: little elongation complex subunit 2 [Drosophila n
>gi|1037068904|ref|XP_017142439.1| PREDICTED: SRR1-like protein [Drosophila miranda]
>gi|1037068873|ref|XP_017142437.1| PREDICTED: coiled-coil domain-containing protein 174 [Drosop
>gi|1037068889|ref|XP_017142438.1| PREDICTED: uncharacterized protein LOC108155838 [Drosophila
>gi|1037104463|ref|XP_017145217.1| PREDICTED: hexosaminidase D [Drosophila miranda]
>gi|1037082963|ref|XP_017143428.1| PREDICTED: single-stranded DNA-binding protein 2 [Drosophila
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>gi|1037062830|ref|XP_017142035.1| PREDICTED: uncharacterized protein LOC108155617 [Drosophila
>gi|1037062786|ref|XP_017142032.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1037062816|ref|XP_017142034.1| PREDICTED: proteasome assembly chaperone 2 [Drosophila mira
>gi|1037062770|ref|XP_017142031.1| PREDICTED: actin-related protein 5 [Drosophila miranda]
>gi|1037062801|ref|XP_017142033.1| PREDICTED: uncharacterized protein LOC108155615 [Drosophila
>gi|1037062754|ref|XP_017142030.1| PREDICTED: importin-13 [Drosophila miranda]
>gi|1037067872|ref|XP_017142366.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037067856|ref|XP_017142365.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037067840|ref|XP_017142364.1| PREDICTED: serine/threonine-protein phosphatase 2A 56 kDa re
>gi|1037067920|ref|XP_017142370.1| PREDICTED: ubiquitin-like protein 4A [Drosophila miranda]
>gi|1037067888|ref|XP_017142367.1| PREDICTED: peroxiredoxin-5, mitochondrial [Drosophila miranc
>gi|1037067904|ref|XP_017142369.1| PREDICTED: nuclear cap-binding protein subunit 2 [Drosophila
>gi|1037094430|ref|XP_017144260.1| PREDICTED: protein TAPT1 homolog [Drosophila miranda]
>gi|1037103985|ref|XP_017145153.1| PREDICTED: uncharacterized protein LOC108157558 [Drosophila
>gi|1037103969|ref|XP_017145151.1| PREDICTED: uncharacterized protein LOC108157557 isoform X2
>gi|1037103977|ref|XP_017145152.1| PREDICTED: uncharacterized protein LOC108157557 isoform X3
>gi|1037103961|ref|XP_017145150.1| PREDICTED: uncharacterized protein LOC108157557 isoform X1
>gi|1037033543|ref|XP_017140103.1| PREDICTED: uncharacterized protein LOC108154348 [Drosophila
>gi|1037098435|ref|XP_017144615.1| PREDICTED: hormone receptor 4 isoform X2 [Drosophila miranda
>gi|1037098425|ref|XP_017144614.1| PREDICTED: hormone receptor 4 isoform X1 [Drosophila miranda
>gi|1037098447|ref|XP_017144616.1| PREDICTED: alpha-protein kinase 1 isoform X3 [Drosophila mi
>gi|1037099902|ref|XP_017144744.1| PREDICTED: uncharacterized protein LOC108157267 [Drosophila
>gi|1037086058|ref|XP_017143660.1| PREDICTED: fibroblast growth factor receptor homolog 1 [Dros
>gi|1037086072|ref|XP_017143661.1| PREDICTED: fibroblast growth factor receptor homolog 1 [Dros
>gi|1037046824|ref|XP_017140944.1| PREDICTED: tubulin alpha-3 chain [Drosophila miranda]
>gi|1037046854|ref|XP_017140947.1| PREDICTED: SPRY domain-containing protein 7 [Drosophila mira
>gi|1037046840|ref|XP_017140945.1| PREDICTED: plancitoxin-1 [Drosophila miranda]
>gi|1037022711|ref|XP_017139439.1| PREDICTED: LOW QUALITY PROTEIN: protein couch potato, partia
>gi|1037046729|ref|XP_017140938.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046808|ref|XP_017140943.1| PREDICTED: uncharacterized protein LOC108154948 isoform X9
>gi|1037046713|ref|XP_017140936.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046745|ref|XP_017140939.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046761|ref|XP_017140940.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046681|ref|XP_017140934.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046697|ref|XP_017140935.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046776|ref|XP_017140941.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046792|ref|XP_017140942.1| PREDICTED: regulating synaptic membrane exocytosis protein 2
>gi|1037046868|ref|XP_017140948.1| PREDICTED: uncharacterized protein LOC108154952 [Drosophila
>gi|1037030467|ref|XP_017139919.1| PREDICTED: protein tincar isoform X1 [Drosophila miranda]
>gi|1037030483|ref|XP_017139920.1| PREDICTED: protein tincar isoform X2 [Drosophila miranda]
>gi|1037030517|ref|XP_017139923.1| PREDICTED: chitobiosyldiphosphodolichol beta-mannosyltransfe
>gi|1037030743|ref|XP_017139938.1| PREDICTED: uncharacterized protein LOC108154222 [Drosophila
>gi|1037030653|ref|XP_017139932.1| PREDICTED: arfaptin-2 [Drosophila miranda]
>gi|1037030781|ref|XP_017139940.1| PREDICTED: DPH3 homolog [Drosophila miranda]
>gi|1037030686|ref|XP_017139934.1| PREDICTED: dnaJ homolog subfamily C member 17 [Drosophila mi
>gi|1037030501|ref|XP_017139921.1| PREDICTED: GRIP and coiled-coil domain-containing protein 1
>gi|1037030533|ref|XP_017139924.1| PREDICTED: NADH dehydrogenase [ubiquinone] complex I, assemb
>gi|1037030707|ref|XP_017139935.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
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>gi|1037030725|ref|XP_017139936.1| PREDICTED: venom allergen 3-like [Drosophila miranda]
>gi|1037030548|ref|XP_017139925.1| PREDICTED: 60S ribosomal protein L3 isoform X1 [Drosophila r
>gi|1037030566|ref|XP_017139926.1| PREDICTED: 60S ribosomal protein L3 isoform X2 [Drosophila r
>gi|1037030599|ref|XP_017139928.1| PREDICTED: zinc finger protein 544-like [Drosophila miranda]
>gi|1037030671|ref|XP_017139933.1| PREDICTED: J domain-containing protein CG6693 [Drosophila mi
>gi|1037030617|ref|XP_017139929.1| PREDICTED: uncharacterized protein LOC108154215 [Drosophila
>gi|1037030582|ref|XP_017139927.1| PREDICTED: uncharacterized protein LOC108154213 [Drosophila
>gi|1037022729|ref|XP_017139440.1| PREDICTED: uncharacterized protein LOC108153853 [Drosophila
>gi|1037030638|ref|XP_017139931.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037030763|ref|XP_017139939.1| PREDICTED: 40S ribosomal protein S25 [Drosophila miranda]
>gi|1037094813|ref|XP_017144288.1| PREDICTED: zinc transporter 7 isoform X1 [Drosophila miranda
>gi|1037094827|ref|XP_017144289.1| PREDICTED: zinc transporter 7 isoform X2 [Drosophila miranda
>gi|1037094841|ref|XP_017144291.1| PREDICTED: zinc transporter 7 isoform X2 [Drosophila miranda
>gi|1037096743|ref|XP_017144452.1| PREDICTED: translationally-controlled tumor protein homolog
>gi|1037089986|ref|XP_017143937.1| PREDICTED: succinate dehydrogenase cytochrome b560 subunit,
>gi|1037089972|ref|XP_017143935.1| PREDICTED: nocturnin isoform X5 [Drosophila miranda]
>gi|1037089918|ref|XP_017143931.1| PREDICTED: nocturnin isoform X1 [Drosophila miranda]
>gi|1037089958|ref|XP_017143934.1| PREDICTED: nocturnin isoform X4 [Drosophila miranda]
>gi|1037089931|ref|XP_017143932.1| PREDICTED: nocturnin isoform X2 [Drosophila miranda]
>gi|1037089944|ref|XP_017143933.1| PREDICTED: nocturnin isoform X3 [Drosophila miranda]
>gi|1037090000|ref|XP_017143938.1| PREDICTED: uncharacterized protein LOC108156777 [Drosophila
>gi|1037071025|ref|XP_017142586.1| PREDICTED: UDP-glucuronosyltransferase 2C1 isoform X2 [Droso
>gi|1037071009|ref|XP_017142585.1| PREDICTED: UDP-glucuronosyltransferase 1-9 isoform X1 [Droso
>gi|1037071059|ref|XP_017142589.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1037070957|ref|XP_017142582.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1037071043|ref|XP_017142587.1| PREDICTED: UDP-glucuronosyltransferase 2A3 [Drosophila mirar
>gi|1037070993|ref|XP_017142584.1| PREDICTED: UDP-glucuronosyltransferase 2A3-like [Drosophila
>gi|1037070941|ref|XP_017142581.1| PREDICTED: venom carboxylesterase-6 [Drosophila miranda]
>gi|1037070975|ref|XP_017142583.1| PREDICTED: UDP-glucuronosyltransferase [Drosophila miranda]
>gi|1037102241|ref|XP_017144957.1| PREDICTED: UDP-glucuronosyltransferase 2B15-like [Drosophila
>gi|1037106057|ref|XP_017145444.1| PREDICTED: LOW QUALITY PROTEIN: UDP-glucuronosyltransferase
>gi|1037105360|ref|XP_017145347.1| PREDICTED: UDP-glucuronosyltransferase 2B31 [Drosophila mira
>gi|1037103921|ref|XP_017145145.1| PREDICTED: probable aconitate hydratase, mitochondrial [Dros
>gi|1037105260|ref|XP_017145331.1| PREDICTED: UDP-glucuronosyltransferase 2B15 [Drosophila mira
>gi|1037033526|ref|XP_017140102.1| PREDICTED: uncharacterized protein LOC108154347 [Drosophila
>gi|1037085126|ref|XP_017143595.1| PREDICTED: nephrin [Drosophila miranda]
>gi|1037085179|ref|XP_017143599.1| PREDICTED: uncharacterized protein LOC108156576 [Drosophila
>gi|1037085153|ref|XP_017143597.1| PREDICTED: uncharacterized protein LOC108156575 isoform X1
>gi|1037085167|ref|XP_017143598.1| PREDICTED: uncharacterized protein LOC108156575 isoform X2
>gi|1037085207|ref|XP_017143602.1| PREDICTED: uncharacterized protein LOC108156578 [Drosophila
>gi|1037085193|ref|XP_017143601.1| PREDICTED: uncharacterized protein LOC108156577 [Drosophila
>gi|1037085140|ref|XP_017143596.1| PREDICTED: proteasome subunit alpha type-6-like [Drosophila
>gi|1037074191|ref|XP_017142804.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037074175|ref|XP_017142803.1| PREDICTED: tetratricopeptide repeat protein 14 homolog [Dros
>gi|1037074206|ref|XP_017142805.1| PREDICTED: GTP:AMP phosphotransferase AK3, mitochondrial [Dr
>gi|1037074220|ref|XP_017142806.1| PREDICTED: GTP:AMP phosphotransferase AK3, mitochondrial [Dr
>gi|1037095949|ref|XP_017144378.1| PREDICTED: arrestin domain-containing protein 1 [Drosophila
>gi|1037095961|ref|XP_017144379.1| PREDICTED: mitochondrial thiamine pyrophosphate carrier [Dro
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>gi|1037099638|ref|XP_017144720.1| PREDICTED: uncharacterized protein LOC108157253 [Drosophila
>gi|1037099626|ref|XP_017144719.1| PREDICTED: sorbitol dehydrogenase isoform X3 [Drosophila mi
>gi|1037099614|ref|XP_017144718.1| PREDICTED: sorbitol dehydrogenase isoform X2 [Drosophila mi
>gi|1037099602|ref|XP_017144717.1| PREDICTED: sorbitol dehydrogenase isoform X1 [Drosophila mi
>gi|1037105340|ref|XP_017145343.1| PREDICTED: uncharacterized protein LOC108157687 [Drosophila
>gi|1037062657|ref|XP_017142023.1| PREDICTED: alcohol dehydrogenase class-3 [Drosophila miranda
>gi|1037062673|ref|XP_017142024.1| PREDICTED: tetraspanin-33 [Drosophila miranda]
>gi|1037062687|ref|XP_017142025.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X1 [
>gi|1037062705|ref|XP_017142026.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X2 [
>gi|1037062721|ref|XP_017142028.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X3 [
>gi|1037062736|ref|XP_017142029.1| PREDICTED: methionine-R-sulfoxide reductase B1 isoform X4 [
>gi|1037062596|ref|XP_017142019.1| PREDICTED: thiamine transporter 1 isoform X1 [Drosophila mi
>gi|1037062612|ref|XP_017142020.1| PREDICTED: thiamine transporter 1 isoform X2 [Drosophila mi
>gi|1037062627|ref|XP_017142021.1| PREDICTED: thiamine transporter 2 isoform X1 [Drosophila mi
>gi|1037062641|ref|XP_017142022.1| PREDICTED: thiamine transporter 2 isoform X2 [Drosophila mi
>gi|1037074797|ref|XP_017142845.1| PREDICTED: uncharacterized protein LOC108156072 [Drosophila
>gi|1037074781|ref|XP_017142844.1| PREDICTED: protein AF-9 [Drosophila miranda]
>gi|1037074839|ref|XP_017142849.1| PREDICTED: uncharacterized protein LOC108156074 [Drosophila
>gi|1037074823|ref|XP_017142846.1| PREDICTED: nuclear transcription factor Y subunit beta [Dros
>gi|1037103175|ref|XP_017145055.1| PREDICTED: tropomyosin-2 isoform X1 [Drosophila miranda]
>gi|1037103185|ref|XP_017145056.1| PREDICTED: tropomyosin-2 isoform X2 [Drosophila miranda]
>gi|1037055526|ref|XP_017141539.1| PREDICTED: golgin-84 isoform X11 [Drosophila miranda]
>gi|1037055509|ref|XP_017141538.1| PREDICTED: golgin-84 isoform X10 [Drosophila miranda]
>gi|1037055649|ref|XP_017141546.1| PREDICTED: tropomyosin-1, isoforms 9A/A/B isoform X18 [Drosc
>gi|1037055665|ref|XP_017141547.1| PREDICTED: tropomyosin-1, isoforms 9A/A/B isoform X19 [Drosc
>gi|1037055584|ref|XP_017141542.1| PREDICTED: tropomyosin-2 isoform X14 [Drosophila miranda]
>gi|1037055600|ref|XP_017141543.1| PREDICTED: tropomyosin-2 isoform X15 [Drosophila miranda]
>gi|1037055465|ref|XP_017141534.1| PREDICTED: golgin-84 isoform X7 [Drosophila miranda]
>gi|1037055451|ref|XP_017141533.1| PREDICTED: golgin-84 isoform X6 [Drosophila miranda]
>gi|1037055383|ref|XP_017141529.1| PREDICTED: formin-binding protein 4 isoform X2 [Drosophila r
>gi|1037055367|ref|XP_017141528.1| PREDICTED: formin-binding protein 4 isoform X1 [Drosophila r
>gi|1037055399|ref|XP_017141530.1| PREDICTED: formin-binding protein 4 isoform X3 [Drosophila r
>gi|1037055568|ref|XP_017141541.1| PREDICTED: tropomyosin-1, isoforms 33/34 isoform X13 [Drosop
>gi|1037055543|ref|XP_017141540.1| PREDICTED: tropomyosin-2 isoform X12 [Drosophila miranda]
>gi|1037055415|ref|XP_017141531.1| PREDICTED: actin cytoskeleton-regulatory complex protein pan
>gi|1037055435|ref|XP_017141532.1| PREDICTED: actin cytoskeleton-regulatory complex protein pan
>gi|1037055614|ref|XP_017141544.1| PREDICTED: tropomyosin-1, isoforms 33/34 isoform X16 [Drosop
>gi|1037055492|ref|XP_017141537.1| PREDICTED: golgin-84 isoform X9 [Drosophila miranda]
>gi|1037055476|ref|XP_017141536.1| PREDICTED: golgin-84 isoform X8 [Drosophila miranda]
>gi|1037055633|ref|XP_017141545.1| PREDICTED: tropomyosin-1, isoforms 9A/A/B isoform X17 [Drosc
>gi|1037043237|ref|XP_017140706.1| PREDICTED: cytochrome c oxidase assembly protein COX16 homol
>gi|1037043147|ref|XP_017140700.1| PREDICTED: nuA4 complex subunit EAF3 homolog [Drosophila mi
>gi|1037043212|ref|XP_017140704.1| PREDICTED: ribosome biogenesis protein TSR3 homolog [Drosoph
>gi|1037043055|ref|XP_017140693.1| PREDICTED: centrosome-associated zinc finger protein CP190
>gi|1037043069|ref|XP_017140694.1| PREDICTED: centrosome-associated zinc finger protein CP190
>gi|1037043085|ref|XP_017140695.1| PREDICTED: centrosome-associated zinc finger protein CP190
>gi|1037043180|ref|XP_017140702.1| PREDICTED: protein zntD isoform X2 [Drosophila miranda]
>gi|1037043164|ref|XP_017140701.1| PREDICTED: protein zntD isoform X1 [Drosophila miranda]
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>gi|1037043101|ref|XP_017140696.1| PREDICTED: uncharacterized protein LOC108154808 [Drosophila
>gi|1037043224|ref|XP_017140705.1| PREDICTED: ATP synthase subunit O, mitochondrial [Drosophila
>gi|1037043196|ref|XP_017140703.1| PREDICTED: protein SET [Drosophila miranda]
>gi|1037043132|ref|XP_017140698.1| PREDICTED: eukaryotic translation initiation factor 2 subuni
>gi|1037043117|ref|XP_017140697.1| PREDICTED: histone-lysine N-methyltransferase Su(var)3-9 iso
>gi|1037043252|ref|XP_017140707.1| PREDICTED: AF4/FMR2 family member 4 [Drosophila miranda]
>gi|1037043264|ref|XP_017140708.1| PREDICTED: uncharacterized protein LOC108154819 [Drosophila
>gi|1037081134|ref|XP_017143291.1| PREDICTED: putative ammonium transporter 2 [Drosophila miran
>gi|1037081120|ref|XP_017143289.1| PREDICTED: heat shock 70 kDa protein cognate 4 [Drosophila r
>gi|1037052045|ref|XP_017141299.1| PREDICTED: serine/threonine-protein kinase ATM [Drosophila r
>gi|1037052138|ref|XP_017141304.1| PREDICTED: uncharacterized protein LOC108155155 [Drosophila
>gi|1037052154|ref|XP_017141306.1| PREDICTED: 28S ribosomal protein S10, mitochondrial isoform
>gi|1037052170|ref|XP_017141307.1| PREDICTED: 28S ribosomal protein S10, mitochondrial isoform
>gi|1037052074|ref|XP_017141300.1| PREDICTED: protein arginine N-methyltransferase 1 isoform X1
>gi|1037052090|ref|XP_017141301.1| PREDICTED: protein arginine N-methyltransferase 1 isoform X2
>gi|1037052106|ref|XP_017141302.1| PREDICTED: probable histone-binding protein Caf1 [Drosophila
>gi|1037052122|ref|XP_017141303.1| PREDICTED: uncharacterized protein LOC108155154 [Drosophila
>gi|1037089715|ref|XP_017143914.1| PREDICTED: rho GTPase-activating protein 92B [Drosophila mir
>gi|1037106167|ref|XP_017145459.1| PREDICTED: uncharacterized protein LOC108157781 [Drosophila
>gi|1037106159|ref|XP_017145458.1| PREDICTED: uncharacterized protein LOC108157779 [Drosophila
>gi|1037034204|ref|XP_017140143.1| PREDICTED: solute carrier family 22 member 3 [Drosophila mir
>gi|1037098529|ref|XP_017144623.1| PREDICTED: putative fatty acyl-CoA reductase CG5065 [Drosopl
>gi|1037098505|ref|XP_017144621.1| PREDICTED: solute carrier family 22 member 13 [Drosophila m:
>gi|1037098517|ref|XP_017144622.1| PREDICTED: solute carrier family 22 member 13 [Drosophila m:
>gi|1037098495|ref|XP_017144620.1| PREDICTED: solute carrier family 22 member 1 [Drosophila mir
>gi|1037071205|ref|XP_017142599.1| PREDICTED: heparan-sulfate 6-O-sulfotransferase 2 [Drosophil
>gi|1037071189|ref|XP_017142598.1| PREDICTED: FYVE and coiled-coil domain-containing protein 1
>gi|1037071171|ref|XP_017142597.1| PREDICTED: E3 ubiquitin-protein ligase BRE1 isoform X1 [Dros
>gi|1037071236|ref|XP_017142600.1| PREDICTED: uncharacterized protein LOC108155946 [Drosophila
>gi|1037071268|ref|XP_017142602.1| PREDICTED: uncharacterized protein LOC108155949 [Drosophila
>gi|1037071252|ref|XP_017142601.1| PREDICTED: uncharacterized protein LOC108155948 [Drosophila
>gi|1037033234|ref|XP_017140086.1| PREDICTED: uncharacterized protein LOC108154336 [Drosophila
>gi|1037060596|ref|XP_017141883.1| PREDICTED: uncharacterized protein LOC108155531 isoform X2
>gi|1037060582|ref|XP_017141882.1| PREDICTED: uncharacterized protein LOC108155531 isoform X1
>gi|1037060610|ref|XP_017141884.1| PREDICTED: uncharacterized protein LOC108155531 isoform X3
>gi|1037060626|ref|XP_017141886.1| PREDICTED: uncharacterized protein LOC108155531 isoform X4
>gi|1037060642|ref|XP_017141887.1| PREDICTED: uncharacterized protein LOC108155531 isoform X5
>gi|1037060706|ref|XP_017141891.1| PREDICTED: N-acetylglucosaminyl-phosphatidylinositol de-N-ac
>gi|1037060658|ref|XP_017141888.1| PREDICTED: phagocyte signaling-impaired protein isoform X1
>gi|1037060672|ref|XP_017141889.1| PREDICTED: phagocyte signaling-impaired protein isoform X2
>gi|1037060736|ref|XP_017141894.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037060752|ref|XP_017141895.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037060720|ref|XP_017141892.1| PREDICTED: zinc finger protein 287 [Drosophila miranda]
>gi|1037060690|ref|XP_017141890.1| PREDICTED: zinc finger protein 62 homolog [Drosophila miranc
>gi|1037090709|ref|XP_017143995.1| PREDICTED: zinc finger protein 771 [Drosophila miranda]
>gi|1037090695|ref|XP_017143994.1| PREDICTED: probable ribonuclease ZC3H12C [Drosophila miranda
>gi|1037103772|ref|XP_017145126.1| PREDICTED: uncharacterized protein LOC108157534 [Drosophila
>gi|1037103748|ref|XP_017145122.1| PREDICTED: protein I'm not dead yet 2-like [Drosophila miran
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>gi|1037103756|ref|XP_017145123.1| PREDICTED: protein I'm not dead yet 2-like [Drosophila mira
>gi|1037103764|ref|XP_017145125.1| PREDICTED: uncharacterized protein LOC108157532 [Drosophila
>gi|1037095312|ref|XP_017144328.1| PREDICTED: S-formylglutathione hydrolase [Drosophila miranda
>gi|1037095326|ref|XP_017144329.1| PREDICTED: S-formylglutathione hydrolase [Drosophila miranda
>gi|1037095341|ref|XP_017144330.1| PREDICTED: S-formylglutathione hydrolase [Drosophila miranda
>gi|1037095298|ref|XP_017144327.1| PREDICTED: RING finger protein 113A [Drosophila miranda]
>gi|1037104656|ref|XP_017145244.1| PREDICTED: opsin Rh3 [Drosophila miranda]
>gi|1037097219|ref|XP_017144495.1| PREDICTED: heat shock 70 kDa protein cognate 4-like [Drosoph
>gi|1037105251|ref|XP_017145330.1| PREDICTED: muscle-specific protein 20 [Drosophila miranda]
>gi|1037103359|ref|XP_017145075.1| PREDICTED: ureidoglycolate dehydrogenase (NAD(+)) [Drosophil
>gi|1037103339|ref|XP_017145073.1| PREDICTED: neuroligin 4-like [Drosophila miranda]
>gi|1037103349|ref|XP_017145074.1| PREDICTED: neuroligin 4-like [Drosophila miranda]
>gi|1037066448|ref|XP_017142272.1| PREDICTED: putative transcription factor capicua isoform X1
>gi|1037066496|ref|XP_017142275.1| PREDICTED: putative transcription factor capicua isoform X4
>gi|1037066464|ref|XP_017142273.1| PREDICTED: putative transcription factor capicua isoform X2
>gi|1037066511|ref|XP_017142276.1| PREDICTED: putative transcription factor capicua isoform X5
>gi|1037066480|ref|XP_017142274.1| PREDICTED: putative transcription factor capicua isoform X3
>gi|1037066529|ref|XP_017142277.1| PREDICTED: putative transcription factor capicua isoform X6
>gi|1037066561|ref|XP_017142279.1| PREDICTED: putative transcription factor capicua isoform X8
>gi|1037066546|ref|XP_017142278.1| PREDICTED: putative transcription factor capicua isoform X7
>gi|1037037294|ref|XP_017140318.1| PREDICTED: uncharacterized protein LOC108154526 [Drosophila
>gi|1037099154|ref|XP_017144674.1| PREDICTED: uncharacterized protein LOC108157228 [Drosophila
>gi|1037099087|ref|XP_017144672.1| PREDICTED: titin homolog [Drosophila miranda]
>gi|1037099105|ref|XP_017144673.1| PREDICTED: uncharacterized family 31 glucosidase KIAA1161 [D
>gi|1037049314|ref|XP_017141109.1| PREDICTED: furin-like protease 1 isoform X2 [Drosophila mira
>gi|1037049284|ref|XP_017141107.1| PREDICTED: furin-like protease 1 isoform X1 [Drosophila mira
>gi|1037049300|ref|XP_017141108.1| PREDICTED: furin-like protease 1 isoform X1 [Drosophila mira
>gi|1037049328|ref|XP_017141110.1| PREDICTED: NADH-quinone oxidoreductase subunit D [Drosophila
>gi|1037049347|ref|XP_017141112.1| PREDICTED: carboxypeptidase N subunit 2 [Drosophila miranda]
>gi|1037049361|ref|XP_017141113.1| PREDICTED: collagen alpha-1(V) chain [Drosophila miranda]
>gi|1037049374|ref|XP_017141114.1| PREDICTED: protein translation factor SUI1 homolog [Drosoph
>gi|1037094349|ref|XP_017144253.1| PREDICTED: vacuolar protein sorting-associated protein 33B
>gi|1037094363|ref|XP_017144254.1| PREDICTED: vacuolar protein sorting-associated protein 33B
>gi|1037094376|ref|XP_017144255.1| PREDICTED: uncharacterized protein LOC108156975 [Drosophila
>gi|1037031400|ref|XP_017139977.1| PREDICTED: mediator of RNA polymerase II transcription subu
>gi|1037106628|ref|XP_017145523.1| PREDICTED: regulator of chromosome condensation-like [Drosop
>gi|1037057418|ref|XP_017141659.1| PREDICTED: transcriptional regulator ATRX homolog [Drosophil
>gi|1037057402|ref|XP_017141658.1| PREDICTED: nuclear factor related to kappa-B-binding protei
>gi|1037094855|ref|XP_017144292.1| PREDICTED: zinc finger protein 423 [Drosophila miranda]
>gi|1037091031|ref|XP_017144019.1| PREDICTED: WD repeat domain phosphoinositide-interacting pro
>gi|1037090949|ref|XP_017144012.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037090963|ref|XP_017144013.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037090976|ref|XP_017144014.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037090990|ref|XP_017144015.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037091004|ref|XP_017144016.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037091018|ref|XP_017144017.1| PREDICTED: facilitated trehalose transporter Tret1 isoform
>gi|1037091047|ref|XP_017144020.1| PREDICTED: venom allergen 3 [Drosophila miranda]
>gi|1037053318|ref|XP_017141385.1| PREDICTED: UPF0160 protein [Drosophila miranda]
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>gi|1037053356|ref|XP_017141386.1| PREDICTED: proteasome subunit beta type-3 [Drosophila mirand
>gi|1037053304|ref|XP_017141384.1| PREDICTED: E3 ubiquitin-protein ligase RNF126-B [Drosophila
>gi|1037053276|ref|XP_017141382.1| PREDICTED: uncharacterized protein LOC108155209 [Drosophila
>gi|1037053250|ref|XP_017141380.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X2 [Dro
>gi|1037053264|ref|XP_017141381.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X3 [Dro
>gi|1037053202|ref|XP_017141377.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1037053218|ref|XP_017141378.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1037053234|ref|XP_017141379.1| PREDICTED: E3 ubiquitin-protein ligase KCMF1 isoform X1 [Dro
>gi|1037053288|ref|XP_017141383.1| PREDICTED: embryonic stem cell-specific 5-hydroxymethylcytos
>gi|1037053176|ref|XP_017141374.1| PREDICTED: aryl hydrocarbon receptor nuclear translocator ho
>gi|1037053196|ref|XP_017141376.1| PREDICTED: aryl hydrocarbon receptor nuclear translocator ho
>gi|1037076200|ref|XP_017142943.1| PREDICTED: protein neuralized isoform X2 [Drosophila miranda
>gi|1037076184|ref|XP_017142942.1| PREDICTED: protein neuralized isoform X1 [Drosophila miranda
>gi|1037065545|ref|XP_017142211.1| PREDICTED: parafibromin [Drosophila miranda]
>gi|1037065577|ref|XP_017142213.1| PREDICTED: bifunctional methylenetetrahydrofolate dehydroger
>gi|1037065561|ref|XP_017142212.1| PREDICTED: bifunctional methylenetetrahydrofolate dehydroger
>gi|1037065593|ref|XP_017142214.1| PREDICTED: bifunctional methylenetetrahydrofolate dehydroger
>gi|1037065527|ref|XP_017142210.1| PREDICTED: nuclear factor NF-kappa-B p110 subunit [Drosophil
>gi|1037065609|ref|XP_017142215.1| PREDICTED: uncharacterized protein LOC108155722 [Drosophila
>gi|1037077240|ref|XP_017143010.1| PREDICTED: putative protein TPRXL [Drosophila miranda]
>gi|1037072769|ref|XP_017142705.1| PREDICTED: uncharacterized protein LOC108156011 [Drosophila
>gi|1037072753|ref|XP_017142703.1| PREDICTED: uncharacterized protein LOC108156010 [Drosophila
>gi|1037072737|ref|XP_017142702.1| PREDICTED: T-complex protein 1 subunit eta [Drosophila miran
>gi|1037072722|ref|XP_017142701.1| PREDICTED: uncharacterized protein LOC108156008 [Drosophila
>gi|1037070676|ref|XP_017142562.1| PREDICTED: cytospin-A isoform X1 [Drosophila miranda]
>gi|1037070690|ref|XP_017142563.1| PREDICTED: cytospin-A isoform X2 [Drosophila miranda]
>gi|1037070765|ref|XP_017142569.1| PREDICTED: uncharacterized protein LOC108155927 [Drosophila
>gi|1037070705|ref|XP_017142564.1| PREDICTED: intracellular protein transport protein USO1 isof
>gi|1037070733|ref|XP_017142566.1| PREDICTED: uncharacterized protein LOC108155925 [Drosophila
>gi|1037070719|ref|XP_017142565.1| PREDICTED: uncharacterized protein LOC108155924 [Drosophila
>gi|1037022747|ref|XP_017139441.1| PREDICTED: uncharacterized protein LOC108153854 [Drosophila
>gi|1037070749|ref|XP_017142568.1| PREDICTED: uncharacterized protein LOC108155926 [Drosophila
>gi|1037105227|ref|XP_017145327.1| PREDICTED: chymotrypsinogen A isoform X1 [Drosophila miranda
>gi|1037105235|ref|XP_017145328.1| PREDICTED: chymotrypsinogen A isoform X2 [Drosophila miranda
>gi|1037105219|ref|XP_017145326.1| PREDICTED: uncharacterized protein LOC108157673 [Drosophila
>gi|1037090668|ref|XP_017143991.1| PREDICTED: uncharacterized protein LOC108156809 [Drosophila
>gi|1037090682|ref|XP_017143992.1| PREDICTED: protein NASP homolog [Drosophila miranda]
>gi|1037048525|ref|XP_017141055.1| PREDICTED: PP2C-like domain-containing protein CG9801 [Droso
>gi|1037048509|ref|XP_017141054.1| PREDICTED: UPF0505 protein CG8202 isoform X2 [Drosophila mir
>gi|1037048496|ref|XP_017141053.1| PREDICTED: UPF0505 protein CG8202 isoform X1 [Drosophila mir
>gi|1037048573|ref|XP_017141058.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037048589|ref|XP_017141059.1| PREDICTED: transcription factor Ouib [Drosophila miranda]
>gi|1037048667|ref|XP_017141065.1| PREDICTED: transcription factor Ouib-like [Drosophila mirand
>gi|1037048651|ref|XP_017141063.1| PREDICTED: transcription factor Ouib isoform X4 [Drosophila
>gi|1037048619|ref|XP_017141061.1| PREDICTED: transcription factor Ouib isoform X2 [Drosophila
>gi|1037048635|ref|XP_017141062.1| PREDICTED: transcription factor Ouib isoform X3 [Drosophila
>gi|1037048605|ref|XP_017141060.1| PREDICTED: transcription factor Ouib isoform X1 [Drosophila
>gi|1037048557|ref|XP_017141057.1| PREDICTED: uncharacterized protein LOC108155016 [Drosophila
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>gi|1037048683|ref|XP_017141066.1| PREDICTED: transmembrane protein 216 [Drosophila miranda]
>gi|1037048699|ref|XP_017141067.1| PREDICTED: uncharacterized protein LOC108155023 [Drosophila
>gi|1037048715|ref|XP_017141068.1| PREDICTED: cyclin-dependent kinases regulatory subunit [Dros
>gi|1037048541|ref|XP_017141056.1| PREDICTED: sterol O-acyltransferase 1 [Drosophila miranda]
>gi|1037088751|ref|XP_017143837.1| PREDICTED: protein hunchback [Drosophila miranda]
>gi|1037088761|ref|XP_017143838.1| PREDICTED: protein hunchback [Drosophila miranda]
>gi|1037044068|ref|XP_017140757.1| PREDICTED: nuclear RNA export factor 2-like isoform X2 [Dros
>gi|1037044053|ref|XP_017140756.1| PREDICTED: nuclear RNA export factor 2-like isoform X1 [Dros
>gi|1037044084|ref|XP_017140758.1| PREDICTED: nuclear RNA export factor 2-like isoform X3 [Dros
>gi|1037044207|ref|XP_017140766.1| PREDICTED: protein unc-50 homolog [Drosophila miranda]
>gi|1037044173|ref|XP_017140764.1| PREDICTED: 39S ribosomal protein L19, mitochondrial [Drosoph
>gi|1037044191|ref|XP_017140765.1| PREDICTED: 39S ribosomal protein L19, mitochondrial [Drosoph
>gi|1037044109|ref|XP_017140760.1| PREDICTED: zinc finger protein 2 homolog isoform X2 [Drosoph
>gi|1037044125|ref|XP_017140761.1| PREDICTED: zinc finger protein 2 homolog isoform X2 [Drosoph
>gi|1037044098|ref|XP_017140759.1| PREDICTED: oocyte zinc finger protein XlCOF6-like isoform X1
>gi|1037044223|ref|XP_017140767.1| PREDICTED: putative riboflavin kinase [Drosophila miranda]
>gi|1037044141|ref|XP_017140762.1| PREDICTED: amidophosphoribosyltransferase [Drosophila mirand
>gi|1037044157|ref|XP_017140763.1| PREDICTED: transcription initiation factor TFIID subunit 7
>gi|1037044039|ref|XP_017140754.1| PREDICTED: ER membrane protein complex subunit 1 [Drosophila
>gi|1037068662|ref|XP_017142422.1| PREDICTED: 28S ribosomal protein S9, mitochondrial [Drosophi
>gi|1037068406|ref|XP_017142405.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068376|ref|XP_017142402.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068391|ref|XP_017142403.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068437|ref|XP_017142407.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068422|ref|XP_017142406.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068451|ref|XP_017142408.1| PREDICTED: transient receptor potential channel pyrexia iso
>gi|1037068646|ref|XP_017142421.1| PREDICTED: probable arginine--tRNA ligase, mitochondrial [D
>gi|1037068467|ref|XP_017142409.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X
>gi|1037068483|ref|XP_017142410.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X
>gi|1037068581|ref|XP_017142416.1| PREDICTED: uncharacterized protein LOC108155825 isoform X7
>gi|1037068515|ref|XP_017142412.1| PREDICTED: formin-J isoform X3 [Drosophila miranda]
>gi|1037068598|ref|XP_017142417.1| PREDICTED: uncharacterized protein LOC108155825 isoform X8
>gi|1037068549|ref|XP_017142414.1| PREDICTED: chromatin modification-related protein EAF1 isof
>gi|1037068498|ref|XP_017142411.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X2
>gi|1037068565|ref|XP_017142415.1| PREDICTED: chromatin modification-related protein EAF1 isof
>gi|1037068531|ref|XP_017142413.1| PREDICTED: RNA polymerase II degradation factor 1 isoform X4
>gi|1037068614|ref|XP_017142418.1| PREDICTED: uncharacterized protein LOC108155825 isoform X9
>gi|1037068630|ref|XP_017142419.1| PREDICTED: uncharacterized protein LOC108155825 isoform X10
>gi|1037101801|ref|XP_017144910.1| PREDICTED: diacylglycerol kinase eta isoform X6 [Drosophila
>gi|1037101781|ref|XP_017144908.1| PREDICTED: diacylglycerol kinase eta isoform X4 [Drosophila
>gi|1037101761|ref|XP_017144906.1| PREDICTED: diacylglycerol kinase eta isoform X2 [Drosophila
>gi|1037101811|ref|XP_017144911.1| PREDICTED: diacylglycerol kinase eta isoform X7 [Drosophila
>gi|1037101771|ref|XP_017144907.1| PREDICTED: diacylglycerol kinase eta isoform X3 [Drosophila
>gi|1037101751|ref|XP_017144905.1| PREDICTED: diacylglycerol kinase eta isoform X1 [Drosophila
>gi|1037101791|ref|XP_017144909.1| PREDICTED: diacylglycerol kinase eta isoform X5 [Drosophila
>gi|1037100400|ref|XP_017144779.1| PREDICTED: pyridoxine/pyridoxamine 5'-phosphate oxidase [Dro
>gi|1037100384|ref|XP_017144778.1| PREDICTED: pyridoxine-5'-phosphate oxidase [Drosophila miran
>gi|1037090225|ref|XP_017143956.1| PREDICTED: arrestin domain-containing protein 1 [Drosophila
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>gi|1037104913|ref|XP_017145280.1| PREDICTED: uncharacterized protein LOC108157644 [Drosophila
>gi|1037104905|ref|XP_017145279.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037100488|ref|XP_017144787.1| PREDICTED: arrestin domain-containing protein 5 [Drosophila
>gi|1037100476|ref|XP_017144786.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037100460|ref|XP_017144785.1| PREDICTED: arrestin domain-containing protein 2 [Drosophila
>gi|1037033073|ref|XP_017140076.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037033090|ref|XP_017140077.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037106440|ref|XP_017145497.1| PREDICTED: arrestin domain-containing protein 15 [Drosophila
>gi|1037106175|ref|XP_017145460.1| PREDICTED: arrestin domain-containing protein 2 [Drosophila
>gi|1037106015|ref|XP_017145439.1| PREDICTED: arrestin domain-containing protein 3 [Drosophila
>gi|1037035323|ref|XP_017140207.1| PREDICTED: uncharacterized protein LOC108154447 [Drosophila
>gi|1037105720|ref|XP_017145397.1| PREDICTED: serine protease easter isoform X1 [Drosophila mi
>gi|1037105726|ref|XP_017145398.1| PREDICTED: serine protease easter isoform X2 [Drosophila mi
>gi|1037034038|ref|XP_017140133.1| PREDICTED: uncharacterized protein LOC108154380 [Drosophila
>gi|1037034018|ref|XP_017140132.1| PREDICTED: uncharacterized protein LOC108154379 [Drosophila
>gi|1037037416|ref|XP_017140326.1| PREDICTED: uncharacterized protein LOC108154534 [Drosophila
>gi|1037033196|ref|XP_017140084.1| PREDICTED: solute carrier family 25 member 40 [Drosophila mi
>gi|1037099720|ref|XP_017144728.1| PREDICTED: uncharacterized protein LOC108157261 [Drosophila
>gi|1037099734|ref|XP_017144729.1| PREDICTED: uncharacterized protein LOC108157261 [Drosophila
>gi|1037103536|ref|XP_017145095.1| PREDICTED: cytochrome c oxidase subunit 4 isoform 1, mitocho
>gi|1037022765|ref|XP_017139443.1| PREDICTED: neuroligin-4, Y-linked [Drosophila miranda]
>gi|1037022781|ref|XP_017139444.1| PREDICTED: uncharacterized protein LOC108153856 [Drosophila
>gi|1037064400|ref|XP_017142143.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037064414|ref|XP_017142144.1| PREDICTED: esterase B1 isoform X1 [Drosophila miranda]
>gi|1037064430|ref|XP_017142145.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1037064446|ref|XP_017142146.1| PREDICTED: esterase B1-like isoform X1 [Drosophila miranda]
>gi|1037064460|ref|XP_017142148.1| PREDICTED: esterase B1-like isoform X2 [Drosophila miranda]
>gi|1037022805|ref|XP_017139445.1| PREDICTED: uncharacterized protein LOC108153857 [Drosophila
>gi|1037064476|ref|XP_017142149.1| PREDICTED: reticulon-3 [Drosophila miranda]
>gi|1037097936|ref|XP_017144562.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037097947|ref|XP_017144565.1| PREDICTED: ubiquitin-conjugating enzyme E2-18 kDa [Drosophil
>gi|1037088826|ref|XP_017143844.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037088840|ref|XP_017143845.1| PREDICTED: esterase B1 isoform X1 [Drosophila miranda]
>gi|1037088880|ref|XP_017143848.1| PREDICTED: growth hormone-inducible transmembrane protein [
>gi|1037088854|ref|XP_017143846.1| PREDICTED: esterase B1 isoform X2 [Drosophila miranda]
>gi|1037088867|ref|XP_017143847.1| PREDICTED: esterase B1 [Drosophila miranda]
>gi|1037036647|ref|XP_017140281.1| PREDICTED: uncharacterized protein LOC108154492 [Drosophila
>gi|1037035357|ref|XP_017140209.1| PREDICTED: neural cell adhesion molecule 1-B isoform X1 [Dr
>gi|1037035375|ref|XP_017140210.1| PREDICTED: neural cell adhesion molecule 1-B isoform X2 [Dr
>gi|1037106477|ref|XP_017145503.1| PREDICTED: uncharacterized protein LOC108157815 [Drosophila
>gi|1037022823|ref|XP_017139446.1| PREDICTED: low-density lipoprotein receptor-related protein
>gi|1037022840|ref|XP_017139447.1| PREDICTED: glutamate receptor U1 [Drosophila miranda]
>gi|1037035824|ref|XP_017140231.1| PREDICTED: uncharacterized protein LOC108154471 isoform X1
>gi|1037035840|ref|XP_017140232.1| PREDICTED: uncharacterized protein LOC108154471 isoform X2
>gi|1037106378|ref|XP_017145488.1| PREDICTED: uncharacterized protein LOC108157805 [Drosophila
>gi|1037105594|ref|XP_017145380.1| PREDICTED: uncharacterized protein LOC108157722 [Drosophila
>gi|1037105586|ref|XP_017145378.1| PREDICTED: classical arabinogalactan protein 9 [Drosophila
>gi|1037022884|ref|XP_017139448.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
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>gi|1037106815|ref|XP_017145550.1| PREDICTED: endoplasmic reticulum resident protein 44 [Droso
>gi|1037095102|ref|XP_017144312.1| PREDICTED: LOW QUALITY PROTEIN: zinc finger protein rotund
>gi|1037095186|ref|XP_017144319.1| PREDICTED: GTPase-activating protein RacGAP84C-like [Drosopl
>gi|1037022481|ref|XP_017139423.1| PREDICTED: LOW QUALITY PROTEIN: GTPase-activating protein Ra
>gi|1037095172|ref|XP_017144318.1| PREDICTED: transcription activator MSS11-like [Drosophila m:
>gi|1037095158|ref|XP_017144317.1| PREDICTED: very low-density lipoprotein receptor [Drosophila
>gi|1037095116|ref|XP_017144313.1| PREDICTED: synaptotagmin-4 [Drosophila miranda]
>gi|1037095130|ref|XP_017144314.1| PREDICTED: synaptotagmin-4 [Drosophila miranda]
>gi|1037095144|ref|XP_017144315.1| PREDICTED: synaptotagmin-4 [Drosophila miranda]
>gi|1037066760|ref|XP_017142289.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1037066776|ref|XP_017142290.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1037066901|ref|XP_017142299.1| PREDICTED: general odorant-binding protein 84a isoform X2 [I
>gi|1037066886|ref|XP_017142298.1| PREDICTED: general odorant-binding protein 84a isoform X1 [I
>gi|1037066917|ref|XP_017142300.1| PREDICTED: general odorant-binding protein 84a isoform X3 [I
>gi|1037066870|ref|XP_017142297.1| PREDICTED: DTW domain-containing protein 2 [Drosophila mirar
>gi|1037066822|ref|XP_017142294.1| PREDICTED: serine/threonine-protein kinase 16 [Drosophila m:
>gi|1037066854|ref|XP_017142296.1| PREDICTED: G patch domain-containing protein 11 isoform X2
>gi|1037066838|ref|XP_017142295.1| PREDICTED: G patch domain-containing protein 11 isoform X1
>gi|1037066744|ref|XP_017142288.1| PREDICTED: nucleolar complex protein 3 homolog [Drosophila r
>gi|1037066808|ref|XP_017142292.1| PREDICTED: uncharacterized protein LOC108155760 isoform X2
>gi|1037066792|ref|XP_017142291.1| PREDICTED: uncharacterized protein LOC108155760 isoform X1
>gi|1037066700|ref|XP_017142285.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037066714|ref|XP_017142286.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037066729|ref|XP_017142287.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037105553|ref|XP_017145374.1| PREDICTED: facilitated trehalose transporter Tret1-like [Dr
>gi|1037106033|ref|XP_017145441.1| PREDICTED: uncharacterized protein LOC108157769 isoform X1
>gi|1037106041|ref|XP_017145442.1| PREDICTED: uncharacterized protein LOC108157769 isoform X2
>gi|1037106049|ref|XP_017145443.1| PREDICTED: uncharacterized protein LOC108157769 isoform X3
>gi|1037105999|ref|XP_017145436.1| PREDICTED: uncharacterized protein LOC108157765 [Drosophila
>gi|1037084425|ref|XP_017143541.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084413|ref|XP_017143540.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084467|ref|XP_017143545.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084374|ref|XP_017143537.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084387|ref|XP_017143538.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084360|ref|XP_017143536.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084480|ref|XP_017143546.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084439|ref|XP_017143543.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084401|ref|XP_017143539.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084494|ref|XP_017143547.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037084453|ref|XP_017143544.1| PREDICTED: phosphatidylinositol-binding clathrin assembly p1
>gi|1037096367|ref|XP_017144418.1| PREDICTED: protein SHQ1 homolog [Drosophila miranda]
>gi|1037072709|ref|XP_017142700.1| PREDICTED: putative epidermal cell surface receptor isoform
>gi|1037072694|ref|XP_017142699.1| PREDICTED: putative epidermal cell surface receptor isoform
>gi|1037072664|ref|XP_017142697.1| PREDICTED: putative epidermal cell surface receptor isoform
>gi|1037072680|ref|XP_017142698.1| PREDICTED: putative epidermal cell surface receptor isoform
>gi|1037066134|ref|XP_017142250.1| PREDICTED: uncharacterized protein LOC108155743 [Drosophila
>gi|1037066150|ref|XP_017142251.1| PREDICTED: uncharacterized protein LOC108155743 [Drosophila
>gi|1037066165|ref|XP_017142252.1| PREDICTED: uncharacterized protein LOC108155743 [Drosophila
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>gi|1037066106|ref|XP_017142248.1| PREDICTED: centromere protein J [Drosophila miranda]
>gi|1037066120|ref|XP_017142249.1| PREDICTED: GPN-loop GTPase 3 [Drosophila miranda]
>gi|1037066075|ref|XP_017142246.1| PREDICTED: uncharacterized protein LOC108155740 isoform X1
>gi|1037066089|ref|XP_017142247.1| PREDICTED: glutathione S-transferase 1-1 isoform X2 [Drosop
>gi|1037104477|ref|XP_017145219.1| PREDICTED: uncharacterized protein LOC108157601 isoform X2
>gi|1037104469|ref|XP_017145218.1| PREDICTED: protein twist isoform X1 [Drosophila miranda]
>gi|1037099187|ref|XP_017144676.1| PREDICTED: uncharacterized protein LOC108157230 [Drosophila
>gi|1037099166|ref|XP_017144675.1| PREDICTED: uncharacterized protein LOC108157229 [Drosophila
>gi|1037050467|ref|XP_017141189.1| PREDICTED: ELMO domain-containing protein 2 [Drosophila mira
>gi|1037050452|ref|XP_017141188.1| PREDICTED: secernin-3 [Drosophila miranda]
>gi|1037050273|ref|XP_017141176.1| PREDICTED: protein AF-17 isoform X1 [Drosophila miranda]
>gi|1037050289|ref|XP_017141177.1| PREDICTED: protein AF-17 isoform X2 [Drosophila miranda]
>gi|1037050309|ref|XP_017141178.1| PREDICTED: protein AF-17 isoform X3 [Drosophila miranda]
>gi|1037050323|ref|XP_017141179.1| PREDICTED: protein AF-10 isoform X4 [Drosophila miranda]
>gi|1037050339|ref|XP_017141180.1| PREDICTED: protein AF-10 isoform X5 [Drosophila miranda]
>gi|1037050356|ref|XP_017141182.1| PREDICTED: protein AF-17 isoform X6 [Drosophila miranda]
>gi|1037050370|ref|XP_017141183.1| PREDICTED: protein AF-10 isoform X7 [Drosophila miranda]
>gi|1037050420|ref|XP_017141186.1| PREDICTED: muscle LIM protein Mlp84B [Drosophila miranda]
>gi|1037050436|ref|XP_017141187.1| PREDICTED: muscle LIM protein Mlp84B [Drosophila miranda]
>gi|1037050388|ref|XP_017141184.1| PREDICTED: mucin-5AC isoform X8 [Drosophila miranda]
>gi|1037050404|ref|XP_017141185.1| PREDICTED: mucin-19 isoform X9 [Drosophila miranda]
>gi|1037050481|ref|XP_017141191.1| PREDICTED: enhancer of yellow 2b transcription factor [Dros
>gi|1037052639|ref|XP_017141339.1| PREDICTED: tubulin alpha-1 chain [Drosophila miranda]
>gi|1037052705|ref|XP_017141344.1| PREDICTED: rRNA methyltransferase 2, mitochondrial [Drosoph
>gi|1037052625|ref|XP_017141338.1| PREDICTED: venom serine carboxypeptidase [Drosophila mirand
>gi|1037052721|ref|XP_017141345.1| PREDICTED: transmembrane protein 256 homolog [Drosophila mi
>gi|1037052607|ref|XP_017141337.1| PREDICTED: LOW QUALITY PROTEIN: probable multidrug resistanc
>gi|1037052672|ref|XP_017141342.1| PREDICTED: zinc finger homeobox protein 3 [Drosophila miranc
>gi|1037052656|ref|XP_017141341.1| PREDICTED: short-chain specific acyl-CoA dehydrogenase, mitc
>gi|1037052687|ref|XP_017141343.1| PREDICTED: zinc finger protein 420-like [Drosophila miranda]
>gi|1037070024|ref|XP_017142518.1| PREDICTED: mitogen-activated protein kinase kinase kinase 15
>gi|1037070068|ref|XP_017142522.1| PREDICTED: opsin Rh1 [Drosophila miranda]
>gi|1037070038|ref|XP_017142520.1| PREDICTED: serine/threonine protein phosphatase 2A regulato
>gi|1037070054|ref|XP_017142521.1| PREDICTED: serine/threonine protein phosphatase 2A regulato
>gi|1037042973|ref|XP_017140688.1| PREDICTED: ATP-dependent Clp protease ATP-binding subunit cl
>gi|1037043007|ref|XP_017140690.1| PREDICTED: surfeit locus protein 6 homolog [Drosophila mirar
>gi|1037043023|ref|XP_017140691.1| PREDICTED: DNA-directed RNA polymerase II subunit RPB7 [Dros
>gi|1037042955|ref|XP_017140686.1| PREDICTED: small G protein signaling modulator 3 homolog [Dr
>gi|1037042874|ref|XP_017140681.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042892|ref|XP_017140682.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042908|ref|XP_017140683.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042922|ref|XP_017140684.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037042938|ref|XP_017140685.1| PREDICTED: trafficking protein particle complex subunit 10 i
>gi|1037043039|ref|XP_017140692.1| PREDICTED: diamine acetyltransferase 2 [Drosophila miranda]
>gi|1037042827|ref|XP_017140678.1| PREDICTED: box A-binding factor isoform X2 [Drosophila mirar
>gi|1037042813|ref|XP_017140677.1| PREDICTED: box A-binding factor isoform X1 [Drosophila mirar
>gi|1037042843|ref|XP_017140679.1| PREDICTED: box A-binding factor isoform X3 [Drosophila mirar
>gi|1037042859|ref|XP_017140680.1| PREDICTED: box A-binding factor isoform X4 [Drosophila mirar
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>gi|1037042991|ref|XP_017140689.1| PREDICTED: box A-binding factor, partial [Drosophila miranda
>gi|1037076802|ref|XP_017142981.1| PREDICTED: GATA-binding factor A isoform X1 [Drosophila mira
>gi|1037076818|ref|XP_017142982.1| PREDICTED: GATA-binding factor A isoform X2 [Drosophila mira
>gi|1037076834|ref|XP_017142983.1| PREDICTED: GATA-binding factor A isoform X3 [Drosophila mira
>gi|1037076868|ref|XP_017142986.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037076883|ref|XP_017142987.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037076850|ref|XP_017142984.1| PREDICTED: inhibitor of nuclear factor kappa-B kinase subuni
>gi|1037076899|ref|XP_017142988.1| PREDICTED: methyltransferase-like protein 23 [Drosophila mi
>gi|1037060005|ref|XP_017141843.1| PREDICTED: LOW QUALITY PROTEIN: cytoskeleton-associated prot
>gi|1037059987|ref|XP_017141842.1| PREDICTED: LOW QUALITY PROTEIN: NACHT and WD repeat domain-c
>gi|1037062249|ref|XP_017141995.1| PREDICTED: unconventional myosin-XVIIIa isoform X1 [Drosophi
>gi|1037062279|ref|XP_017141997.1| PREDICTED: unconventional myosin-XVIIIa isoform X3 [Drosophi
>gi|1037062263|ref|XP_017141996.1| PREDICTED: unconventional myosin-XVIIIa isoform X2 [Drosophi
>gi|1037062295|ref|XP_017141998.1| PREDICTED: unconventional myosin-XVIIIa isoform X4 [Drosophi
>gi|1037062311|ref|XP_017141999.1| PREDICTED: unconventional myosin-XVIIIa isoform X5 [Drosophi
>gi|1037062327|ref|XP_017142000.1| PREDICTED: uncharacterized protein LOC108155593 [Drosophila
>gi|1037062341|ref|XP_017142001.1| PREDICTED: lipase member H [Drosophila miranda]
>gi|1037091445|ref|XP_017144050.1| PREDICTED: RAC serine/threonine-protein kinase [Drosophila m
>gi|1037097366|ref|XP_017144508.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1037097378|ref|XP_017144509.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1037097390|ref|XP_017144510.1| PREDICTED: serine proteinase stubble isoform X1 [Drosophila
>gi|1037097402|ref|XP_017144511.1| PREDICTED: serine proteinase stubble isoform X2 [Drosophila
>gi|1037097413|ref|XP_017144513.1| PREDICTED: uncharacterized protein LOC108157128 [Drosophila
>gi|1037094063|ref|XP_017144232.1| PREDICTED: growth hormone-regulated TBC protein 1-A [Drosop
>gi|1037094077|ref|XP_017144233.1| PREDICTED: growth hormone-regulated TBC protein 1-A [Drosop
>gi|1037094121|ref|XP_017144236.1| PREDICTED: uncharacterized protein LOC108156965 isoform X3
>gi|1037094107|ref|XP_017144235.1| PREDICTED: circumsporozoite protein isoform X2 [Drosophila m
>gi|1037094091|ref|XP_017144234.1| PREDICTED: circumsporozoite protein isoform X1 [Drosophila m
>gi|1037097460|ref|XP_017144517.1| PREDICTED: organic cation transporter protein isoform X1 [Dr
>gi|1037097471|ref|XP_017144518.1| PREDICTED: organic cation transporter protein isoform X2 [Dr
>gi|1037097482|ref|XP_017144519.1| PREDICTED: organic cation transporter protein isoform X2 [Dr
>gi|1037097449|ref|XP_017144516.1| PREDICTED: organic cation transporter protein isoform X3 [Dr
>gi|1037097425|ref|XP_017144514.1| PREDICTED: organic cation transporter protein isoform X1 [Dr
>gi|1037097437|ref|XP_017144515.1| PREDICTED: organic cation transporter protein isoform X2 [Dr
>gi|1037100870|ref|XP_017144827.1| PREDICTED: caprin homolog [Drosophila miranda]
>gi|1037100880|ref|XP_017144828.1| PREDICTED: histone deacetylase complex subunit SAP18 [Drosop
>gi|1037091678|ref|XP_017144066.1| PREDICTED: protein sarah [Drosophila miranda]
>gi|1037091664|ref|XP_017144065.1| PREDICTED: organic cation transporter protein [Drosophila mi
>gi|1037076111|ref|XP_017142936.1| PREDICTED: uncharacterized protein LOC108156135 [Drosophila
>gi|1037076127|ref|XP_017142937.1| PREDICTED: single-stranded DNA-binding protein, mitochondria
>gi|1037076171|ref|XP_017142941.1| PREDICTED: GPI ethanolamine phosphate transferase 1 [Drosoph
>gi|1037076142|ref|XP_017142939.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037076156|ref|XP_017142940.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037022900|ref|XP_017139449.1| PREDICTED: GPI ethanolamine phosphate transferase 1 [Drosoph
>gi|1037080982|ref|XP_017143279.1| PREDICTED: TBC1 domain family member 10A isoform X1 [Drosoph
>gi|1037080995|ref|XP_017143280.1| PREDICTED: TBC1 domain family member 10A isoform X2 [Drosoph
>gi|1037080968|ref|XP_017143277.1| PREDICTED: zinc finger protein 850 [Drosophila miranda]
>gi|1037078498|ref|XP_017143100.1| PREDICTED: microtubule-associated protein Jupiter isoform X0
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>gi|1037078466|ref|XP_017143098.1| PREDICTED: microtubule-associated protein Jupiter isoform X4
>gi|1037078333|ref|XP_017143096.1| PREDICTED: microtubule-associated protein Jupiter isoform X2
>gi|1037078484|ref|XP_017143099.1| PREDICTED: microtubule-associated protein Jupiter isoform X5
>gi|1037078452|ref|XP_017143097.1| PREDICTED: microtubule-associated protein Jupiter isoform X3
>gi|1037078319|ref|XP_017143095.1| PREDICTED: microtubule-associated protein Jupiter isoform X1
>gi|1037095425|ref|XP_017144336.1| PREDICTED: inhibitor of growth protein 2 [Drosophila miranda
>gi|1037095441|ref|XP_017144337.1| PREDICTED: uncharacterized protein LOC108157036 [Drosophila
>gi|1037093434|ref|XP_017144188.1| PREDICTED: uncharacterized protein LOC108156938 isoform X1
>gi|1037093450|ref|XP_017144190.1| PREDICTED: rho guanine nucleotide exchange factor 17 isoform
>gi|1037098889|ref|XP_017144656.1| PREDICTED: alpha-1D adrenergic receptor isoform X2 [Drosophi
>gi|1037098901|ref|XP_017144657.1| PREDICTED: alpha-1D adrenergic receptor isoform X3 [Drosophi
>gi|1037098858|ref|XP_017144654.1| PREDICTED: tyramine receptor 1 isoform X1 [Drosophila mirand
>gi|1037098871|ref|XP_017144655.1| PREDICTED: tyramine receptor 1 isoform X1 [Drosophila mirand
>gi|1037098913|ref|XP_017144659.1| PREDICTED: 5-hydroxytryptamine receptor 1A isoform X1 [Droso
>gi|1037098925|ref|XP_017144660.1| PREDICTED: 5-hydroxytryptamine receptor 1A isoform X1 [Droso
>gi|1037098937|ref|XP_017144661.1| PREDICTED: uncharacterized protein LOC108157216 isoform X2
>gi|1037098949|ref|XP_017144662.1| PREDICTED: uncharacterized protein LOC108157216 isoform X2
>gi|1037102468|ref|XP_017144980.1| PREDICTED: lutropin-choriogonadotropic hormone receptor [Dro
>gi|1037102478|ref|XP_017144981.1| PREDICTED: gamma-aminobutyric acid receptor-associated prote
>gi|1037063268|ref|XP_017142066.1| PREDICTED: trithorax group protein osa isoform X2 [Drosophil
>gi|1037063298|ref|XP_017142068.1| PREDICTED: trithorax group protein osa isoform X4 [Drosophil
>gi|1037063254|ref|XP_017142065.1| PREDICTED: trithorax group protein osa isoform X1 [Drosophil
>gi|1037063284|ref|XP_017142067.1| PREDICTED: trithorax group protein osa isoform X3 [Drosophil
>gi|1037083433|ref|XP_017143464.1| PREDICTED: protein YSC84 [Drosophila miranda]
>gi|1037083447|ref|XP_017143465.1| PREDICTED: protein YSC84 [Drosophila miranda]
>gi|1037083461|ref|XP_017143466.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083475|ref|XP_017143467.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083489|ref|XP_017143468.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083503|ref|XP_017143469.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037083517|ref|XP_017143470.1| PREDICTED: stabilizer of axonemal microtubules 1 [Drosophila
>gi|1037053454|ref|XP_017141394.1| PREDICTED: zinc finger protein 420 [Drosophila miranda]
>gi|1037053671|ref|XP_017141409.1| PREDICTED: transcription initiation factor TFIID subunit 12
>gi|1037053681|ref|XP_017141410.1| PREDICTED: transcription initiation factor TFIID subunit 12
>gi|1037053597|ref|XP_017141404.1| PREDICTED: heme oxygenase 1 [Drosophila miranda]
>gi|1037053470|ref|XP_017141395.1| PREDICTED: uncharacterized protein LOC108155222 [Drosophila
>gi|1037053549|ref|XP_017141401.1| PREDICTED: uncharacterized protein LOC108155225 [Drosophila
>gi|1037053581|ref|XP_017141403.1| PREDICTED: uncharacterized protein LOC108155227 [Drosophila
>gi|1037053629|ref|XP_017141406.1| PREDICTED: protein Fer3 [Drosophila miranda]
>gi|1037053643|ref|XP_017141407.1| PREDICTED: protein Fer3 [Drosophila miranda]
>gi|1037053659|ref|XP_017141408.1| PREDICTED: protein Fer3 [Drosophila miranda]
>gi|1037053565|ref|XP_017141402.1| PREDICTED: uncharacterized protein LOC108155226 [Drosophila
>gi|1037053613|ref|XP_017141405.1| PREDICTED: uncharacterized protein LOC108155229 [Drosophila
>gi|1037053485|ref|XP_017141396.1| PREDICTED: uncharacterized protein LOC108155224 [Drosophila
>gi|1037053501|ref|XP_017141397.1| PREDICTED: uncharacterized protein LOC108155224 [Drosophila
>gi|1037053517|ref|XP_017141398.1| PREDICTED: uncharacterized protein LOC108155224 [Drosophila
>gi|1037053533|ref|XP_017141399.1| PREDICTED: uncharacterized protein LOC108155224 [Drosophila
>gi|1037059111|ref|XP_017141778.1| PREDICTED: thiamin pyrophosphokinase 1 [Drosophila miranda]
>gi|1037059081|ref|XP_017141776.1| PREDICTED: Bloom syndrome protein homolog [Drosophila mirand
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>gi|1037059097|ref|XP_017141777.1| PREDICTED: WD repeat-containing protein 55 homolog [Drosophi
>gi|1037059067|ref|XP_017141774.1| PREDICTED: putative uncharacterized protein DDB_G0291608 [Dr
>gi|1037068272|ref|XP_017142395.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1037068284|ref|XP_017142396.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1037061788|ref|XP_017141963.1| PREDICTED: zinc finger protein 92 homolog [Drosophila miran
>gi|1037061736|ref|XP_017141959.1| PREDICTED: uncharacterized protein LOC108155570 [Drosophila
>gi|1037061754|ref|XP_017141960.1| PREDICTED: glycine receptor subunit alpha-4 isoform X1 [Dros
>gi|1037061770|ref|XP_017141962.1| PREDICTED: glycine receptor subunit alpha-4 isoform X2 [Dros
>gi|1037074159|ref|XP_017142802.1| PREDICTED: myotubularin-related protein 13 isoform X2 [Droso
>gi|1037074143|ref|XP_017142801.1| PREDICTED: myotubularin-related protein 13 isoform X1 [Droso
>gi|1037097317|ref|XP_017144504.1| PREDICTED: chloride channel protein 2 isoform X8 [Drosophila
>gi|1037097280|ref|XP_017144500.1| PREDICTED: chloride channel protein 2 isoform X5 [Drosophila
>gi|1037097292|ref|XP_017144501.1| PREDICTED: chloride channel protein 2 isoform X6 [Drosophila
>gi|1037097256|ref|XP_017144498.1| PREDICTED: chloride channel protein 2 isoform X3 [Drosophila
>gi|1037097245|ref|XP_017144497.1| PREDICTED: chloride channel protein 2 isoform X2 [Drosophila
>gi|1037097231|ref|XP_017144496.1| PREDICTED: chloride channel protein 2 isoform X1 [Drosophila
>gi|1037097306|ref|XP_017144503.1| PREDICTED: chloride channel protein 2 isoform X7 [Drosophila
>gi|1037097268|ref|XP_017144499.1| PREDICTED: chloride channel protein 2 isoform X4 [Drosophila
>gi|1037097329|ref|XP_017144505.1| PREDICTED: chloride channel protein 2 isoform X9 [Drosophila
>gi|1037089082|ref|XP_017143864.1| PREDICTED: heterogeneous nuclear ribonucleoprotein H2 isofor
>gi|1037089096|ref|XP_017143865.1| PREDICTED: heterogeneous nuclear ribonucleoprotein H2 isofor
>gi|1037085476|ref|XP_017143618.1| PREDICTED: uncharacterized protein LOC108156590 [Drosophila
>gi|1037085462|ref|XP_017143617.1| PREDICTED: LOW QUALITY PROTEIN: nuclear pore complex protein
>gi|1037096594|ref|XP_017144441.1| PREDICTED: protein vreteno-like [Drosophila miranda]
>gi|1037103575|ref|XP_017145099.1| PREDICTED: dynein heavy chain 5, axonemal isoform X1 [Drosop
>gi|1037103583|ref|XP_017145100.1| PREDICTED: dynein heavy chain 5, axonemal isoform X2 [Drosop
>gi|1037103591|ref|XP_017145101.1| PREDICTED: uncharacterized protein LOC108157511 [Drosophila
>gi|1037060274|ref|XP_017141861.1| PREDICTED: uncharacterized protein LOC108155523 isoform X2
>gi|1037060241|ref|XP_017141859.1| PREDICTED: uncharacterized protein LOC108155523 isoform X1
>gi|1037060259|ref|XP_017141860.1| PREDICTED: uncharacterized protein LOC108155523 isoform X1
>gi|1037060289|ref|XP_017141862.1| PREDICTED: proteoglycan 4 [Drosophila miranda]
>gi|1037060179|ref|XP_017141854.1| PREDICTED: polycomb protein Scm [Drosophila miranda]
>gi|1037060305|ref|XP_017141863.1| PREDICTED: probable fatty acid-binding protein [Drosophila
>gi|1037060225|ref|XP_017141858.1| PREDICTED: putative tricarboxylate transport protein, mitoc
>gi|1037060194|ref|XP_017141855.1| PREDICTED: 8-oxo-dGDP phosphatase NUDT18 [Drosophila miranda
>gi|1037060209|ref|XP_017141856.1| PREDICTED: limbic system-associated membrane protein [Drosop
>gi|1037033057|ref|XP_017140075.1| PREDICTED: hemicentin-2 [Drosophila miranda]
>gi|1037059627|ref|XP_017141813.1| PREDICTED: cilia- and flagella-associated protein 53 [Drosop
>gi|1037059641|ref|XP_017141815.1| PREDICTED: cilia- and flagella-associated protein 53 [Drosop
>gi|1037059693|ref|XP_017141819.1| PREDICTED: peptidoglycan-recognition protein LB isoform X1
>gi|1037059719|ref|XP_017141821.1| PREDICTED: peptidoglycan-recognition protein LB isoform X3
>gi|1037059705|ref|XP_017141820.1| PREDICTED: peptidoglycan-recognition protein LB isoform X2
>gi|1037059737|ref|XP_017141822.1| PREDICTED: peptidoglycan-recognition protein LB isoform X4
>gi|1037059601|ref|XP_017141811.1| PREDICTED: probable serine/threonine-protein kinase kinX iso
>gi|1037059562|ref|XP_017141808.1| PREDICTED: glutamic acid-rich protein isoform X1 [Drosophila
>gi|1037059576|ref|XP_017141809.1| PREDICTED: glutamic acid-rich protein isoform X1 [Drosophila
>gi|1037059587|ref|XP_017141810.1| PREDICTED: glutamic acid-rich protein isoform X1 [Drosophila
>gi|1037059667|ref|XP_017141817.1| PREDICTED: solute carrier family 35 member G1 [Drosophila mi
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>gi|1037059614|ref|XP_017141812.1| PREDICTED: exonuclease 3'-5' domain-containing protein 2 [D
>gi|1037059654|ref|XP_017141816.1| PREDICTED: soluble calcium-activated nucleotidase 1 [Drosopl
>gi|1037059679|ref|XP_017141818.1| PREDICTED: solute carrier family 35 member B1 homolog [Dros
>gi|1037056045|ref|XP_017141566.1| PREDICTED: uncharacterized protein LOC108155333 isoform X2
>gi|1037056171|ref|XP_017141574.1| PREDICTED: nephrocan [Drosophila miranda]
>gi|1037056187|ref|XP_017141575.1| PREDICTED: leucine-rich repeat-containing protein 15 [Droso
>gi|1037056155|ref|XP_017141573.1| PREDICTED: sensory neuron membrane protein 1 [Drosophila mi
>gi|1037056030|ref|XP_017141565.1| PREDICTED: uncharacterized protein LOC108155333 isoform X1
>gi|1037056125|ref|XP_017141571.1| PREDICTED: uncharacterized protein LOC108155333 isoform X5
>gi|1037056062|ref|XP_017141567.1| PREDICTED: uncharacterized protein LOC108155333 isoform X3
>gi|1037056077|ref|XP_017141568.1| PREDICTED: uncharacterized protein LOC108155333 isoform X3
>gi|1037056090|ref|XP_017141569.1| PREDICTED: uncharacterized protein LOC108155333 isoform X3
>gi|1037056107|ref|XP_017141570.1| PREDICTED: uncharacterized protein LOC108155333 isoform X4
>gi|1037056141|ref|XP_017141572.1| PREDICTED: 5'-AMP-activated protein kinase subunit gamma-1
>gi|1037051148|ref|XP_017141239.1| PREDICTED: SET and MYND domain-containing protein 5 [Drosopl
>gi|1037051180|ref|XP_017141241.1| PREDICTED: transmembrane emp24 domain-containing protein eca
>gi|1037051197|ref|XP_017141242.1| PREDICTED: DDRGK domain-containing protein 1 [Drosophila mi
>gi|1037051211|ref|XP_017141243.1| PREDICTED: transmembrane emp24 domain-containing protein eca
>gi|1037051004|ref|XP_017141229.1| PREDICTED: solute carrier family 15 member 1 [Drosophila mi
>gi|1037051162|ref|XP_017141240.1| PREDICTED: nucleoside diphosphate kinase 7 [Drosophila mira
>gi|1037051275|ref|XP_017141247.1| PREDICTED: uncharacterized protein CG16817 isoform X4 [Dros
>gi|1037051260|ref|XP_017141246.1| PREDICTED: uncharacterized protein CG16817 isoform X3 [Dros
>gi|1037051246|ref|XP_017141245.1| PREDICTED: uncharacterized protein CG16817 isoform X2 [Dros
>gi|1037051231|ref|XP_017141244.1| PREDICTED: uncharacterized protein CG16817 isoform X1 [Dros
>gi|1037051287|ref|XP_017141249.1| PREDICTED: cytochrome c oxidase subunit 5A, mitochondrial [
>gi|1037051036|ref|XP_017141231.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform
>gi|1037051020|ref|XP_017141230.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform
>gi|1037051052|ref|XP_017141232.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform
>gi|1037051068|ref|XP_017141233.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform
>gi|1037051082|ref|XP_017141234.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform
>gi|1037051098|ref|XP_017141235.1| PREDICTED: kynurenine--oxoglutarate transaminase 3 isoform
>gi|1037051114|ref|XP_017141237.1| PREDICTED: trophoblast glycoprotein [Drosophila miranda]
>gi|1037051132|ref|XP_017141238.1| PREDICTED: trophoblast glycoprotein [Drosophila miranda]
>gi|1037091720|ref|XP_017144070.1| PREDICTED: cytochrome P450 315a1, mitochondrial isoform X1
>gi|1037091734|ref|XP_017144071.1| PREDICTED: cytochrome P450 315a1, mitochondrial isoform X2
>gi|1037091706|ref|XP_017144069.1| PREDICTED: sphingomyelin phosphodiesterase 4 [Drosophila mi
>gi|1037068299|ref|XP_017142397.1| PREDICTED: intron-binding protein aquarius isoform X1 [Dros
>gi|1037068346|ref|XP_017142400.1| PREDICTED: probable G-protein coupled receptor Mth-like 5 is
>gi|1037068329|ref|XP_017142399.1| PREDICTED: probable G-protein coupled receptor Mth-like 5 is
>gi|1037068314|ref|XP_017142398.1| PREDICTED: intron-binding protein aquarius isoform X2 [Dros
>gi|1037068361|ref|XP_017142401.1| PREDICTED: putative inner dynein arm light chain, axonemal
>gi|1037104540|ref|XP_017145228.1| PREDICTED: histone-like protein 18C [Drosophila miranda]
>gi|1037062089|ref|XP_017141984.1| PREDICTED: cadherin-87A [Drosophila miranda]
>gi|1037062108|ref|XP_017141985.1| PREDICTED: uncharacterized protein LOC108155587 isoform X1
>gi|1037062124|ref|XP_017141986.1| PREDICTED: uncharacterized protein LOC108155587 isoform X2
>gi|1037077061|ref|XP_017142999.1| PREDICTED: hormone receptor 4 isoform X1 [Drosophila miranda
>gi|1037077075|ref|XP_017143000.1| PREDICTED: hormone receptor 4 isoform X2 [Drosophila miranda
>gi|1037082883|ref|XP_017143421.1| PREDICTED: double-strand-break repair protein rad21-like pro
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>gi|1037103848|ref|XP_017145135.1| PREDICTED: uncharacterized protein LOC108157543 [Drosophila
>gi|1037095355|ref|XP_017144331.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037093420|ref|XP_017144187.1| PREDICTED: piezo-type mechanosensitive ion channel component
>gi|1037100163|ref|XP_017144758.1| PREDICTED: homeobox protein Nkx-2.1 isoform X3 [Drosophila m
>gi|1037100151|ref|XP_017144756.1| PREDICTED: homeobox protein Nkx-2.1 isoform X2 [Drosophila m
>gi|1037100127|ref|XP_017144754.1| PREDICTED: homeobox protein Nkx-2.1 isoform X1 [Drosophila m
>gi|1037100139|ref|XP_017144755.1| PREDICTED: homeobox protein Nkx-2.1 isoform X1 [Drosophila m
>gi|1037083866|ref|XP_017143498.1| PREDICTED: poly [ADP-ribose] polymerase [Drosophila miranda]
>gi|1037099067|ref|XP_017144671.1| PREDICTED: programmed cell death protein 6 [Drosophila miran
>gi|1037096627|ref|XP_017144444.1| PREDICTED: ubiquitin-conjugating enzyme E2 G1 [Drosophila mi
>gi|1037102488|ref|XP_017144982.1| PREDICTED: uncharacterized protein LOC108157439 [Drosophila
>gi|1037092806|ref|XP_017144143.1| PREDICTED: calcium/calmodulin-dependent protein kinase kinas
>gi|1037092820|ref|XP_017144144.1| PREDICTED: calcium/calmodulin-dependent protein kinase kinas
>gi|1037092834|ref|XP_017144145.1| PREDICTED: uncharacterized protein LOC108156906 [Drosophila
>gi|1037037576|ref|XP_017140336.1| PREDICTED: uncharacterized protein LOC108154544 [Drosophila
>gi|1037047298|ref|XP_017140975.1| PREDICTED: rabankyrin-5 [Drosophila miranda]
>gi|1037047360|ref|XP_017140980.1| PREDICTED: uncharacterized protein LOC108154973 isoform X1
>gi|1037047376|ref|XP_017140981.1| PREDICTED: uncharacterized protein LOC108154973 isoform X2
>gi|1037047420|ref|XP_017140984.1| PREDICTED: cytochrome b-c1 complex subunit 6, mitochondrial
>gi|1037047284|ref|XP_017140974.1| PREDICTED: histone-lysine N-methyltransferase SETD1 [Drosoph
>gi|1037047408|ref|XP_017140983.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1037047392|ref|XP_017140982.1| PREDICTED: mediator of RNA polymerase II transcription subun
>gi|1037047312|ref|XP_017140976.1| PREDICTED: uncharacterized protein LOC108154972 isoform X1
>gi|1037047328|ref|XP_017140977.1| PREDICTED: uncharacterized protein LOC108154972 isoform X2
>gi|1037047344|ref|XP_017140978.1| PREDICTED: uncharacterized protein LOC108154972 isoform X2
>gi|1037047251|ref|XP_017140972.1| PREDICTED: fatty acid synthase isoform X1 [Drosophila miran
>gi|1037047269|ref|XP_017140973.1| PREDICTED: fatty acid synthase isoform X2 [Drosophila miran
>gi|1037106221|ref|XP_017145467.1| PREDICTED: uncharacterized protein LOC108157788 [Drosophila
>gi|1037106229|ref|XP_017145468.1| PREDICTED: uncharacterized protein LOC108157788 [Drosophila
>gi|1037106235|ref|XP_017145469.1| PREDICTED: uncharacterized protein LOC108157788 [Drosophila
>gi|1037106578|ref|XP_017145515.1| PREDICTED: tensin-2-like [Drosophila miranda]
>gi|1037106586|ref|XP_017145516.1| PREDICTED: tensin-2-like [Drosophila miranda]
>gi|1037102963|ref|XP_017145030.1| PREDICTED: chymotrypsin-like protease CTRL-1 [Drosophila mi
>gi|1037102973|ref|XP_017145031.1| PREDICTED: chymotrypsin-like protease CTRL-1 [Drosophila mi
>gi|1037095244|ref|XP_017144323.1| PREDICTED: uncharacterized protein LOC108157024 isoform X2
>gi|1037095216|ref|XP_017144321.1| PREDICTED: uncharacterized protein LOC108157024 isoform X1
>gi|1037095230|ref|XP_017144322.1| PREDICTED: uncharacterized protein LOC108157024 isoform X1
>gi|1037102509|ref|XP_017144984.1| PREDICTED: uncharacterized protein LOC108157441 isoform X1
>gi|1037102521|ref|XP_017144985.1| PREDICTED: uncharacterized protein LOC108157441 isoform X1
>gi|1037102529|ref|XP_017144986.1| PREDICTED: uncharacterized protein LOC108157441 isoform X2
>gi|1037038977|ref|XP_017140420.1| PREDICTED: uncharacterized protein LOC108154614 [Drosophila
>gi|1037096789|ref|XP_017144456.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096824|ref|XP_017144459.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096800|ref|XP_017144457.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096836|ref|XP_017144460.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096812|ref|XP_017144458.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096848|ref|XP_017144461.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
>gi|1037096862|ref|XP_017144462.1| PREDICTED: glutamine--fructose-6-phosphate aminotransferase
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>gi|1037038253|ref|XP_017140374.1| PREDICTED: mucin-5AC isoform X2 [Drosophila miranda]
>gi|1037038239|ref|XP_017140373.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037038269|ref|XP_017140375.1| PREDICTED: uncharacterized protein LOC108154582 isoform X3
>gi|1037106485|ref|XP_017145504.1| PREDICTED: alpha-N-acetylgalactosaminidase-like isoform X1
>gi|1037106493|ref|XP_017145505.1| PREDICTED: alpha-N-acetylgalactosaminidase-like isoform X2
>gi|1037078991|ref|XP_017143138.1| PREDICTED: uncharacterized protein LOC108156277 isoform X1
>gi|1037079002|ref|XP_017143139.1| PREDICTED: uncharacterized protein LOC108156277 isoform X2
>gi|1037079030|ref|XP_017143141.1| PREDICTED: drosulfakinins [Drosophila miranda]
>gi|1037079016|ref|XP_017143140.1| PREDICTED: beta-mannosidase-like [Drosophila miranda]
>gi|1037093464|ref|XP_017144191.1| PREDICTED: beta-mannosidase [Drosophila miranda]
>gi|1037093478|ref|XP_017144192.1| PREDICTED: uncharacterized protein LOC108156940 [Drosophila
>gi|1037093492|ref|XP_017144193.1| PREDICTED: uncharacterized protein LOC108156940 [Drosophila
>gi|1037106743|ref|XP_017145538.1| PREDICTED: uncharacterized protein LOC108157846 [Drosophila
>gi|1037068068|ref|XP_017142381.1| PREDICTED: cyclin-G-associated kinase [Drosophila miranda]
>gi|1037068083|ref|XP_017142382.1| PREDICTED: guanine deaminase [Drosophila miranda]
>gi|1037068099|ref|XP_017142383.1| PREDICTED: pre-mRNA-splicing factor RBM22 [Drosophila miranda
>gi|1037081173|ref|XP_017143294.1| PREDICTED: ATP-dependent RNA helicase abstrakt [Drosophila m
>gi|1037081160|ref|XP_017143293.1| PREDICTED: gelsolin isoform X2 [Drosophila miranda]
>gi|1037081148|ref|XP_017143292.1| PREDICTED: gelsolin isoform X1 [Drosophila miranda]
>gi|1037081187|ref|XP_017143295.1| PREDICTED: serine protease snake isoform X1 [Drosophila mira
>gi|1037081201|ref|XP_017143296.1| PREDICTED: uncharacterized protein LOC108156389 isoform X2
>gi|1037081215|ref|XP_017143297.1| PREDICTED: uncharacterized protein LOC108156390 [Drosophila
>gi|1037036907|ref|XP_017140295.1| PREDICTED: uncharacterized protein LOC108154506 [Drosophila
>gi|1037038787|ref|XP_017140408.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037035752|ref|XP_017140227.1| PREDICTED: adenylate cyclase, terminal-differentiation speci
>gi|1037038088|ref|XP_017140362.1| PREDICTED: uncharacterized protein LOC108154573 [Drosophila
>gi|1037035631|ref|XP_017140220.1| PREDICTED: odorant receptor 82a [Drosophila miranda]
>gi|1037033818|ref|XP_017140120.1| PREDICTED: omega-amidase NIT2-like [Drosophila miranda]
>gi|1037090527|ref|XP_017143981.1| PREDICTED: uncharacterized protein LOC108156805 [Drosophila
>gi|1037090543|ref|XP_017143982.1| PREDICTED: ankyrin repeat, PH and SEC7 domain containing pro
>gi|1037090557|ref|XP_017143983.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090571|ref|XP_017143985.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090585|ref|XP_017143986.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090599|ref|XP_017143987.1| PREDICTED: complexin isoform X1 [Drosophila miranda]
>gi|1037090613|ref|XP_017143988.1| PREDICTED: complexin isoform X2 [Drosophila miranda]
>gi|1037090627|ref|XP_017143989.1| PREDICTED: complexin isoform X3 [Drosophila miranda]
>gi|1037090641|ref|XP_017143990.1| PREDICTED: complexin isoform X4 [Drosophila miranda]
>gi|1037090514|ref|XP_017143980.1| PREDICTED: uncharacterized protein LOC108156804 [Drosophila
>gi|1037096934|ref|XP_017144468.1| PREDICTED: serine protease easter [Drosophila miranda]
>gi|1037096946|ref|XP_017144469.1| PREDICTED: charged multivesicular body protein 3 [Drosophila
>gi|1037092548|ref|XP_017144128.1| PREDICTED: ATP-dependent RNA helicase SUV3 homolog, mitochon
>gi|1037092562|ref|XP_017144129.1| PREDICTED: synaptotagmin-16 [Drosophila miranda]
>gi|1037071455|ref|XP_017142614.1| PREDICTED: uncharacterized protein LOC108155960 isoform X1
>gi|1037071471|ref|XP_017142615.1| PREDICTED: uncharacterized protein LOC108155960 isoform X1
>gi|1037071487|ref|XP_017142617.1| PREDICTED: uncharacterized protein LOC108155960 isoform X1
>gi|1037071501|ref|XP_017142618.1| PREDICTED: uncharacterized protein LOC108155960 isoform X2
>gi|1037071549|ref|XP_017142621.1| PREDICTED: AF4/FMR2 family member 4 isoform X5 [Drosophila m
>gi|1037071533|ref|XP_017142620.1| PREDICTED: GATA zinc finger domain-containing protein 6 isof
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>gi|1037071567|ref|XP_017142622.1| PREDICTED: AF4/FMR2 family member 4 isoform X6 [Drosophila r
>gi|1037071517|ref|XP_017142619.1| PREDICTED: uncharacterized protein LOC108155960 isoform X3
>gi|1037071423|ref|XP_017142612.1| PREDICTED: zinc finger matrin-type protein CG9776 isoform X
>gi|1037071439|ref|XP_017142613.1| PREDICTED: zinc finger matrin-type protein CG9776 isoform X2
>gi|1037076263|ref|XP_017142946.1| PREDICTED: uncharacterized protein LOC108156143 [Drosophila
>gi|1037076216|ref|XP_017142944.1| PREDICTED: S-phase kinase-associated protein 2 [Drosophila r
>gi|1037076233|ref|XP_017142945.1| PREDICTED: uncharacterized protein LOC108156141 [Drosophila
>gi|1037103629|ref|XP_017145107.1| PREDICTED: Krueppel-like factor 16 [Drosophila miranda]
>gi|1037100943|ref|XP_017144834.1| PREDICTED: probable sodium/potassium/calcium exchanger CG109
>gi|1037101911|ref|XP_017144922.1| PREDICTED: sideroflexin-1 [Drosophila miranda]
>gi|1037079115|ref|XP_017143148.1| PREDICTED: contactin [Drosophila miranda]
>gi|1037085089|ref|XP_017143592.1| PREDICTED: protein Tube [Drosophila miranda]
>gi|1037085101|ref|XP_017143593.1| PREDICTED: E3 ubiquitin-protein ligase TM129 [Drosophila mi
>gi|1037085115|ref|XP_017143594.1| PREDICTED: uncharacterized protein LOC108156571 [Drosophila
>gi|1037097506|ref|XP_017144521.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 [Drosophila m
>gi|1037097520|ref|XP_017144522.1| PREDICTED: E3 ubiquitin-protein ligase MARCH5 [Drosophila m
>gi|1037022918|ref|XP_017139450.1| PREDICTED: Golgi to ER traffic protein 4 homolog, partial [
>gi|1037093255|ref|XP_017144174.1| PREDICTED: Golgi to ER traffic protein 4 homolog [Drosophila
>gi|1037093269|ref|XP_017144175.1| PREDICTED: Golgi to ER traffic protein 4 homolog [Drosophila
>gi|1037093283|ref|XP_017144177.1| PREDICTED: BTB/POZ domain-containing protein KCTD9 isoform
>gi|1037093297|ref|XP_017144178.1| PREDICTED: BTB/POZ domain-containing protein KCTD9 isoform
>gi|1037080299|ref|XP_017143226.1| PREDICTED: methenyltetrahydrofolate synthase domain-contain
>gi|1037080265|ref|XP_017143224.1| PREDICTED: serine/threonine-protein kinase 32A isoform X1 [
>gi|1037080279|ref|XP_017143225.1| PREDICTED: serine/threonine-protein kinase 32A isoform X2 [
>gi|1037080331|ref|XP_017143228.1| PREDICTED: uncharacterized protein LOC108156340 [Drosophila
>gi|1037080313|ref|XP_017143227.1| PREDICTED: ubiquilin [Drosophila miranda]
>gi|1037060524|ref|XP_017141879.1| PREDICTED: uncharacterized protein LOC108155529 [Drosophila
>gi|1037060551|ref|XP_017141880.1| PREDICTED: uncharacterized protein LOC108155530 isoform X1
>gi|1037060565|ref|XP_017141881.1| PREDICTED: uncharacterized protein LOC108155530 isoform X2
>gi|1037060508|ref|XP_017141878.1| PREDICTED: eukaryotic translation initiation factor 3 subun
>gi|1037060066|ref|XP_017141846.1| PREDICTED: tRNA (guanine(10)-N2)-methyltransferase homolog
>gi|1037060163|ref|XP_017141853.1| PREDICTED: putative lipoyltransferase 2, mitochondrial [Dros
>gi|1037060034|ref|XP_017141844.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037060050|ref|XP_017141845.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037060084|ref|XP_017141847.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060102|ref|XP_017141849.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060118|ref|XP_017141850.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060133|ref|XP_017141851.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037060149|ref|XP_017141852.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078713|ref|XP_017143116.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078725|ref|XP_017143117.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078737|ref|XP_017143118.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037078749|ref|XP_017143119.1| PREDICTED: elongation of very long chain fatty acids protei
>gi|1037069646|ref|XP_017142493.1| PREDICTED: uncharacterized protein LOC108155878 isoform X2
>gi|1037069674|ref|XP_017142494.1| PREDICTED: uncharacterized protein LOC108155878 isoform X2
>gi|1037069690|ref|XP_017142495.1| PREDICTED: uncharacterized protein LOC108155879 [Drosophila
>gi|1037069706|ref|XP_017142496.1| PREDICTED: uncharacterized protein LOC108155879 [Drosophila
>gi|1037069614|ref|XP_017142490.1| PREDICTED: uncharacterized protein LOC108155878 isoform X1
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>gi|1037069632|ref|XP_017142491.1| PREDICTED: uncharacterized protein LOC108155878 isoform X1
>gi|1037048108|ref|XP_017141026.1| PREDICTED: maternal protein pumilio isoform X1 [Drosophila n
>gi|1037048124|ref|XP_017141027.1| PREDICTED: maternal protein pumilio isoform X2 [Drosophila n
>gi|1037048189|ref|XP_017141032.1| PREDICTED: maternal protein pumilio isoform X6 [Drosophila n
>gi|1037048158|ref|XP_017141029.1| PREDICTED: maternal protein pumilio isoform X4 [Drosophila n
>gi|1037048174|ref|XP_017141031.1| PREDICTED: maternal protein pumilio isoform X5 [Drosophila n
>gi|1037048142|ref|XP_017141028.1| PREDICTED: maternal protein pumilio isoform X3 [Drosophila n
>gi|1037048205|ref|XP_017141033.1| PREDICTED: uncharacterized protein LOC108154995 [Drosophila
>gi|1037048221|ref|XP_017141034.1| PREDICTED: uncharacterized protein LOC108154996 [Drosophila
>gi|1037071687|ref|XP_017142632.1| PREDICTED: chromosomal protein D1 [Drosophila miranda]
>gi|1037071701|ref|XP_017142633.1| PREDICTED: chromosomal protein D1 [Drosophila miranda]
>gi|1037071673|ref|XP_017142631.1| PREDICTED: dipeptidyl peptidase 9 [Drosophila miranda]
>gi|1037071712|ref|XP_017142634.1| PREDICTED: acylphosphatase-2 [Drosophila miranda]
>gi|1037039892|ref|XP_017140482.1| PREDICTED: uncharacterized protein LOC108154656 [Drosophila
>gi|1037039878|ref|XP_017140481.1| PREDICTED: putative uncharacterized protein DDB_G0271606 [Dr
>gi|1037039864|ref|XP_017140480.1| PREDICTED: uncharacterized protein LOC108154654 [Drosophila
>gi|1037039756|ref|XP_017140471.1| PREDICTED: HEAT repeat-containing protein 5B isoform X2 [Dro
>gi|1037039740|ref|XP_017140470.1| PREDICTED: HEAT repeat-containing protein 5B isoform X1 [Dro
>gi|1037039772|ref|XP_017140472.1| PREDICTED: HEAT repeat-containing protein 5B isoform X3 [Dro
>gi|1037039914|ref|XP_017140483.1| PREDICTED: uncharacterized protein LOC108154657 [Drosophila
>gi|1037039944|ref|XP_017140485.1| PREDICTED: probable 18S rRNA (guanine-N(7))-methyltransferas
>gi|1037039958|ref|XP_017140486.1| PREDICTED: dnaJ homolog subfamily C member 30 [Drosophila mi
>gi|1037039848|ref|XP_017140479.1| PREDICTED: protein unc-45 homolog B [Drosophila miranda]
>gi|1037039928|ref|XP_017140484.1| PREDICTED: aquaporin-2 [Drosophila miranda]
>gi|1037022936|ref|XP_017139451.1| PREDICTED: uncharacterized protein LOC108153865 [Drosophila
>gi|1037022952|ref|XP_017139453.1| PREDICTED: uncharacterized protein LOC108153866 [Drosophila
>gi|1037039786|ref|XP_017140473.1| PREDICTED: transcription factor mef2A isoform X1 [Drosophila
>gi|1037039800|ref|XP_017140476.1| PREDICTED: transcription factor mef2A isoform X1 [Drosophila
>gi|1037039816|ref|XP_017140477.1| PREDICTED: transcription factor mef2A isoform X1 [Drosophila
>gi|1037039832|ref|XP_017140478.1| PREDICTED: ataxin-2 homolog isoform X2 [Drosophila miranda]
>gi|1037022972|ref|XP_017139454.1| PREDICTED: uncharacterized protein LOC108153867 [Drosophila
>gi|1037022990|ref|XP_017139455.1| PREDICTED: uncharacterized protein LOC108153868 [Drosophila
>gi|1037098096|ref|XP_017144578.1| PREDICTED: uncharacterized protein LOC108157170 [Drosophila
>gi|1037076471|ref|XP_017142959.1| PREDICTED: transforming acidic coiled-coil-containing prote
>gi|1037076437|ref|XP_017142957.1| PREDICTED: transforming acidic coiled-coil-containing prote
>gi|1037076421|ref|XP_017142956.1| PREDICTED: transforming acidic coiled-coil-containing prote
>gi|1037076453|ref|XP_017142958.1| PREDICTED: transforming acidic coiled-coil-containing prote
>gi|1037101075|ref|XP_017144847.1| PREDICTED: membrane metallo-endopeptidase-like 1 isoform X1
>gi|1037101083|ref|XP_017144848.1| PREDICTED: membrane metallo-endopeptidase-like 1 isoform X2
>gi|1037093749|ref|XP_017144207.1| PREDICTED: eukaryotic translation initiation factor 3 subuni
>gi|1037093707|ref|XP_017144203.1| PREDICTED: kelch-like protein 7 isoform X1 [Drosophila miran
>gi|1037093721|ref|XP_017144205.1| PREDICTED: kelch-like protein 7 isoform X2 [Drosophila miran
>gi|1037093735|ref|XP_017144206.1| PREDICTED: uncharacterized protein LOC108156948 isoform X3
>gi|1037069800|ref|XP_017142503.1| PREDICTED: formin-J isoform X2 [Drosophila miranda]
>gi|1037069768|ref|XP_017142500.1| PREDICTED: formin-J isoform X1 [Drosophila miranda]
>gi|1037069784|ref|XP_017142502.1| PREDICTED: formin-J isoform X1 [Drosophila miranda]
>gi|1037069834|ref|XP_017142505.1| PREDICTED: dual specificity protein kinase splA isoform X4
>gi|1037069850|ref|XP_017142506.1| PREDICTED: dual specificity protein kinase splA isoform X5
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>gi|1037069818|ref|XP_017142504.1| PREDICTED: formin-J isoform X3 [Drosophila miranda]
>gi|1037069862|ref|XP_017142507.1| PREDICTED: uncharacterized protein CG43427 isoform X6 [Drosc
>gi|1037023008|ref|XP_017139457.1| PREDICTED: uncharacterized protein LOC108153869 [Drosophila
>gi|1037023025|ref|XP_017139458.1| PREDICTED: uncharacterized protein LOC108153870 [Drosophila
>gi|1037023043|ref|XP_017139459.1| PREDICTED: uncharacterized protein LOC108153871 [Drosophila
>gi|1037104073|ref|XP_017145163.1| PREDICTED: zinc finger protein 878 isoform X1 [Drosophila mi
>gi|1037104081|ref|XP_017145165.1| PREDICTED: oocyte zinc finger protein XlCOF22 isoform X2 [Di
>gi|1037095369|ref|XP_017144332.1| PREDICTED: protein SON [Drosophila miranda]
>gi|1037023061|ref|XP_017139461.1| PREDICTED: uncharacterized protein LOC108153872 [Drosophila
>gi|1037091250|ref|XP_017144035.1| PREDICTED: importin subunit alpha-4 [Drosophila miranda]
>gi|1037102314|ref|XP_017144964.1| PREDICTED: mitochondrial pyruvate carrier 2 [Drosophila mira
>gi|1037106362|ref|XP_017145486.1| PREDICTED: mitochondrial pyruvate carrier 2-like [Drosophila
>gi|1037106370|ref|XP_017145487.1| PREDICTED: mitochondrial pyruvate carrier 2-like [Drosophila
>gi|1037058944|ref|XP_017141766.1| PREDICTED: vacuolar protein sorting-associated protein 45 [I
>gi|1037058914|ref|XP_017141764.1| PREDICTED: IQ and AAA domain-containing protein 1 isoform XI
>gi|1037058930|ref|XP_017141765.1| PREDICTED: IQ and AAA domain-containing protein 1 isoform X2
>gi|1037059051|ref|XP_017141773.1| PREDICTED: U6 snRNA phosphodiesterase [Drosophila miranda]
>gi|1037059021|ref|XP_017141771.1| PREDICTED: metaxin-1 homolog isoform X1 [Drosophila miranda]
>gi|1037059037|ref|XP_017141772.1| PREDICTED: metaxin-1 homolog isoform X2 [Drosophila miranda]
>gi|1037059006|ref|XP_017141770.1| PREDICTED: methyl-CpG-binding domain protein 2 isoform X2 [I
>gi|1037058990|ref|XP_017141769.1| PREDICTED: methyl-CpG-binding domain protein 2 isoform X1 [I
>gi|1037058976|ref|XP_017141768.1| PREDICTED: AP-1 complex subunit mu-1 [Drosophila miranda]
>gi|1037058960|ref|XP_017141767.1| PREDICTED: 2-oxoisovalerate dehydrogenase subunit alpha, mit
>gi|1037092948|ref|XP_017144153.1| PREDICTED: probable tRNA (uracil-O(2)-)-methyltransferase [I
>gi|1037092990|ref|XP_017144157.1| PREDICTED: RNA-binding protein with serine-rich domain 1-B i
>gi|1037092976|ref|XP_017144155.1| PREDICTED: RNA-binding protein with serine-rich domain 1-B i
>gi|1037092962|ref|XP_017144154.1| PREDICTED: RNA-binding protein with serine-rich domain 1-B i
>gi|1037102333|ref|XP_017144966.1| PREDICTED: uncharacterized protein LOC108157428 isoform X1
>gi|1037102343|ref|XP_017144967.1| PREDICTED: uncharacterized protein LOC108157428 isoform X2
>gi|1037081782|ref|XP_017143342.1| PREDICTED: DNA polymerase alpha catalytic subunit [Drosophil
>gi|1037081795|ref|XP_017143343.1| PREDICTED: uncharacterized protein LOC108156409 [Drosophila
>gi|1037063920|ref|XP_017142111.1| PREDICTED: uncharacterized protein LOC108155660 [Drosophila
>gi|1037063933|ref|XP_017142112.1| PREDICTED: protein archease-like [Drosophila miranda]
>gi|1037063858|ref|XP_017142107.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037063873|ref|XP_017142108.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037063889|ref|XP_017142109.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037063906|ref|XP_017142110.1| PREDICTED: transcription factor E2f1 [Drosophila miranda]
>gi|1037023078|ref|XP_017139462.1| PREDICTED: GTPase-activating protein RacGAP84C-like, partial
>gi|1037097794|ref|XP_017144549.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037097784|ref|XP_017144548.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037097803|ref|XP_017144550.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037034374|ref|XP_017140155.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-1
>gi|1037034391|ref|XP_017140156.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-1
>gi|1037035081|ref|XP_017140196.1| PREDICTED: V-type proton ATPase 16 kDa proteolipid subunit-1
>gi|1037062140|ref|XP_017141987.1| PREDICTED: insulin-like receptor [Drosophila miranda]
>gi|1037062156|ref|XP_017141988.1| PREDICTED: insulin-like receptor [Drosophila miranda]
>gi|1037023095|ref|XP_017139463.1| PREDICTED: acyl-CoA Delta(11) desaturase-like [Drosophila mi
>gi|1037096110|ref|XP_017144395.1| PREDICTED: adenosine 3'-phospho 5'-phosphosulfate transporte
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>gi|1037096123|ref|XP_017144396.1| PREDICTED: uncharacterized protein LOC108157072 [Drosophila
>gi|1037038062|ref|XP_017140360.1| PREDICTED: pupal cuticle protein Edg-91 isoform X2 [Drosophi
>gi|1037038050|ref|XP_017140359.1| PREDICTED: pupal cuticle protein Edg-91 isoform X1 [Drosophi
>gi|1037038961|ref|XP_017140419.1| PREDICTED: keratin-associated protein 19-2 [Drosophila mira
>gi|1037037891|ref|XP_017140352.1| PREDICTED: pupal cuticle protein Edg-91 [Drosophila miranda]
>gi|1037040857|ref|XP_017140548.1| PREDICTED: acanthoscurrin-1 [Drosophila miranda]
>gi|1037023113|ref|XP_017139464.1| PREDICTED: translation initiation factor IF-2 [Drosophila m
>gi|1037023131|ref|XP_017139465.1| PREDICTED: uncharacterized protein LOC108153876 [Drosophila
>gi|1037035649|ref|XP_017140221.1| PREDICTED: attacin-A [Drosophila miranda]
>gi|1037106689|ref|XP_017145531.1| PREDICTED: uncharacterized protein LOC108157839 [Drosophila
>gi|1037050510|ref|XP_017141192.1| PREDICTED: uncharacterized protein LOC108155085 isoform X1
>gi|1037050524|ref|XP_017141193.1| PREDICTED: uncharacterized protein LOC108155085 isoform X1
>gi|1037050540|ref|XP_017141194.1| PREDICTED: uncharacterized protein LOC108155085 isoform X2
>gi|1037050634|ref|XP_017141201.1| PREDICTED: bax inhibitor 1-like [Drosophila miranda]
>gi|1037050618|ref|XP_017141200.1| PREDICTED: transmembrane protein 192 [Drosophila miranda]
>gi|1037050602|ref|XP_017141198.1| PREDICTED: serine/threonine-protein kinase shk1/pak1 [Droso
>gi|1037050586|ref|XP_017141197.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037050554|ref|XP_017141195.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037050570|ref|XP_017141196.1| PREDICTED: putative mediator of RNA polymerase II transcript
>gi|1037052014|ref|XP_017141297.1| PREDICTED: splicing factor 3A subunit 3 [Drosophila miranda]
>gi|1037052029|ref|XP_017141298.1| PREDICTED: mitotic spindle assembly checkpoint protein MAD2l
>gi|1037051884|ref|XP_017141288.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1037051900|ref|XP_017141289.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dros
>gi|1037051936|ref|XP_017141291.1| PREDICTED: dihydropyrimidinase isoform X2 [Drosophila miranc
>gi|1037051918|ref|XP_017141290.1| PREDICTED: dihydropyrimidinase isoform X1 [Drosophila miranc
>gi|1037051952|ref|XP_017141292.1| PREDICTED: dihydropyrimidinase 2 isoform X3 [Drosophila mira
>gi|1037051998|ref|XP_017141296.1| PREDICTED: dihydropyrimidinase-related protein 3 isoform X6
>gi|1037051968|ref|XP_017141294.1| PREDICTED: dihydropyrimidinase isoform X4 [Drosophila miranc
>gi|1037051984|ref|XP_017141295.1| PREDICTED: dihydropyrimidinase isoform X5 [Drosophila miranc
>gi|1037032364|ref|XP_017140031.1| PREDICTED: uncharacterized protein LOC108154292 isoform X1
>gi|1037032382|ref|XP_017140032.1| PREDICTED: uncharacterized protein LOC108154292 isoform X2
>gi|1037093311|ref|XP_017144179.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037093325|ref|XP_017144180.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037093339|ref|XP_017144181.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037093353|ref|XP_017144182.1| PREDICTED: uncharacterized protein LOC108156932 [Drosophila
>gi|1037076079|ref|XP_017142934.1| PREDICTED: uncharacterized protein LOC108156132 [Drosophila
>gi|1037076064|ref|XP_017142933.1| PREDICTED: LOW QUALITY PROTEIN: tankyrase [Drosophila miranc
>gi|1037089110|ref|XP_017143866.1| PREDICTED: probable serine/threonine-protein kinase cdc7 [Di
>gi|1037089128|ref|XP_017143867.1| PREDICTED: probable serine/threonine-protein kinase cdc7 [Di
>gi|1037089142|ref|XP_017143868.1| PREDICTED: probable serine/threonine-protein kinase cdc7 [Di
>gi|1037080415|ref|XP_017143235.1| PREDICTED: relaxin receptor 2 [Drosophila miranda]
>gi|1037080471|ref|XP_017143239.1| PREDICTED: zinc finger protein OZF [Drosophila miranda]
>gi|1037080499|ref|XP_017143241.1| PREDICTED: uncharacterized protein LOC108156351 [Drosophila
>gi|1037080457|ref|XP_017143238.1| PREDICTED: S-adenosylmethionine mitochondrial carrier prote:
>gi|1037080443|ref|XP_017143237.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit gamma, mit
>gi|1037080429|ref|XP_017143236.1| PREDICTED: isocitrate dehydrogenase [NAD] subunit gamma, mit
>gi|1037080485|ref|XP_017143240.1| PREDICTED: 60S ribosomal protein L27 [Drosophila miranda]
>gi|1037098387|ref|XP_017144610.1| PREDICTED: probable cardiolipin synthase (CMP-forming) [Dros
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>gi|1037071807|ref|XP_017142640.1| PREDICTED: low-density lipoprotein receptor isoform X3 [Dros
>gi|1037071838|ref|XP_017142641.1| PREDICTED: low-density lipoprotein receptor isoform X4 [Dros
>gi|1037071775|ref|XP_017142638.1| PREDICTED: low-density lipoprotein receptor isoform X1 [Dros
>gi|1037071854|ref|XP_017142642.1| PREDICTED: low-density lipoprotein receptor isoform X5 [Dros
>gi|1037071791|ref|XP_017142639.1| PREDICTED: low-density lipoprotein receptor isoform X2 [Dros
>gi|1037071868|ref|XP_017142643.1| PREDICTED: low-density lipoprotein receptor isoform X6 [Dros
>gi|1037071923|ref|XP_017142645.1| PREDICTED: low-density lipoprotein receptor isoform X8 [Dros
>gi|1037071892|ref|XP_017142644.1| PREDICTED: low-density lipoprotein receptor isoform X7 [Dros
>gi|1037071941|ref|XP_017142646.1| PREDICTED: LOW QUALITY PROTEIN: very low-density lipoprotei
>gi|1037096671|ref|XP_017144445.1| PREDICTED: myosin-9-like [Drosophila miranda]
>gi|1037032969|ref|XP_017140070.1| PREDICTED: probable palmitoyltransferase ZDHHC24 [Drosophila
>gi|1037105991|ref|XP_017145435.1| PREDICTED: uncharacterized protein LOC108157764 [Drosophila
>gi|1037036977|ref|XP_017140299.1| PREDICTED: uncharacterized protein LOC108154509 isoform X2
>gi|1037036960|ref|XP_017140298.1| PREDICTED: uncharacterized protein LOC108154509 isoform X1
>gi|1037106386|ref|XP_017145489.1| PREDICTED: uncharacterized protein LOC108157806 [Drosophila
>gi|1037104422|ref|XP_017145211.1| PREDICTED: calmodulin-like [Drosophila miranda]
>gi|1037104432|ref|XP_017145212.1| PREDICTED: calmodulin-like [Drosophila miranda]
>gi|1037104440|ref|XP_017145213.1| PREDICTED: calmodulin-related protein 97A-like [Drosophila r
>gi|1037104448|ref|XP_017145214.1| PREDICTED: calmodulin-related protein 97A-like [Drosophila r
>gi|1037103667|ref|XP_017145112.1| PREDICTED: protein TolA [Drosophila miranda]
>gi|1037023147|ref|XP_017139466.1| PREDICTED: uncharacterized protein LOC108153877 [Drosophila
>gi|1037094502|ref|XP_017144266.1| PREDICTED: uncharacterized protein LOC108156984 [Drosophila
>gi|1037094516|ref|XP_017144267.1| PREDICTED: uncharacterized protein LOC108156984 [Drosophila
>gi|1037094488|ref|XP_017144264.1| PREDICTED: beta-taxilin [Drosophila miranda]
>gi|1037104804|ref|XP_017145265.1| PREDICTED: Kv channel-interacting protein 1 [Drosophila mira
>gi|1037104812|ref|XP_017145267.1| PREDICTED: Kv channel-interacting protein 1 [Drosophila mira
>gi|1037084074|ref|XP_017143514.1| PREDICTED: CD81 antigen [Drosophila miranda]
>gi|1037064178|ref|XP_017142128.1| PREDICTED: V-type proton ATPase subunit E [Drosophila mirano
>gi|1037064132|ref|XP_017142124.1| PREDICTED: protein krasavietz [Drosophila miranda]
>gi|1037064148|ref|XP_017142125.1| PREDICTED: protein krasavietz [Drosophila miranda]
>gi|1037064164|ref|XP_017142127.1| PREDICTED: uncharacterized protein LOC108155674 [Drosophila
>gi|1037064194|ref|XP_017142129.1| PREDICTED: protein KRTCAP2 homolog [Drosophila miranda]
>gi|1037094708|ref|XP_017144281.1| PREDICTED: 28S ribosomal protein S18a, mitochondrial [Drosop
>gi|1037094692|ref|XP_017144280.1| PREDICTED: pachytene checkpoint protein 2 homolog [Drosophil
>gi|1037023163|ref|XP_017139467.1| PREDICTED: putative odorant receptor 85d [Drosophila miranda
>gi|1037037628|ref|XP_017140338.1| PREDICTED: uncharacterized protein LOC108154546 [Drosophila
>gi|1037106137|ref|XP_017145455.1| PREDICTED: transmembrane protein 135-like [Drosophila mirano
>gi|1037096683|ref|XP_017144446.1| PREDICTED: transmembrane protein 135 [Drosophila miranda]
>gi|1037096697|ref|XP_017144447.1| PREDICTED: transmembrane protein 135 [Drosophila miranda]
>gi|1037037380|ref|XP_017140324.1| PREDICTED: odorant receptor 85c [Drosophila miranda]
>gi|1037106602|ref|XP_017145519.1| PREDICTED: odorant receptor 85b-like [Drosophila miranda]
>gi|1037046571|ref|XP_017140927.1| PREDICTED: irregular chiasm C-roughest protein isoform X3 [D
>gi|1037046539|ref|XP_017140925.1| PREDICTED: uncharacterized protein LOC108154943 isoform X1
>gi|1037046555|ref|XP_017140926.1| PREDICTED: uncharacterized protein LOC108154943 isoform X2
>gi|1037046605|ref|XP_017140929.1| PREDICTED: hemicentin-2 isoform X5 [Drosophila miranda]
>gi|1037046587|ref|XP_017140928.1| PREDICTED: hemicentin-2 isoform X4 [Drosophila miranda]
>gi|1037022499|ref|XP_017139424.1| PREDICTED: uncharacterized protein LOC108153837 [Drosophila
>gi|1037046633|ref|XP_017140931.1| PREDICTED: zinc finger protein 383 isoform X1 [Drosophila mi
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>gi|1037046649|ref|XP_017140932.1| PREDICTED: zinc finger protein 383 isoform X2 [Drosophila m
>gi|1037046663|ref|XP_017140933.1| PREDICTED: adenosine deaminase-like protein [Drosophila mira
>gi|1037046619|ref|XP_017140930.1| PREDICTED: protein Red [Drosophila miranda]
>gi|1037046473|ref|XP_017140920.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037046489|ref|XP_017140921.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037046505|ref|XP_017140922.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037046521|ref|XP_017140924.1| PREDICTED: jmjC domain-containing histone demethylation prot
>gi|1037102426|ref|XP_017144975.1| PREDICTED: vasotab [Drosophila miranda]
>gi|1037102414|ref|XP_017144974.1| PREDICTED: iron-sulfur cluster assembly scaffold protein Is
>gi|1037038601|ref|XP_017140397.1| PREDICTED: UPF0668 protein C10orf76 [Drosophila miranda]
>gi|1037038396|ref|XP_017140383.1| PREDICTED: tight junction protein ZO-1 isoform X1 [Drosophil
>gi|1037038412|ref|XP_017140384.1| PREDICTED: tight junction protein ZO-1 isoform X2 [Drosophil
>gi|1037038430|ref|XP_017140385.1| PREDICTED: tight junction protein ZO-1 isoform X3 [Drosophil
>gi|1037038446|ref|XP_017140386.1| PREDICTED: tight junction protein ZO-1 isoform X4 [Drosophil
>gi|1037038462|ref|XP_017140387.1| PREDICTED: tight junction protein ZO-1 isoform X5 [Drosophil
>gi|1037038478|ref|XP_017140389.1| PREDICTED: uncharacterized protein LOC108154591 isoform X6
>gi|1037038490|ref|XP_017140390.1| PREDICTED: uncharacterized protein LOC108154591 isoform X7
>gi|1037038538|ref|XP_017140393.1| PREDICTED: tight junction protein ZO-1 isoform X10 [Drosoph
>gi|1037038504|ref|XP_017140391.1| PREDICTED: tight junction protein ZO-2 isoform X8 [Drosophil
>gi|1037038520|ref|XP_017140392.1| PREDICTED: tight junction protein ZO-2 isoform X9 [Drosophil
>gi|1037038553|ref|XP_017140394.1| PREDICTED: tight junction protein ZO-2 isoform X11 [Drosoph
>gi|1037038663|ref|XP_017140401.1| PREDICTED: uncharacterized protein LOC108154597 isoform X2
>gi|1037038651|ref|XP_017140400.1| PREDICTED: uncharacterized protein LOC108154597 isoform X1
>gi|1037038619|ref|XP_017140398.1| PREDICTED: maternal effect protein oskar isoform X1 [Drosop
>gi|1037038636|ref|XP_017140399.1| PREDICTED: maternal effect protein oskar isoform X2 [Drosop
>gi|1037038569|ref|XP_017140395.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1037038583|ref|XP_017140396.1| PREDICTED: prominin-like protein [Drosophila miranda]
>gi|1037038772|ref|XP_017140407.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, 
>gi|1037038724|ref|XP_017140404.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, 
>gi|1037038740|ref|XP_017140405.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, 
>gi|1037038756|ref|XP_017140406.1| PREDICTED: succinyl-CoA ligase [ADP-forming] subunit beta, 
>gi|1037038708|ref|XP_017140403.1| PREDICTED: beta-catenin-like protein 1 [Drosophila miranda]
>gi|1037103808|ref|XP_017145130.1| PREDICTED: uncharacterized protein LOC108157538 [Drosophila
>gi|1037078833|ref|XP_017143126.1| PREDICTED: uncharacterized protein LOC108156269 [Drosophila
>gi|1037078847|ref|XP_017143127.1| PREDICTED: uncharacterized protein LOC108156269 [Drosophila
>gi|1037078889|ref|XP_017143130.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
>gi|1037078861|ref|XP_017143128.1| PREDICTED: UPF0528 protein CG10038 isoform X1 [Drosophila m
>gi|1037078875|ref|XP_017143129.1| PREDICTED: UPF0528 protein CG10038 isoform X2 [Drosophila m
>gi|1037104388|ref|XP_017145207.1| PREDICTED: uncharacterized protein LOC108157591 [Drosophila
>gi|1037092345|ref|XP_017144112.1| PREDICTED: steroid receptor seven-up, isoforms B/C isoform 
>gi|1037092271|ref|XP_017144111.1| PREDICTED: steroid receptor seven-up isoform X1 [Drosophila
>gi|1037035011|ref|XP_017140192.1| PREDICTED: uncharacterized protein F12A10.7 [Drosophila mira
>gi|1037105578|ref|XP_017145377.1| PREDICTED: glycerophosphocholine phosphodiesterase GPCPD1 [
>gi|1037033490|ref|XP_017140100.1| PREDICTED: probable cytochrome P450 313a3 [Drosophila miran
>gi|1037105620|ref|XP_017145383.1| PREDICTED: probable cytochrome P450 313a1 [Drosophila miran
>gi|1037105628|ref|XP_017145384.1| PREDICTED: phosphoglycerate mutase 2 [Drosophila miranda]
>gi|1037023179|ref|XP_017139469.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037023195|ref|XP_017139470.1| PREDICTED: chymotrypsin-2 [Drosophila miranda]
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>gi|1037023213|ref|XP_017139471.1| PREDICTED: chymotrypsin-1 [Drosophila miranda]
>gi|1037106780|ref|XP_017145544.1| PREDICTED: LIM domain-binding protein 3 [Drosophila miranda]
>gi|1037106213|ref|XP_017145466.1| PREDICTED: short-chain specific acyl-CoA dehydrogenase, mit
>gi|1037090820|ref|XP_017144003.1| PREDICTED: conserved oligomeric Golgi complex subunit 1 [Dro
>gi|1037090836|ref|XP_017144004.1| PREDICTED: glycine-rich protein 5 [Drosophila miranda]
>gi|1037033984|ref|XP_017140130.1| PREDICTED: 4-coumarate--CoA ligase 1 [Drosophila miranda]
>gi|1037106671|ref|XP_017145529.1| PREDICTED: adenosine kinase [Drosophila miranda]
>gi|1037105844|ref|XP_017145414.1| PREDICTED: eukaryotic translation initiation factor 3 subun
>gi|1037105276|ref|XP_017145334.1| PREDICTED: calphotin [Drosophila miranda]
>gi|1037105852|ref|XP_017145416.1| PREDICTED: mucin-5AC [Drosophila miranda]
>gi|1037074127|ref|XP_017142800.1| PREDICTED: uncharacterized protein LOC108156041 isoform X4
>gi|1037074061|ref|XP_017142795.1| PREDICTED: uncharacterized protein LOC108156041 isoform X1
>gi|1037074077|ref|XP_017142796.1| PREDICTED: uncharacterized protein LOC108156041 isoform X1
>gi|1037074093|ref|XP_017142797.1| PREDICTED: uncharacterized protein LOC108156041 isoform X2
>gi|1037074109|ref|XP_017142798.1| PREDICTED: uncharacterized protein LOC108156041 isoform X3
>gi|1037074029|ref|XP_017142793.1| PREDICTED: uncharacterized protein LOC108156040 [Drosophila
>gi|1037074045|ref|XP_017142794.1| PREDICTED: uncharacterized protein LOC108156040 [Drosophila
>gi|1037073931|ref|XP_017142786.1| PREDICTED: phospholipid-transporting ATPase ID isoform X5 [
>gi|1037073947|ref|XP_017142787.1| PREDICTED: phospholipid-transporting ATPase ID isoform X6 [
>gi|1037073979|ref|XP_017142790.1| PREDICTED: phospholipid-transporting ATPase ID isoform X8 [
>gi|1037073995|ref|XP_017142791.1| PREDICTED: phospholipid-transporting ATPase ID isoform X9 [
>gi|1037074011|ref|XP_017142792.1| PREDICTED: phospholipid-transporting ATPase ID isoform X10 [
>gi|1037073839|ref|XP_017142780.1| PREDICTED: phospholipid-transporting ATPase ID isoform X1 [
>gi|1037073854|ref|XP_017142781.1| PREDICTED: phospholipid-transporting ATPase ID isoform X1 [
>gi|1037073868|ref|XP_017142782.1| PREDICTED: phospholipid-transporting ATPase ID isoform X1 [
>gi|1037073883|ref|XP_017142783.1| PREDICTED: phospholipid-transporting ATPase ID isoform X2 [
>gi|1037073915|ref|XP_017142785.1| PREDICTED: phospholipid-transporting ATPase ID isoform X4 [
>gi|1037073899|ref|XP_017142784.1| PREDICTED: phospholipid-transporting ATPase ID isoform X3 [
>gi|1037073963|ref|XP_017142788.1| PREDICTED: phospholipid-transporting ATPase ID isoform X7 [
>gi|1037105941|ref|XP_017145428.1| PREDICTED: probable citrate synthase, mitochondrial [Drosopl
>gi|1037035169|ref|XP_017140201.1| PREDICTED: ubiquitin-conjugating enzyme E2 6 [Drosophila mi
>gi|1037102021|ref|XP_017144934.1| PREDICTED: band 7 protein AGAP004871 [Drosophila miranda]
>gi|1037101966|ref|XP_017144929.1| PREDICTED: uncharacterized protein LOC108157399 [Drosophila
>gi|1037101978|ref|XP_017144930.1| PREDICTED: uncharacterized protein LOC108157400 isoform X1
>gi|1037101991|ref|XP_017144931.1| PREDICTED: band 7 protein AGAP004871-like isoform X2 [Drosop
>gi|1037102002|ref|XP_017144932.1| PREDICTED: band 7 protein AGAP004871-like isoform X3 [Drosop
>gi|1037102011|ref|XP_017144933.1| PREDICTED: band 7 protein AGAP004871-like isoform X4 [Drosop
>gi|1037102031|ref|XP_017144935.1| PREDICTED: uncharacterized protein LOC108157402 [Drosophila
>gi|1037100552|ref|XP_017144793.1| PREDICTED: transcription initiation protein SPT3 homolog [Di
>gi|1037076930|ref|XP_017142990.1| PREDICTED: DDB1- and CUL4-associated factor 12 [Drosophila n
>gi|1037076915|ref|XP_017142989.1| PREDICTED: GRIP and coiled-coil domain-containing protein 2
>gi|1037094722|ref|XP_017144282.1| PREDICTED: kelch-like protein 18 [Drosophila miranda]
>gi|1037058204|ref|XP_017141715.1| PREDICTED: beta-galactosidase [Drosophila miranda]
>gi|1037058251|ref|XP_017141719.1| PREDICTED: tachykinins isoform X1 [Drosophila miranda]
>gi|1037058265|ref|XP_017141720.1| PREDICTED: tachykinins isoform X2 [Drosophila miranda]
>gi|1037058147|ref|XP_017141711.1| PREDICTED: transforming growth factor-beta-induced protein
>gi|1037058162|ref|XP_017141712.1| PREDICTED: transforming growth factor-beta-induced protein
>gi|1037058116|ref|XP_017141709.1| PREDICTED: transforming growth factor-beta-induced protein
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>gi|1037058132|ref|XP_017141710.1| PREDICTED: transforming growth factor-beta-induced protein
>gi|1037058177|ref|XP_017141713.1| PREDICTED: transforming growth factor-beta-induced protein
>gi|1037058187|ref|XP_017141714.1| PREDICTED: transforming growth factor-beta-induced protein
>gi|1037058236|ref|XP_017141718.1| PREDICTED: L-galactose dehydrogenase [Drosophila miranda]
>gi|1037058218|ref|XP_017141717.1| PREDICTED: L-galactose dehydrogenase [Drosophila miranda]
>gi|1037058279|ref|XP_017141721.1| PREDICTED: uncharacterized protein LOC108155425 [Drosophila
>gi|1037032056|ref|XP_017140013.1| PREDICTED: solute carrier family 35 member F6 [Drosophila mi
>gi|1037032119|ref|XP_017140018.1| PREDICTED: mitochondrial glutamate carrier 1 [Drosophila mir
>gi|1037032136|ref|XP_017140019.1| PREDICTED: mitochondrial glutamate carrier 1-like [Drosophil
>gi|1037032092|ref|XP_017140015.1| PREDICTED: mitochondrial glutamate carrier 1-like [Drosophil
>gi|1037032106|ref|XP_017140017.1| PREDICTED: mitochondrial glutamate carrier 1-like [Drosophil
>gi|1037031807|ref|XP_017139998.1| PREDICTED: uncharacterized protein LOC108154271 isoform X1
>gi|1037031841|ref|XP_017140000.1| PREDICTED: protein IWS1 homolog isoform X3 [Drosophila mira
>gi|1037031823|ref|XP_017139999.1| PREDICTED: uncharacterized protein LOC108154271 isoform X2
>gi|1037032037|ref|XP_017140012.1| PREDICTED: aurora kinase C [Drosophila miranda]
>gi|1037031773|ref|XP_017139996.1| PREDICTED: zinc finger protein 829 isoform X1 [Drosophila mi
>gi|1037031791|ref|XP_017139997.1| PREDICTED: uncharacterized protein LOC108154270 isoform X2
>gi|1037031612|ref|XP_017139991.1| PREDICTED: arginine/serine-rich protein PNISR [Drosophila mi
>gi|1037031648|ref|XP_017139994.1| PREDICTED: ATPase family AAA domain-containing protein 3 [D
>gi|1037031757|ref|XP_017139995.1| PREDICTED: ATPase family AAA domain-containing protein 3 [D
>gi|1037031629|ref|XP_017139992.1| PREDICTED: protein asunder [Drosophila miranda]
>gi|1037032074|ref|XP_017140014.1| PREDICTED: uncharacterized protein LOC108154276 [Drosophila
>gi|1037031596|ref|XP_017139990.1| PREDICTED: nucleolar protein 6 [Drosophila miranda]
>gi|1037031954|ref|XP_017140007.1| PREDICTED: casein kinase I isoform X6 [Drosophila miranda]
>gi|1037031970|ref|XP_017140008.1| PREDICTED: casein kinase I isoform X6 [Drosophila miranda]
>gi|1037031886|ref|XP_017140003.1| PREDICTED: casein kinase I isoform X2 [Drosophila miranda]
>gi|1037031936|ref|XP_017140006.1| PREDICTED: casein kinase I isoform X5 [Drosophila miranda]
>gi|1037031986|ref|XP_017140009.1| PREDICTED: casein kinase I isoform X7 [Drosophila miranda]
>gi|1037031869|ref|XP_017140001.1| PREDICTED: casein kinase I isoform X1 [Drosophila miranda]
>gi|1037031922|ref|XP_017140005.1| PREDICTED: casein kinase I isoform X4 [Drosophila miranda]
>gi|1037031904|ref|XP_017140004.1| PREDICTED: casein kinase I isoform X3 [Drosophila miranda]
>gi|1037032001|ref|XP_017140010.1| PREDICTED: casein kinase I isoform X8 [Drosophila miranda]
>gi|1037032019|ref|XP_017140011.1| PREDICTED: casein kinase I isoform X9 [Drosophila miranda]
>gi|1037101928|ref|XP_017144924.1| PREDICTED: zinc transporter ZIP1 [Drosophila miranda]
>gi|1037082760|ref|XP_017143412.1| PREDICTED: extracellular sulfatase SULF-1 homolog [Drosophil
>gi|1037082774|ref|XP_017143413.1| PREDICTED: extracellular sulfatase SULF-1 homolog [Drosophil
>gi|1037082801|ref|XP_017143415.1| PREDICTED: facilitated trehalose transporter Tret1-like [Dro
>gi|1037082788|ref|XP_017143414.1| PREDICTED: probable plastidic glucose transporter 2 [Drosoph
>gi|1037073595|ref|XP_017142764.1| PREDICTED: serine/arginine-rich splicing factor 1B [Drosophi
>gi|1037073611|ref|XP_017142765.1| PREDICTED: transcription factor Sp4 [Drosophila miranda]
>gi|1037073661|ref|XP_017142768.1| PREDICTED: uncharacterized protein LOC108156027 isoform X3
>gi|1037073627|ref|XP_017142766.1| PREDICTED: uncharacterized protein LOC108156027 isoform X1
>gi|1037073643|ref|XP_017142767.1| PREDICTED: uncharacterized protein LOC108156027 isoform X2
>gi|1037097565|ref|XP_017144527.1| PREDICTED: mesencephalic astrocyte-derived neurotrophic fact
>gi|1037097575|ref|XP_017144528.1| PREDICTED: uncharacterized protein LOC108157138 [Drosophila
>gi|1037102456|ref|XP_017144979.1| PREDICTED: putative SERF-like protein [Drosophila miranda]
>gi|1037104768|ref|XP_017145260.1| PREDICTED: uncharacterized protein LOC108157631 [Drosophila
>gi|1037104760|ref|XP_017145259.1| PREDICTED: uncharacterized protein LOC108157630 [Drosophila
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>gi|1037105570|ref|XP_017145376.1| PREDICTED: uncharacterized protein LOC108157719 [Drosophila
>gi|1037102165|ref|XP_017144948.1| PREDICTED: circadian locomoter output cycles protein kaput
>gi|1037102144|ref|XP_017144946.1| PREDICTED: single-minded homolog 2 isoform X1 [Drosophila m
>gi|1037102154|ref|XP_017144947.1| PREDICTED: single-minded homolog 2 isoform X1 [Drosophila m
>gi|1037033470|ref|XP_017140099.1| PREDICTED: uncharacterized protein LOC108154344 isoform X4
>gi|1037033414|ref|XP_017140095.1| PREDICTED: uncharacterized protein LOC108154344 isoform X1
>gi|1037033452|ref|XP_017140098.1| PREDICTED: uncharacterized protein LOC108154344 isoform X3
>gi|1037033434|ref|XP_017140097.1| PREDICTED: uncharacterized protein LOC108154344 isoform X2
>gi|1037050839|ref|XP_017141217.1| PREDICTED: protein farnesyltransferase subunit beta [Drosop
>gi|1037050883|ref|XP_017141220.1| PREDICTED: uncharacterized protein LOC108155102 isoform X2
>gi|1037050871|ref|XP_017141219.1| PREDICTED: armadillo repeat-containing protein 7 isoform X1
>gi|1037050901|ref|XP_017141221.1| PREDICTED: 28S ribosomal protein S33, mitochondrial [Drosop
>gi|1037050793|ref|XP_017141212.1| PREDICTED: protein CLEC16A isoform X2 [Drosophila miranda]
>gi|1037050777|ref|XP_017141211.1| PREDICTED: protein CLEC16A isoform X1 [Drosophila miranda]
>gi|1037050855|ref|XP_017141218.1| PREDICTED: tetratricopeptide repeat protein 1 [Drosophila m
>gi|1037050825|ref|XP_017141216.1| PREDICTED: methionine synthase reductase [Drosophila miranda
>gi|1037050809|ref|XP_017141215.1| PREDICTED: serine/threonine-protein kinase PAK 2 [Drosophila
>gi|1037054821|ref|XP_017141488.1| PREDICTED: techylectin-5A [Drosophila miranda]
>gi|1037054761|ref|XP_017141484.1| PREDICTED: uncharacterized protein LOC108155281 isoform X4
>gi|1037054746|ref|XP_017141483.1| PREDICTED: uncharacterized protein LOC108155281 isoform X3
>gi|1037054732|ref|XP_017141482.1| PREDICTED: uncharacterized protein LOC108155281 isoform X2
>gi|1037054697|ref|XP_017141479.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037054714|ref|XP_017141481.1| PREDICTED: mucin-5AC isoform X1 [Drosophila miranda]
>gi|1037054789|ref|XP_017141486.1| PREDICTED: mucin-5AC isoform X6 [Drosophila miranda]
>gi|1037054777|ref|XP_017141485.1| PREDICTED: mucin-5AC isoform X5 [Drosophila miranda]
>gi|1037054852|ref|XP_017141491.1| PREDICTED: acylphosphatase-2 [Drosophila miranda]
>gi|1037054836|ref|XP_017141490.1| PREDICTED: uroporphyrinogen-III synthase [Drosophila miranda
>gi|1037054805|ref|XP_017141487.1| PREDICTED: protein FAM114A2 [Drosophila miranda]
>gi|1037036661|ref|XP_017140282.1| PREDICTED: uncharacterized protein LOC108154493 [Drosophila
>gi|1037026831|ref|XP_017139698.1| PREDICTED: uncharacterized protein LOC108154093 isoform X1
>gi|1037026849|ref|XP_017139699.1| PREDICTED: uncharacterized protein LOC108154093 isoform X2
>gi|1037027074|ref|XP_017139713.1| PREDICTED: uncharacterized protein LOC108154102 [Drosophila
>gi|1037027164|ref|XP_017139719.1| PREDICTED: syntaxin-18 [Drosophila miranda]
>gi|1037027281|ref|XP_017139727.1| PREDICTED: ras-like GTP-binding protein RhoL [Drosophila mi
>gi|1037027213|ref|XP_017139722.1| PREDICTED: uncharacterized protein LOC108154110 isoform X1
>gi|1037027230|ref|XP_017139724.1| PREDICTED: SAP domain-containing ribonucleoprotein isoform X
>gi|1037027025|ref|XP_017139710.1| PREDICTED: heterogeneous nuclear ribonucleoprotein M isoform
>gi|1037027040|ref|XP_017139711.1| PREDICTED: glycine-rich RNA-binding protein isoform X2 [Dros
>gi|1037027299|ref|XP_017139728.1| PREDICTED: ras-like protein 1 [Drosophila miranda]
>gi|1037027147|ref|XP_017139718.1| PREDICTED: serum response factor-binding protein 1 [Drosophi
>gi|1037027264|ref|XP_017139726.1| PREDICTED: 39S ribosomal protein L47, mitochondrial [Drosop
>gi|1037026957|ref|XP_017139705.1| PREDICTED: lysine-specific demethylase 3B [Drosophila miranc
>gi|1037026975|ref|XP_017139706.1| PREDICTED: lysine-specific demethylase 3B [Drosophila miranc
>gi|1037026866|ref|XP_017139700.1| PREDICTED: F-BAR domain only protein 2 isoform X1 [Drosophil
>gi|1037026881|ref|XP_017139701.1| PREDICTED: F-BAR domain only protein 2 isoform X1 [Drosophil
>gi|1037026899|ref|XP_017139702.1| PREDICTED: F-BAR domain only protein 2 isoform X1 [Drosophil
>gi|1037026915|ref|XP_017139703.1| PREDICTED: F-BAR domain only protein 2 isoform X2 [Drosophil
>gi|1037027010|ref|XP_017139709.1| PREDICTED: tensin [Drosophila miranda]
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>gi|1037026993|ref|XP_017139708.1| PREDICTED: solute carrier family 22 member 3 [Drosophila mi
>gi|1037027092|ref|XP_017139714.1| PREDICTED: RNA polymerase II-associated factor 1 homolog [D
>gi|1037027112|ref|XP_017139716.1| PREDICTED: zinc finger protein 628 [Drosophila miranda]
>gi|1037026933|ref|XP_017139704.1| PREDICTED: coiled-coil domain-containing protein 39 [Drosop
>gi|1037026813|ref|XP_017139697.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X5 [D
>gi|1037026796|ref|XP_017139696.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X4 [D
>gi|1037026776|ref|XP_017139695.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X3 [D
>gi|1037026758|ref|XP_017139693.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X2 [D
>gi|1037026724|ref|XP_017139691.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X1 [D
>gi|1037026740|ref|XP_017139692.1| PREDICTED: E3 ubiquitin-protein ligase TRIP12 isoform X1 [D
>gi|1037027249|ref|XP_017139725.1| PREDICTED: checkpoint protein HUS1 [Drosophila miranda]
>gi|1037027195|ref|XP_017139721.1| PREDICTED: mannose-1-phosphate guanyltransferase beta [Dros
>gi|1037027180|ref|XP_017139720.1| PREDICTED: Bardet-Biedl syndrome 5 protein homolog [Drosoph
>gi|1037027130|ref|XP_017139717.1| PREDICTED: serine/arginine repetitive matrix protein 2 [Dro
>gi|1037027317|ref|XP_017139729.1| PREDICTED: uncharacterized protein LOC108154115 [Drosophila
>gi|1037027335|ref|XP_017139730.1| PREDICTED: uncharacterized protein LOC108154116 [Drosophila
>gi|1037027057|ref|XP_017139712.1| PREDICTED: pair-rule protein odd-paired [Drosophila miranda
>gi|1037105559|ref|XP_017145375.1| PREDICTED: uncharacterized protein LOC108157716 [Drosophila
>gi|1037040931|ref|XP_017140554.1| PREDICTED: LOW QUALITY PROTEIN: replication factor C subuni
>gi|1037040873|ref|XP_017140550.1| PREDICTED: FERM, RhoGEF and pleckstrin domain-containing pr
>gi|1037040887|ref|XP_017140551.1| PREDICTED: FERM, RhoGEF and pleckstrin domain-containing pr
>gi|1037040903|ref|XP_017140552.1| PREDICTED: microtubule-associated protein futsch isoform X3
>gi|1037040917|ref|XP_017140553.1| PREDICTED: ubiquitin-conjugating enzyme E2-17 kDa [Drosophi
>gi|1037023229|ref|XP_017139472.1| PREDICTED: uncharacterized protein LOC108153883, partial [D
>gi|1037023247|ref|XP_017139473.1| PREDICTED: lachesin, partial [Drosophila miranda]
>gi|1037105292|ref|XP_017145336.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037106432|ref|XP_017145496.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037105268|ref|XP_017145332.1| PREDICTED: uncharacterized protein LOC108157678 [Drosophila
>gi|1037106594|ref|XP_017145518.1| PREDICTED: uncharacterized protein LOC108157828 [Drosophila
>gi|1037104929|ref|XP_017145284.1| PREDICTED: uncharacterized protein LOC108157646 [Drosophila
>gi|1037104561|ref|XP_017145231.1| PREDICTED: mpv17-like protein [Drosophila miranda]
>gi|1037104576|ref|XP_017145233.1| PREDICTED: mpv17-like protein [Drosophila miranda]
>gi|1037081808|ref|XP_017143344.1| PREDICTED: centrosomal and chromosomal factor [Drosophila m
>gi|1037081819|ref|XP_017143345.1| PREDICTED: uncharacterized protein LOC108156411 [Drosophila
>gi|1037106711|ref|XP_017145534.1| PREDICTED: 4-coumarate--CoA ligase 1-like [Drosophila miran
>gi|1037102251|ref|XP_017144958.1| PREDICTED: LOW QUALITY PROTEIN: luciferin 4-monooxygenase-l
>gi|1037102261|ref|XP_017144960.1| PREDICTED: luciferin 4-monooxygenase-like [Drosophila miran
>gi|1037073267|ref|XP_017142741.1| PREDICTED: LOW QUALITY PROTEIN: geranylgeranyl transferase
>gi|1037073251|ref|XP_017142740.1| PREDICTED: protein argonaute-2-like [Drosophila miranda]
>gi|1037103413|ref|XP_017145079.1| PREDICTED: uncharacterized protein LOC108157494 isoform X1
>gi|1037103432|ref|XP_017145081.1| PREDICTED: uncharacterized protein LOC108157495 isoform X1
>gi|1037103444|ref|XP_017145082.1| PREDICTED: uncharacterized protein LOC108157495 isoform X2
>gi|1037103422|ref|XP_017145080.1| PREDICTED: uncharacterized protein LOC108157494 isoform X2
>gi|1037058322|ref|XP_017141724.1| PREDICTED: afadin isoform X2 [Drosophila miranda]
>gi|1037058308|ref|XP_017141722.1| PREDICTED: afadin isoform X1 [Drosophila miranda]
>gi|1037058338|ref|XP_017141725.1| PREDICTED: uncharacterized protein LOC108155427 isoform X3
>gi|1037101251|ref|XP_017144861.1| PREDICTED: RAB6A-GEF complex partner protein 2 [Drosophila
>gi|1037054219|ref|XP_017141446.1| PREDICTED: vacuolar protein sorting-associated protein 37B
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>gi|1037054170|ref|XP_017141443.1| PREDICTED: katanin p60 ATPase-containing subunit A-like 1 is
>gi|1037054188|ref|XP_017141444.1| PREDICTED: katanin p60 ATPase-containing subunit A-like 1 is
>gi|1037054123|ref|XP_017141439.1| PREDICTED: MMS19 nucleotide excision repair protein homolog
>gi|1037054203|ref|XP_017141445.1| PREDICTED: MORN repeat-containing protein 4 [Drosophila mira
>gi|1037054154|ref|XP_017141441.1| PREDICTED: putative cysteine proteinase CG12163 isoform X2
>gi|1037054139|ref|XP_017141440.1| PREDICTED: putative cysteine proteinase CG12163 isoform X1
>gi|1037054235|ref|XP_017141447.1| PREDICTED: uncharacterized protein LOC108155260 [Drosophila
>gi|1037100842|ref|XP_017144824.1| PREDICTED: uncharacterized protein LOC108157320 [Drosophila
>gi|1037100850|ref|XP_017144825.1| PREDICTED: uncharacterized protein LOC108157320 [Drosophila
>gi|1037100860|ref|XP_017144826.1| PREDICTED: enolase-phosphatase E1 [Drosophila miranda]
>gi|1037046300|ref|XP_017140908.1| PREDICTED: egl nine homolog 1 isoform X1 [Drosophila miranda
>gi|1037046330|ref|XP_017140910.1| PREDICTED: probable mitochondrial import inner membrane tra
>gi|1037046314|ref|XP_017140909.1| PREDICTED: egl nine homolog 1 isoform X2 [Drosophila miranda
>gi|1037046098|ref|XP_017140892.1| PREDICTED: oxidation resistance protein 1 isoform X9 [Droso
>gi|1037046082|ref|XP_017140891.1| PREDICTED: oxidation resistance protein 1 isoform X8 [Droso
>gi|1037046244|ref|XP_017140904.1| PREDICTED: oxidation resistance protein 1 isoform X20 [Dros
>gi|1037046064|ref|XP_017140890.1| PREDICTED: oxidation resistance protein 1 isoform X7 [Droso
>gi|1037046271|ref|XP_017140906.1| PREDICTED: oxidation resistance protein 1 isoform X22 [Dros
>gi|1037046046|ref|XP_017140889.1| PREDICTED: oxidation resistance protein 1 isoform X6 [Droso
>gi|1037046258|ref|XP_017140905.1| PREDICTED: oxidation resistance protein 1 isoform X21 [Dros
>gi|1037046003|ref|XP_017140886.1| PREDICTED: oxidation resistance protein 1 isoform X3 [Droso
>gi|1037046033|ref|XP_017140888.1| PREDICTED: oxidation resistance protein 1 isoform X5 [Droso
>gi|1037045987|ref|XP_017140885.1| PREDICTED: oxidation resistance protein 1 isoform X2 [Droso
>gi|1037046232|ref|XP_017140903.1| PREDICTED: uncharacterized protein LOC108154930 isoform X19
>gi|1037046017|ref|XP_017140887.1| PREDICTED: oxidation resistance protein 1 isoform X4 [Droso
>gi|1037045967|ref|XP_017140884.1| PREDICTED: oxidation resistance protein 1 isoform X1 [Droso
>gi|1037046154|ref|XP_017140896.1| PREDICTED: oxidation resistance protein 1 isoform X13 [Dros
>gi|1037046128|ref|XP_017140894.1| PREDICTED: oxidation resistance protein 1 isoform X11 [Dros
>gi|1037046142|ref|XP_017140895.1| PREDICTED: oxidation resistance protein 1 isoform X12 [Dros
>gi|1037046114|ref|XP_017140893.1| PREDICTED: oxidation resistance protein 1 isoform X10 [Dros
>gi|1037046178|ref|XP_017140899.1| PREDICTED: oxidation resistance protein 1 isoform X15 [Dros
>gi|1037046286|ref|XP_017140907.1| PREDICTED: oxidation resistance protein 1 isoform X23 [Dros
>gi|1037046164|ref|XP_017140897.1| PREDICTED: oxidation resistance protein 1 isoform X14 [Dros
>gi|1037046204|ref|XP_017140901.1| PREDICTED: TLD domain-containing protein 2 isoform X17 [Dros
>gi|1037046190|ref|XP_017140900.1| PREDICTED: TLD domain-containing protein 2 isoform X16 [Dros
>gi|1037046218|ref|XP_017140902.1| PREDICTED: TLD domain-containing protein 2 isoform X18 [Dros
>gi|1037046347|ref|XP_017140911.1| PREDICTED: probable mitochondrial import inner membrane tra
>gi|1037046363|ref|XP_017140912.1| PREDICTED: probable mitochondrial import inner membrane tra
>gi|1037089816|ref|XP_017143922.1| PREDICTED: stromal cell-derived factor 2 [Drosophila miranda
>gi|1037089802|ref|XP_017143921.1| PREDICTED: uncharacterized protein CG1161 [Drosophila miran
>gi|1037089788|ref|XP_017143920.1| PREDICTED: proteasome subunit beta type-4 [Drosophila miran
>gi|1037096204|ref|XP_017144404.1| PREDICTED: ceramide kinase [Drosophila miranda]
>gi|1037096215|ref|XP_017144405.1| PREDICTED: ceramide kinase [Drosophila miranda]
>gi|1037105284|ref|XP_017145335.1| PREDICTED: transcription factor SPT20 homolog [Drosophila m
>gi|1037105516|ref|XP_017145370.1| PREDICTED: uncharacterized protein LOC108157710 [Drosophila
>gi|1037076786|ref|XP_017142980.1| PREDICTED: DNA-directed RNA polymerases I, II, and III subu
>gi|1037076726|ref|XP_017142976.1| PREDICTED: protein downstream neighbor of son homolog [Dros
>gi|1037076757|ref|XP_017142978.1| PREDICTED: zinc finger protein 17 [Drosophila miranda]
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>gi|1037076770|ref|XP_017142979.1| PREDICTED: neuroendocrine protein 7B2 [Drosophila miranda]
>gi|1037076708|ref|XP_017142975.1| PREDICTED: uncharacterized protein LOC108156158 isoform X2
>gi|1037076662|ref|XP_017142971.1| PREDICTED: uncharacterized protein LOC108156158 isoform X1
>gi|1037076678|ref|XP_017142972.1| PREDICTED: uncharacterized protein LOC108156158 isoform X1
>gi|1037076694|ref|XP_017142974.1| PREDICTED: uncharacterized protein LOC108156158 isoform X1
>gi|1037076742|ref|XP_017142977.1| PREDICTED: uncharacterized protein LOC108156160 [Drosophila
>gi|1037027607|ref|XP_017139746.1| PREDICTED: homeobox protein 5 [Drosophila miranda]
>gi|1037027624|ref|XP_017139747.1| PREDICTED: homeobox protein 5 [Drosophila miranda]
>gi|1037027817|ref|XP_017139761.1| PREDICTED: derlin-2 [Drosophila miranda]
>gi|1037027961|ref|XP_017139768.1| PREDICTED: succinate dehydrogenase assembly factor 3, mitoch
>gi|1037027943|ref|XP_017139767.1| PREDICTED: dihydrofolate reductase [Drosophila miranda]
>gi|1037027575|ref|XP_017139744.1| PREDICTED: soluble guanylate cyclase 89Db [Drosophila mirand
>gi|1037027590|ref|XP_017139745.1| PREDICTED: soluble guanylate cyclase 89Db [Drosophila mirand
>gi|1037027558|ref|XP_017139743.1| PREDICTED: soluble guanylate cyclase 89Da [Drosophila mirand
>gi|1037027800|ref|XP_017139759.1| PREDICTED: COP9 signalosome complex subunit 5 [Drosophila mi
>gi|1037027381|ref|XP_017139732.1| PREDICTED: titin isoform X1 [Drosophila miranda]
>gi|1037027403|ref|XP_017139733.1| PREDICTED: microtubule-associated protein futsch isoform X2
>gi|1037027755|ref|XP_017139756.1| PREDICTED: glycerophosphodiester phosphodiesterase 1 [Drosop
>gi|1037027769|ref|XP_017139757.1| PREDICTED: glycerophosphodiester phosphodiesterase 1 [Drosop
>gi|1037027785|ref|XP_017139758.1| PREDICTED: glycerophosphodiester phosphodiesterase 1 [Drosop
>gi|1037027835|ref|XP_017139762.1| PREDICTED: trypsin-4 [Drosophila miranda]
>gi|1037027421|ref|XP_017139734.1| PREDICTED: alpha-mannosidase 2x isoform X1 [Drosophila miran
>gi|1037027438|ref|XP_017139735.1| PREDICTED: alpha-mannosidase 2x isoform X2 [Drosophila miran
>gi|1037027456|ref|XP_017139736.1| PREDICTED: alpha-mannosidase 2x isoform X2 [Drosophila miran
>gi|1037027488|ref|XP_017139738.1| PREDICTED: DNA replication licensing factor REC isoform X1
>gi|1037027706|ref|XP_017139753.1| PREDICTED: nose resistant to fluoxetine protein 6 [Drosophil
>gi|1037027506|ref|XP_017139740.1| PREDICTED: DNA replication licensing factor REC isoform X2
>gi|1037027524|ref|XP_017139741.1| PREDICTED: DNA replication licensing factor REC isoform X3
>gi|1037027851|ref|XP_017139763.1| PREDICTED: uncharacterized protein LOC108154138 [Drosophila
>gi|1037027641|ref|XP_017139749.1| PREDICTED: pickpocket protein 28 [Drosophila miranda]
>gi|1037027473|ref|XP_017139737.1| PREDICTED: importin-5 [Drosophila miranda]
>gi|1037027658|ref|XP_017139750.1| PREDICTED: protein spartin [Drosophila miranda]
>gi|1037027673|ref|XP_017139751.1| PREDICTED: protein spartin [Drosophila miranda]
>gi|1037027690|ref|XP_017139752.1| PREDICTED: dnaJ homolog subfamily C member 3 [Drosophila mi
>gi|1037027738|ref|XP_017139755.1| PREDICTED: otefin [Drosophila miranda]
>gi|1037027541|ref|XP_017139742.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037027871|ref|XP_017139764.1| PREDICTED: GILT-like protein F37H8.5 isoform X1 [Drosophila
>gi|1037027885|ref|XP_017139765.1| PREDICTED: GILT-like protein F37H8.5 isoform X2 [Drosophila
>gi|1037027925|ref|XP_017139766.1| PREDICTED: uncharacterized protein LOC108154140 [Drosophila
>gi|1037027721|ref|XP_017139754.1| PREDICTED: zinc finger protein 664 [Drosophila miranda]
>gi|1037096911|ref|XP_017144466.1| PREDICTED: ras-related GTP-binding protein A [Drosophila mi
>gi|1037096922|ref|XP_017144467.1| PREDICTED: beta carbonic anhydrase 1 [Drosophila miranda]
>gi|1037093506|ref|XP_017144194.1| PREDICTED: ubiquitin domain-containing protein 1 [Drosophila
>gi|1037093575|ref|XP_017144195.1| PREDICTED: ubiquitin domain-containing protein 1 [Drosophila
>gi|1037023263|ref|XP_017139475.1| PREDICTED: odorant receptor 83a [Drosophila miranda]
>gi|1037032414|ref|XP_017140035.1| PREDICTED: alpha-tocopherol transfer protein-like [Drosophil
>gi|1037104858|ref|XP_017145273.1| PREDICTED: odorant receptor coreceptor [Drosophila miranda]
>gi|1037106800|ref|XP_017145547.1| PREDICTED: casein kinase I [Drosophila miranda]
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>gi|1037075783|ref|XP_017142915.1| PREDICTED: LOW QUALITY PROTEIN: ras-like protein family memb
>gi|1037075749|ref|XP_017142913.1| PREDICTED: eukaryotic translation initiation factor 2-alpha
>gi|1037075767|ref|XP_017142914.1| PREDICTED: eukaryotic translation initiation factor 2-alpha
>gi|1037075799|ref|XP_017142916.1| PREDICTED: 60S ribosomal protein L35a [Drosophila miranda]
>gi|1037054356|ref|XP_017141457.1| PREDICTED: polymerase delta-interacting protein 2 [Drosophil
>gi|1037054383|ref|XP_017141459.1| PREDICTED: 39S ribosomal protein L44, mitochondrial [Drosoph
>gi|1037054368|ref|XP_017141458.1| PREDICTED: SWI/SNF-related matrix-associated actin-dependent
>gi|1037054340|ref|XP_017141456.1| PREDICTED: histone deacetylase 3 [Drosophila miranda]
>gi|1037054251|ref|XP_017141448.1| PREDICTED: biotin--protein ligase isoform X1 [Drosophila mir
>gi|1037054281|ref|XP_017141449.1| PREDICTED: biotin--protein ligase isoform X2 [Drosophila mir
>gi|1037054293|ref|XP_017141450.1| PREDICTED: biotin--protein ligase isoform X3 [Drosophila mir
>gi|1037054313|ref|XP_017141452.1| PREDICTED: biotin--protein ligase isoform X5 [Drosophila mir
>gi|1037054305|ref|XP_017141451.1| PREDICTED: uncharacterized protein LOC108155261 isoform X4
>gi|1037054325|ref|XP_017141453.1| PREDICTED: kinase suppressor of Ras 1 [Drosophila miranda]
>gi|1037083755|ref|XP_017143489.1| PREDICTED: SURP and G-patch domain-containing protein 1 iso
>gi|1037083769|ref|XP_017143490.1| PREDICTED: SURP and G-patch domain-containing protein 1 iso
>gi|1037083783|ref|XP_017143491.1| PREDICTED: formin-like protein 20 isoform X3 [Drosophila mi
>gi|1037083727|ref|XP_017143486.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1037083741|ref|XP_017143488.1| PREDICTED: bumetanide-sensitive sodium-(potassium)-chloride
>gi|1037059827|ref|XP_017141830.1| PREDICTED: glutamate [NMDA] receptor subunit 1 [Drosophila m
>gi|1037059860|ref|XP_017141832.1| PREDICTED: uncharacterized protein LOC108155504 isoform X2
>gi|1037059842|ref|XP_017141831.1| PREDICTED: uncharacterized protein LOC108155504 isoform X1
>gi|1037059751|ref|XP_017141824.1| PREDICTED: inositol 1,4,5-trisphosphate receptor isoform X1
>gi|1037059767|ref|XP_017141825.1| PREDICTED: inositol 1,4,5-trisphosphate receptor isoform X1
>gi|1037059779|ref|XP_017141826.1| PREDICTED: inositol 1,4,5-trisphosphate receptor isoform X2
>gi|1037059811|ref|XP_017141829.1| PREDICTED: uncharacterized protein LOC108155502 isoform X2
>gi|1037059795|ref|XP_017141827.1| PREDICTED: uncharacterized protein LOC108155502 isoform X1
>gi|1037071659|ref|XP_017142629.1| PREDICTED: protein-L-isoaspartate(D-aspartate) O-methyltrans
>gi|1037071581|ref|XP_017142623.1| PREDICTED: DNA cross-link repair 1A protein [Drosophila mira
>gi|1037071594|ref|XP_017142624.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037071607|ref|XP_017142625.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037071632|ref|XP_017142627.1| PREDICTED: phosphatidylinositol 4-kinase type 2-alpha isofor
>gi|1037071646|ref|XP_017142628.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037071620|ref|XP_017142626.1| PREDICTED: phosphatidylinositol 4-kinase type 2-beta isoform
>gi|1037089775|ref|XP_017143918.1| PREDICTED: uncharacterized protein LOC108156765 [Drosophila
>gi|1037089731|ref|XP_017143915.1| PREDICTED: phospholipid phosphatase 5 isoform X1 [Drosophila
>gi|1037089747|ref|XP_017143916.1| PREDICTED: phospholipid phosphatase 5 isoform X2 [Drosophila
>gi|1037089761|ref|XP_017143917.1| PREDICTED: RNA-binding protein 42 [Drosophila miranda]
>gi|1037101065|ref|XP_017144846.1| PREDICTED: GTP-binding protein Rheb homolog [Drosophila mira
>gi|1037098227|ref|XP_017144594.1| PREDICTED: succinate-semialdehyde dehydrogenase [NADP(+)] Ga
>gi|1037049406|ref|XP_017141116.1| PREDICTED: nuclear protein localization protein 4 homolog is
>gi|1037049422|ref|XP_017141117.1| PREDICTED: nuclear protein localization protein 4 homolog is
>gi|1037049388|ref|XP_017141115.1| PREDICTED: nuclear protein localization protein 4 homolog is
>gi|1037049440|ref|XP_017141118.1| PREDICTED: uncharacterized protein LOC108155052 isoform X1
>gi|1037049472|ref|XP_017141121.1| PREDICTED: uncharacterized protein LOC108155052 isoform X3
>gi|1037049456|ref|XP_017141119.1| PREDICTED: uncharacterized protein LOC108155052 isoform X2
>gi|1037049632|ref|XP_017141131.1| PREDICTED: putative uncharacterized protein DDB_G0274435 [D
>gi|1037049614|ref|XP_017141130.1| PREDICTED: uncharacterized protein LOC108155056 isoform X2
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>gi|1037049582|ref|XP_017141128.1| PREDICTED: uncharacterized protein LOC108155056 isoform X1
>gi|1037049598|ref|XP_017141129.1| PREDICTED: uncharacterized protein LOC108155056 isoform X1
>gi|1037049534|ref|XP_017141125.1| PREDICTED: protein gar2 isoform X3 [Drosophila miranda]
>gi|1037049520|ref|XP_017141124.1| PREDICTED: CUGBP Elav-like family member 2 isoform X2 [Dros
>gi|1037049488|ref|XP_017141122.1| PREDICTED: CUGBP Elav-like family member 2 isoform X1 [Dros
>gi|1037049504|ref|XP_017141123.1| PREDICTED: CUGBP Elav-like family member 2 isoform X1 [Dros
>gi|1037049550|ref|XP_017141126.1| PREDICTED: lipoprotein lipase [Drosophila miranda]
>gi|1037049566|ref|XP_017141127.1| PREDICTED: uncharacterized protein LOC108155055 [Drosophila
>gi|1037104163|ref|XP_017145175.1| PREDICTED: protein regulator of cytokinesis 1-like isoform X
>gi|1037104171|ref|XP_017145176.1| PREDICTED: protein regulator of cytokinesis 1-like isoform X
>gi|1037104176|ref|XP_017145177.1| PREDICTED: NEDD8 ultimate buster 1 [Drosophila miranda]
>gi|1037036570|ref|XP_017140276.1| PREDICTED: uncharacterized protein LOC108154486 [Drosophila
>gi|1037100542|ref|XP_017144792.1| PREDICTED: fatty-acid amide hydrolase 2 [Drosophila miranda]
>gi|1037079511|ref|XP_017143164.1| PREDICTED: dihydroxyacetone phosphate acyltransferase [Dros
>gi|1037079524|ref|XP_017143165.1| PREDICTED: putative GTP-binding protein 6 [Drosophila miran
>gi|1037079538|ref|XP_017143166.1| PREDICTED: transmembrane protein 141 [Drosophila miranda]
>gi|1037075515|ref|XP_017142896.1| PREDICTED: zinc finger protein 436 [Drosophila miranda]
>gi|1037075498|ref|XP_017142895.1| PREDICTED: MAU2 chromatid cohesion factor homolog [Drosophil
>gi|1037075531|ref|XP_017142897.1| PREDICTED: serpin B3 [Drosophila miranda]
>gi|1037075545|ref|XP_017142898.1| PREDICTED: serpin B3 [Drosophila miranda]
>gi|1037079498|ref|XP_017143163.1| PREDICTED: transmembrane protein 165 isoform X2 [Drosophila
>gi|1037079470|ref|XP_017143161.1| PREDICTED: transmembrane protein 165 isoform X1 [Drosophila
>gi|1037079482|ref|XP_017143162.1| PREDICTED: transmembrane protein 165 isoform X1 [Drosophila
>gi|1037079456|ref|XP_017143160.1| PREDICTED: E3 ubiquitin-protein ligase RNF123 [Drosophila m
>gi|1037032776|ref|XP_017140058.1| PREDICTED: soluble guanylate cyclase 88E [Drosophila miranda
>gi|1037099815|ref|XP_017144735.1| PREDICTED: uncharacterized protein LOC108157264 isoform X2
>gi|1037099839|ref|XP_017144737.1| PREDICTED: uncharacterized protein LOC108157264 isoform X4
>gi|1037099803|ref|XP_017144734.1| PREDICTED: uncharacterized protein LOC108157264 isoform X1
>gi|1037099827|ref|XP_017144736.1| PREDICTED: uncharacterized protein LOC108157264 isoform X3
>gi|1037099851|ref|XP_017144738.1| PREDICTED: uncharacterized protein LOC108157264 isoform X5
>gi|1037099861|ref|XP_017144740.1| PREDICTED: uncharacterized protein LOC108157264 isoform X6
>gi|1037099872|ref|XP_017144741.1| PREDICTED: uncharacterized protein LOC108157264 isoform X7
>gi|1037099882|ref|XP_017144742.1| PREDICTED: uncharacterized protein LOC108157264 isoform X8
>gi|1037088541|ref|XP_017143821.1| PREDICTED: glycogen [starch] synthase isoform X1 [Drosophila
>gi|1037088555|ref|XP_017143822.1| PREDICTED: glycogen [starch] synthase isoform X2 [Drosophila
>gi|1037088569|ref|XP_017143823.1| PREDICTED: uncharacterized protein LOC108156690 [Drosophila
>gi|1037105433|ref|XP_017145357.1| PREDICTED: vegetative cell wall protein gp1 [Drosophila mira
>gi|1037105438|ref|XP_017145359.1| PREDICTED: uncharacterized protein LOC108157700 [Drosophila
>gi|1037092051|ref|XP_017144094.1| PREDICTED: uncharacterized protein LOC108156877 [Drosophila
>gi|1037092064|ref|XP_017144095.1| PREDICTED: uncharacterized protein LOC108156877 [Drosophila
>gi|1037092078|ref|XP_017144096.1| PREDICTED: uncharacterized protein LOC108156877 [Drosophila
>gi|1037089359|ref|XP_017143885.1| PREDICTED: protein fem-1 homolog CG6966 [Drosophila miranda]
>gi|1037023281|ref|XP_017139476.1| PREDICTED: low choriolytic enzyme [Drosophila miranda]
>gi|1037104354|ref|XP_017145202.1| PREDICTED: uncharacterized protein LOC108157589 isoform X1
>gi|1037104362|ref|XP_017145203.1| PREDICTED: uncharacterized protein LOC108157589 isoform X1
>gi|1037104372|ref|XP_017145205.1| PREDICTED: uncharacterized protein LOC108157589 isoform X2
>gi|1037037786|ref|XP_017140347.1| PREDICTED: uncharacterized protein LOC108154555 [Drosophila
>gi|1037106727|ref|XP_017145536.1| PREDICTED: uncharacterized protein LOC108157843 isoform X2
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>gi|1037106719|ref|XP_017145535.1| PREDICTED: uncharacterized protein LOC108157843 isoform X1
>gi|1037104554|ref|XP_017145230.1| PREDICTED: uncharacterized protein LOC108157612 [Drosophila
>gi|1037104547|ref|XP_017145229.1| PREDICTED: general transcription factor IIF subunit 1-like
>gi|1037092373|ref|XP_017144114.1| PREDICTED: insulin-like peptide receptor [Drosophila miranda
>gi|1037037208|ref|XP_017140312.1| PREDICTED: luciferin 4-monooxygenase-like [Drosophila miranc
>gi|1037023299|ref|XP_017139477.1| PREDICTED: succinate-semialdehyde dehydrogenase [NADP(+)] Ga
>gi|1037072161|ref|XP_017142662.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit
>gi|1037072177|ref|XP_017142663.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit
>gi|1037072116|ref|XP_017142658.1| PREDICTED: 40S ribosomal protein S5b [Drosophila miranda]
>gi|1037072131|ref|XP_017142660.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit :
>gi|1037072147|ref|XP_017142661.1| PREDICTED: V-type proton ATPase 21 kDa proteolipid subunit :
>gi|1037072068|ref|XP_017142655.1| PREDICTED: RRP15-like protein [Drosophila miranda]
>gi|1037072100|ref|XP_017142657.1| PREDICTED: UPF0568 protein C14orf166 homolog [Drosophila mi
>gi|1037072191|ref|XP_017142664.1| PREDICTED: uncharacterized protein LOC108155984 [Drosophila
>gi|1037072052|ref|XP_017142654.1| PREDICTED: nudC domain-containing protein 3 [Drosophila mira
>gi|1037072084|ref|XP_017142656.1| PREDICTED: nurim homolog [Drosophila miranda]
>gi|1037090445|ref|XP_017143974.1| PREDICTED: eukaryotic translation initiation factor 1A, X-cl
>gi|1037090459|ref|XP_017143975.1| PREDICTED: eukaryotic translation initiation factor 1A, X-cl
>gi|1037090473|ref|XP_017143977.1| PREDICTED: eukaryotic translation initiation factor 1A, X-cl
>gi|1037090486|ref|XP_017143978.1| PREDICTED: eukaryotic translation initiation factor 1A, X-cl
>gi|1037090431|ref|XP_017143973.1| PREDICTED: transmembrane protease serine 9 [Drosophila miran
>gi|1037090500|ref|XP_017143979.1| PREDICTED: uncharacterized protein LOC108156803 [Drosophila
>gi|1037087703|ref|XP_017143756.1| PREDICTED: WD repeat-containing protein 3 [Drosophila mirano
>gi|1037087717|ref|XP_017143757.1| PREDICTED: peptidyl-prolyl cis-trans isomerase FKBP2 [Drosop
>gi|1037068919|ref|XP_017142440.1| PREDICTED: flocculation protein FLO11 [Drosophila miranda]
>gi|1037068935|ref|XP_017142441.1| PREDICTED: putative elongator complex protein 1 [Drosophila
>gi|1037064004|ref|XP_017142117.1| PREDICTED: uncharacterized protein LOC108155665 [Drosophila
>gi|1037063989|ref|XP_017142116.1| PREDICTED: nuclear pore complex protein Nup214 isoform X2 [I
>gi|1037063961|ref|XP_017142113.1| PREDICTED: nuclear pore complex protein Nup214 isoform X1 [I
>gi|1037063974|ref|XP_017142114.1| PREDICTED: nuclear pore complex protein Nup214 isoform X1 [I
>gi|1037040551|ref|XP_017140528.1| PREDICTED: zinc finger protein 10 [Drosophila miranda]
>gi|1037040624|ref|XP_017140533.1| PREDICTED: gastrula zinc finger protein XlCGF17.1 isoform X:
>gi|1037040638|ref|XP_017140534.1| PREDICTED: zinc finger protein 211 isoform X2 [Drosophila m:
>gi|1037040539|ref|XP_017140526.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037040597|ref|XP_017140531.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037040565|ref|XP_017140529.1| PREDICTED: zinc finger protein 20 [Drosophila miranda]
>gi|1037040382|ref|XP_017140514.1| PREDICTED: zinc finger protein 229 [Drosophila miranda]
>gi|1037040654|ref|XP_017140535.1| PREDICTED: zinc finger protein 648-like [Drosophila miranda]
>gi|1037040446|ref|XP_017140520.1| PREDICTED: transcription factor Ouib-like isoform X5 [Drosop
>gi|1037040430|ref|XP_017140519.1| PREDICTED: transcription factor Ouib-like isoform X4 [Drosop
>gi|1037040470|ref|XP_017140521.1| PREDICTED: transcription factor Ouib-like isoform X6 [Drosop
>gi|1037040485|ref|XP_017140522.1| PREDICTED: transcription factor Ouib-like isoform X6 [Drosop
>gi|1037040497|ref|XP_017140523.1| PREDICTED: transcription factor Ouib-like isoform X7 [Drosop
>gi|1037040400|ref|XP_017140515.1| PREDICTED: transcription factor Ouib-like isoform X1 [Drosop
>gi|1037040420|ref|XP_017140518.1| PREDICTED: transcription factor Ouib-like isoform X3 [Drosop
>gi|1037040410|ref|XP_017140516.1| PREDICTED: transcription factor Ouib-like isoform X2 [Drosop
>gi|1037040581|ref|XP_017140530.1| PREDICTED: transcription factor Ouib-like [Drosophila mirano
>gi|1037040666|ref|XP_017140536.1| PREDICTED: protein RER1 [Drosophila miranda]
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>gi|1037040608|ref|XP_017140532.1| PREDICTED: 3-ketodihydrosphingosine reductase [Drosophila m:
>gi|1037040680|ref|XP_017140537.1| PREDICTED: peptidyl-prolyl cis-trans isomerase NIMA-interact
>gi|1037040346|ref|XP_017140512.1| PREDICTED: probable glutamine--tRNA ligase [Drosophila miran
>gi|1037040362|ref|XP_017140513.1| PREDICTED: serine/threonine-protein kinase RIO2 [Drosophila
>gi|1037040527|ref|XP_017140525.1| PREDICTED: uncharacterized protein LOC108154688 [Drosophila
>gi|1037040511|ref|XP_017140524.1| PREDICTED: uncharacterized protein LOC108154687 [Drosophila
>gi|1037104456|ref|XP_017145216.1| PREDICTED: uncharacterized protein LOC108157599 [Drosophila
>gi|1037102846|ref|XP_017145017.1| PREDICTED: uncharacterized protein LOC108157456 [Drosophila
>gi|1037102855|ref|XP_017145018.1| PREDICTED: uncharacterized protein LOC108157457 [Drosophila
>gi|1037092862|ref|XP_017144147.1| PREDICTED: uncharacterized protein LOC108156909 [Drosophila
>gi|1037092892|ref|XP_017144149.1| PREDICTED: uncharacterized protein LOC108156912 [Drosophila
>gi|1037092878|ref|XP_017144148.1| PREDICTED: uncharacterized protein LOC108156910 [Drosophila
>gi|1037106153|ref|XP_017145457.1| PREDICTED: uncharacterized protein LOC108157778 [Drosophila
>gi|1037106205|ref|XP_017145465.1| PREDICTED: uncharacterized protein LOC108157786 [Drosophila
>gi|1037105460|ref|XP_017145362.1| PREDICTED: uncharacterized protein LOC108157702 [Drosophila
>gi|1037032813|ref|XP_017140061.1| PREDICTED: uncharacterized protein LOC108154314 isoform X1
>gi|1037032831|ref|XP_017140062.1| PREDICTED: uncharacterized protein LOC108154314 isoform X2
>gi|1037104324|ref|XP_017145198.1| PREDICTED: uncharacterized protein LOC108157585 [Drosophila
>gi|1037104332|ref|XP_017145199.1| PREDICTED: uncharacterized protein LOC108157585 [Drosophila
>gi|1037033108|ref|XP_017140079.1| PREDICTED: uncharacterized protein LOC108154328 [Drosophila
>gi|1037106774|ref|XP_017145543.1| PREDICTED: uncharacterized protein LOC108157850 [Drosophila
>gi|1037106087|ref|XP_017145448.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037106095|ref|XP_017145449.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037106103|ref|XP_017145451.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037106111|ref|XP_017145452.1| PREDICTED: uncharacterized protein LOC108157774 [Drosophila
>gi|1037023315|ref|XP_017139478.1| PREDICTED: uncharacterized protein LOC108153890 [Drosophila
>gi|1037032314|ref|XP_017140029.1| PREDICTED: uncharacterized protein LOC108154289 [Drosophila
>gi|1037032484|ref|XP_017140040.1| PREDICTED: uncharacterized protein LOC108154298 [Drosophila
>gi|1037093392|ref|XP_017144185.1| PREDICTED: uncharacterized protein LOC108156935 [Drosophila
>gi|1037093406|ref|XP_017144186.1| PREDICTED: uncharacterized protein LOC108156936 [Drosophila
>gi|1037093379|ref|XP_017144183.1| PREDICTED: zinc finger protein 62 homolog [Drosophila mirano
>gi|1037086961|ref|XP_017143730.1| PREDICTED: zinc finger protein 91 [Drosophila miranda]
>gi|1037079058|ref|XP_017143144.1| PREDICTED: TATA-binding protein-associated factor 172 [Drosc
>gi|1037058017|ref|XP_017141701.1| PREDICTED: SWI/SNF complex subunit SMARCC1 [Drosophila miran
>gi|1037058031|ref|XP_017141702.1| PREDICTED: retinoblastoma family protein [Drosophila miranda
>gi|1037058106|ref|XP_017141708.1| PREDICTED: uncharacterized protein LOC108155419 [Drosophila
>gi|1037058077|ref|XP_017141707.1| PREDICTED: zinc finger CCHC domain-containing protein 10 [Dr
>gi|1037058047|ref|XP_017141703.1| PREDICTED: uncharacterized protein LOC108155417 [Drosophila
>gi|1037058063|ref|XP_017141704.1| PREDICTED: uncharacterized protein LOC108155417 [Drosophila
>gi|1037061293|ref|XP_017141932.1| PREDICTED: synapse-associated protein of 47 kDa isoform X9
>gi|1037061279|ref|XP_017141931.1| PREDICTED: synapse-associated protein of 47 kDa isoform X8
>gi|1037061264|ref|XP_017141930.1| PREDICTED: synapse-associated protein of 47 kDa isoform X7
>gi|1037061186|ref|XP_017141924.1| PREDICTED: synapse-associated protein of 47 kDa isoform X2
>gi|1037061202|ref|XP_017141925.1| PREDICTED: synapse-associated protein of 47 kDa isoform X3
>gi|1037061216|ref|XP_017141926.1| PREDICTED: synapse-associated protein of 47 kDa isoform X4
>gi|1037061155|ref|XP_017141922.1| PREDICTED: synapse-associated protein of 47 kDa isoform X1
>gi|1037061170|ref|XP_017141923.1| PREDICTED: synapse-associated protein of 47 kDa isoform X1
>gi|1037061309|ref|XP_017141933.1| PREDICTED: synapse-associated protein of 47 kDa isoform X10
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>gi|1037061248|ref|XP_017141929.1| PREDICTED: synapse-associated protein of 47 kDa isoform X6
>gi|1037061232|ref|XP_017141928.1| PREDICTED: synapse-associated protein of 47 kDa isoform X5
>gi|1037061325|ref|XP_017141934.1| PREDICTED: synapse-associated protein of 47 kDa isoform X11
>gi|1037061371|ref|XP_017141935.1| PREDICTED: synapse-associated protein of 47 kDa isoform X12
>gi|1037061414|ref|XP_017141937.1| PREDICTED: mitochondrial import inner membrane translocase s
>gi|1037061398|ref|XP_017141936.1| PREDICTED: uncharacterized protein LOC108155556 [Drosophila
>gi|1037061137|ref|XP_017141921.1| PREDICTED: F-box/LRR-repeat protein 7 [Drosophila miranda]
>gi|1037061430|ref|XP_017141938.1| PREDICTED: PI-actitoxin-Afv2a [Drosophila miranda]
>gi|1037095049|ref|XP_017144307.1| PREDICTED: uncharacterized protein LOC108157013 isoform X3
>gi|1037095033|ref|XP_017144306.1| PREDICTED: serologically defined colon cancer antigen 8 homo
>gi|1037095019|ref|XP_017144305.1| PREDICTED: serologically defined colon cancer antigen 8 homo
>gi|1037095063|ref|XP_017144308.1| PREDICTED: poly(U)-specific endoribonuclease homolog [Drosop
>gi|1037097697|ref|XP_017144540.1| PREDICTED: globin CTT-VI [Drosophila miranda]
>gi|1037097711|ref|XP_017144541.1| PREDICTED: globin CTT-VI [Drosophila miranda]
>gi|1037097836|ref|XP_017144554.1| PREDICTED: atrial natriuretic peptide receptor 3 [Drosophila
>gi|1037097823|ref|XP_017144553.1| PREDICTED: atrial natriuretic peptide receptor 1 [Drosophila
>gi|1037097846|ref|XP_017144555.1| PREDICTED: uncharacterized protein LOC108157157 [Drosophila
>gi|1037052769|ref|XP_017141348.1| PREDICTED: ATP-binding cassette sub-family B member 6, mito
>gi|1037052797|ref|XP_017141349.1| PREDICTED: ATP-binding cassette sub-family B member 6, mito
>gi|1037052737|ref|XP_017141346.1| PREDICTED: uncharacterized protein LOC108155186 [Drosophila
>gi|1037052813|ref|XP_017141350.1| PREDICTED: arginine kinase [Drosophila miranda]
>gi|1037052753|ref|XP_017141347.1| PREDICTED: probable ATP-dependent RNA helicase spindle-E [Dr
>gi|1037052827|ref|XP_017141351.1| PREDICTED: uncharacterized protein LOC108155191 [Drosophila
>gi|1037069046|ref|XP_017142449.1| PREDICTED: NADH-quinone oxidoreductase subunit I [Drosophila
>gi|1037068998|ref|XP_017142446.1| PREDICTED: structure-specific endonuclease subunit SLX1 homo
>gi|1037069012|ref|XP_017142447.1| PREDICTED: mediator of RNA polymerase II transcription subur
>gi|1037069028|ref|XP_017142448.1| PREDICTED: actin-related protein 2/3 complex subunit 3 [Dros
>gi|1037068966|ref|XP_017142443.1| PREDICTED: uncharacterized protein LOC108155844 isoform X1
>gi|1037068982|ref|XP_017142445.1| PREDICTED: uncharacterized protein LOC108155844 isoform X2
>gi|1037092846|ref|XP_017144146.1| PREDICTED: micronuclear linker histone polyprotein [Drosophi
>gi|1037103905|ref|XP_017145143.1| PREDICTED: 5-hydroxytryptamine receptor 2B [Drosophila mirar
>gi|1037103913|ref|XP_017145144.1| PREDICTED: uncharacterized protein LOC108157551 [Drosophila
>gi|1037104673|ref|XP_017145246.1| PREDICTED: uncharacterized protein LOC108157621 [Drosophila
>gi|1037104683|ref|XP_017145247.1| PREDICTED: trichohyalin [Drosophila miranda]
>gi|1037094620|ref|XP_017144275.1| PREDICTED: pre-mRNA 3'-end-processing factor FIP1 [Drosophil
>gi|1037032619|ref|XP_017140048.1| PREDICTED: nuclear receptor subfamily 2 group E member 1 iso
>gi|1037032656|ref|XP_017140050.1| PREDICTED: nuclear receptor subfamily 2 group E member 1 iso
>gi|1037032636|ref|XP_017140049.1| PREDICTED: nuclear receptor subfamily 2 group E member 1 iso
>gi|1037088719|ref|XP_017143835.1| PREDICTED: serine/threonine-protein kinase PAK 1 [Drosophila
>gi|1037088737|ref|XP_017143836.1| PREDICTED: serine/threonine-protein kinase PAK 1 [Drosophila
>gi|1037056216|ref|XP_017141578.1| PREDICTED: transcription elongation regulator 1 isoform X2
>gi|1037056203|ref|XP_017141576.1| PREDICTED: transcription elongation regulator 1 isoform X1
>gi|1037056264|ref|XP_017141581.1| PREDICTED: transcription elongation regulator 1 isoform X5
>gi|1037056232|ref|XP_017141579.1| PREDICTED: transcription elongation regulator 1 isoform X3
>gi|1037056248|ref|XP_017141580.1| PREDICTED: transcription elongation regulator 1 isoform X4
>gi|1037056282|ref|XP_017141583.1| PREDICTED: HEAT repeat-containing protein 3 [Drosophila mira
>gi|1037023331|ref|XP_017139479.1| PREDICTED: uncharacterized protein LOC108153891 [Drosophila
>gi|1037056314|ref|XP_017141585.1| PREDICTED: uncharacterized protein LOC108155341 [Drosophila
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>gi|1037056298|ref|XP_017141584.1| PREDICTED: general transcription factor IIF subunit 1 [Droso
>gi|1037094390|ref|XP_017144257.1| PREDICTED: E3 ubiquitin-protein ligase RNF13 [Drosophila mi
>gi|1037082465|ref|XP_017143392.1| PREDICTED: transmembrane and coiled-coil domains protein 2
>gi|1037060768|ref|XP_017141896.1| PREDICTED: histone-lysine N-methyltransferase, H3 lysine-79
>gi|1037060786|ref|XP_017141897.1| PREDICTED: histone-lysine N-methyltransferase, H3 lysine-79
>gi|1037089564|ref|XP_017143898.1| PREDICTED: uncharacterized protein LOC108156747 [Drosophila
>gi|1037097097|ref|XP_017144484.1| PREDICTED: zinc finger protein 616 [Drosophila miranda]
>gi|1037103062|ref|XP_017145041.1| PREDICTED: neurochondrin homolog [Drosophila miranda]
>gi|1037102740|ref|XP_017145006.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102780|ref|XP_017145011.1| PREDICTED: uncharacterized protein LOC108157454 isoform X5
>gi|1037102758|ref|XP_017145009.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102802|ref|XP_017145013.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102748|ref|XP_017145008.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102792|ref|XP_017145012.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102768|ref|XP_017145010.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102816|ref|XP_017145014.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037102827|ref|XP_017145015.1| PREDICTED: serine/arginine repetitive matrix protein 1 isof
>gi|1037036995|ref|XP_017140300.1| PREDICTED: uncharacterized protein LOC108154510 [Drosophila
>gi|1037035889|ref|XP_017140236.1| PREDICTED: uncharacterized protein LOC108154473 isoform X2
>gi|1037035871|ref|XP_017140235.1| PREDICTED: uncharacterized protein LOC108154473 isoform X1
>gi|1037034002|ref|XP_017140131.1| PREDICTED: uncharacterized protein LOC108154378 [Drosophila
>gi|1037100520|ref|XP_017144790.1| PREDICTED: hemicentin-2 [Drosophila miranda]
>gi|1037100500|ref|XP_017144788.1| PREDICTED: probable histone-binding protein Caf1 [Drosophila
>gi|1037100510|ref|XP_017144789.1| PREDICTED: golgin subfamily A member 6-like protein 22 [Dros
>gi|1037100530|ref|XP_017144791.1| PREDICTED: uncharacterized protein LOC108157301 [Drosophila
>gi|1037033766|ref|XP_017140116.1| PREDICTED: uncharacterized protein LOC108154362 [Drosophila
>gi|1037106808|ref|XP_017145549.1| PREDICTED: cytoplasmic dynein 1 light intermediate chain 1-1
>gi|1037035613|ref|XP_017140219.1| PREDICTED: Golgi-associated plant pathogenesis-related prote
>gi|1037095800|ref|XP_017144367.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Drose
>gi|1037095823|ref|XP_017144369.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Drose
>gi|1037095835|ref|XP_017144370.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Drose
>gi|1037095846|ref|XP_017144371.1| PREDICTED: flocculation protein FLO11-like isoform X1 [Drose
>gi|1037095858|ref|XP_017144372.1| PREDICTED: uncharacterized protein LOC108157054 isoform X2
>gi|1037095870|ref|XP_017144373.1| PREDICTED: uncharacterized protein LOC108157055 [Drosophila
>gi|1037095913|ref|XP_017144374.1| PREDICTED: uncharacterized protein LOC108157056 [Drosophila
>gi|1037092386|ref|XP_017144115.1| PREDICTED: argininosuccinate synthase isoform X1 [Drosophila
>gi|1037092400|ref|XP_017144116.1| PREDICTED: argininosuccinate synthase isoform X2 [Drosophila
>gi|1037092414|ref|XP_017144117.1| PREDICTED: uncharacterized protein LOC108156891 isoform X3
>gi|1037092430|ref|XP_017144119.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037092444|ref|XP_017144120.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037092458|ref|XP_017144121.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037092472|ref|XP_017144122.1| PREDICTED: speckle targeted PIP5K1A-regulated poly(A) polyme
>gi|1037062362|ref|XP_017142003.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062378|ref|XP_017142004.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062392|ref|XP_017142005.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062407|ref|XP_017142006.1| PREDICTED: transcription initiation factor TFIID subunit 1
>gi|1037062455|ref|XP_017142009.1| PREDICTED: probable peptidyl-tRNA hydrolase 2 [Drosophila mi
>gi|1037062439|ref|XP_017142008.1| PREDICTED: signal peptidase complex catalytic subunit SEC11A
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>gi|1037062423|ref|XP_017142007.1| PREDICTED: regulatory protein ada [Drosophila miranda]
>gi|1037102136|ref|XP_017144945.1| PREDICTED: homeotic protein labial [Drosophila miranda]
>gi|1037023347|ref|XP_017139480.1| PREDICTED: pupal cuticle protein Edg-84A [Drosophila miranda
>gi|1037036635|ref|XP_017140280.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037023363|ref|XP_017139481.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037023379|ref|XP_017139483.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037036695|ref|XP_017140285.1| PREDICTED: larval cuticle protein A3A [Drosophila miranda]
>gi|1037037731|ref|XP_017140344.1| PREDICTED: uncharacterized protein LOC108154552 [Drosophila
>gi|1037038864|ref|XP_017140413.1| PREDICTED: cuticle protein 19.8 [Drosophila miranda]
>gi|1037037065|ref|XP_017140305.1| PREDICTED: larval cuticle protein A2B [Drosophila miranda]
>gi|1037034464|ref|XP_017140161.1| PREDICTED: cuticle protein 7-like [Drosophila miranda]
>gi|1037101693|ref|XP_017144899.1| PREDICTED: homeotic protein proboscipedia [Drosophila mirand
>gi|1037101703|ref|XP_017144900.1| PREDICTED: homeotic protein proboscipedia [Drosophila mirand
>gi|1037030845|ref|XP_017139943.1| PREDICTED: protein zerknuellt 2 [Drosophila miranda]
>gi|1037030871|ref|XP_017139945.1| PREDICTED: tetratricopeptide repeat protein 12 [Drosophila m
>gi|1037030973|ref|XP_017139950.1| PREDICTED: protein zerknuellt [Drosophila miranda]
>gi|1037058367|ref|XP_017141727.1| PREDICTED: homeotic protein bicoid isoform X1 [Drosophila mi
>gi|1037058383|ref|XP_017141728.1| PREDICTED: homeotic protein bicoid isoform X2 [Drosophila mi
>gi|1037058431|ref|XP_017141731.1| PREDICTED: protein amalgam [Drosophila miranda]
>gi|1037058447|ref|XP_017141733.1| PREDICTED: uncharacterized protein LOC108155432 [Drosophila
>gi|1037058354|ref|XP_017141726.1| PREDICTED: homeotic protein deformed [Drosophila miranda]
>gi|1037058399|ref|XP_017141729.1| PREDICTED: homeotic protein Sex combs reduced [Drosophila mi
>gi|1037058415|ref|XP_017141730.1| PREDICTED: homeotic protein Sex combs reduced [Drosophila mi
>gi|1037101222|ref|XP_017144860.1| PREDICTED: segmentation protein fushi tarazu [Drosophila mir
>gi|1037032569|ref|XP_017140045.1| PREDICTED: uncharacterized protein LOC108154302 [Drosophila
>gi|1037066419|ref|XP_017142270.1| PREDICTED: homeotic protein antennapedia isoform X4 [Drosoph
>gi|1037066406|ref|XP_017142269.1| PREDICTED: homeotic protein antennapedia isoform X3 [Drosoph
>gi|1037066392|ref|XP_017142268.1| PREDICTED: homeotic protein antennapedia isoform X2 [Drosoph
>gi|1037066362|ref|XP_017142266.1| PREDICTED: homeotic protein antennapedia isoform X1 [Drosoph
>gi|1037066379|ref|XP_017142267.1| PREDICTED: homeotic protein antennapedia isoform X1 [Drosoph
>gi|1037049041|ref|XP_017141091.1| PREDICTED: sorbitol dehydrogenase [Drosophila miranda]
>gi|1037023395|ref|XP_017139484.1| PREDICTED: putative leucine-rich repeat-containing protein l
>gi|1037049012|ref|XP_017141089.1| PREDICTED: axoneme-associated protein mst101(2) [Drosophila
>gi|1037049028|ref|XP_017141090.1| PREDICTED: arrestin domain-containing protein 2 [Drosophila
>gi|1037048891|ref|XP_017141080.1| PREDICTED: PAX3- and PAX7-binding protein 1 isoform X1 [Dros
>gi|1037048923|ref|XP_017141082.1| PREDICTED: uncharacterized protein LOC108155030 isoform X3
>gi|1037048907|ref|XP_017141081.1| PREDICTED: PAX3- and PAX7-binding protein 1 isoform X2 [Dros
>gi|1037048941|ref|XP_017141083.1| PREDICTED: E3 UFM1-protein ligase 1 homolog [Drosophila mira
>gi|1037048980|ref|XP_017141087.1| PREDICTED: TBC1 domain family member 22B [Drosophila miranda
>gi|1037048956|ref|XP_017141084.1| PREDICTED: exocyst complex component 6 [Drosophila miranda]
>gi|1037048996|ref|XP_017141088.1| PREDICTED: major facilitator superfamily domain-containing p
>gi|1037064384|ref|XP_017142142.1| PREDICTED: ras-related protein Rab-11A [Drosophila miranda]
>gi|1037064350|ref|XP_017142140.1| PREDICTED: protein Peter pan [Drosophila miranda]
>gi|1037064366|ref|XP_017142141.1| PREDICTED: peptidyl-prolyl cis-trans isomerase D [Drosophila
>gi|1037064334|ref|XP_017142138.1| PREDICTED: F-box/WD repeat-containing protein 11 [Drosophila
>gi|1037085749|ref|XP_017143636.1| PREDICTED: acylpyruvase FAHD1, mitochondrial [Drosophila mir
>gi|1037085763|ref|XP_017143637.1| PREDICTED: uncharacterized protein LOC108156606 isoform X1
>gi|1037085777|ref|XP_017143638.1| PREDICTED: uncharacterized protein LOC108156606 isoform X2
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>gi|1037085721|ref|XP_017143634.1| PREDICTED: 14-3-3 protein epsilon isoform X1 [Drosophila mi
>gi|1037085735|ref|XP_017143635.1| PREDICTED: 14-3-3 protein epsilon isoform X2 [Drosophila mi
>gi|1037061552|ref|XP_017141947.1| PREDICTED: uncharacterized protein LOC108155565 [Drosophila
>gi|1037061522|ref|XP_017141945.1| PREDICTED: protein AAR2 homolog [Drosophila miranda]
>gi|1037061478|ref|XP_017141942.1| PREDICTED: DNA-directed RNA polymerase II 16 kDa polypeptide
>gi|1037061462|ref|XP_017141941.1| PREDICTED: transcriptional adapter 2A isoform X1 [Drosophila
>gi|1037061542|ref|XP_017141946.1| PREDICTED: uncharacterized protein LOC108155564 [Drosophila
>gi|1037061492|ref|XP_017141943.1| PREDICTED: trimethylguanosine synthase [Drosophila miranda]
>gi|1037061506|ref|XP_017141944.1| PREDICTED: homeobox protein MOX-2 [Drosophila miranda]
>gi|1037061446|ref|XP_017141940.1| PREDICTED: vam6/Vps39-like protein [Drosophila miranda]
>gi|1037090890|ref|XP_017144008.1| PREDICTED: signal peptide peptidase-like 3 isoform X1 [Dros
>gi|1037090903|ref|XP_017144009.1| PREDICTED: signal peptide peptidase-like 3 isoform X2 [Dros
>gi|1037091459|ref|XP_017144051.1| PREDICTED: SH2B adapter protein 1 isoform X1 [Drosophila mi
>gi|1037091473|ref|XP_017144052.1| PREDICTED: SH2B adapter protein 1 isoform X1 [Drosophila mi
>gi|1037091487|ref|XP_017144053.1| PREDICTED: SH2B adapter protein 1 isoform X2 [Drosophila mi
>gi|1037091501|ref|XP_017144054.1| PREDICTED: SH2B adapter protein 1 isoform X3 [Drosophila mi
>gi|1037100115|ref|XP_017144753.1| PREDICTED: protein FAM98B [Drosophila miranda]
>gi|1037023411|ref|XP_017139485.1| PREDICTED: superoxide dismutase [Cu-Zn], chloroplastic [Dros
>gi|1037037083|ref|XP_017140306.1| PREDICTED: laccase [Drosophila miranda]
>gi|1037106636|ref|XP_017145524.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1037032936|ref|XP_017140068.1| PREDICTED: lipase 3 [Drosophila miranda]
>gi|1037033784|ref|XP_017140117.1| PREDICTED: lipase 3-like [Drosophila miranda]
>gi|1037100802|ref|XP_017144819.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100772|ref|XP_017144816.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100822|ref|XP_017144821.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100812|ref|XP_017144820.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100792|ref|XP_017144818.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037100782|ref|XP_017144817.1| PREDICTED: sodium-coupled monocarboxylate transporter 1 iso
>gi|1037104707|ref|XP_017145251.1| PREDICTED: uncharacterized protein LOC108157625 [Drosophila
>gi|1037104713|ref|XP_017145252.1| PREDICTED: uncharacterized protein LOC108157626 isoform X1
>gi|1037104721|ref|XP_017145253.1| PREDICTED: uncharacterized protein LOC108157626 isoform X2
>gi|1037032585|ref|XP_017140046.1| PREDICTED: cardioacceleratory peptide receptor [Drosophila m
>gi|1037105484|ref|XP_017145365.1| PREDICTED: lateral signaling target protein 2 homolog [Dros
>gi|1037103023|ref|XP_017145037.1| PREDICTED: LOW QUALITY PROTEIN: protein bride of sevenless
>gi|1037105332|ref|XP_017145342.1| PREDICTED: uncharacterized protein LOC108157686 [Drosophila
>gi|1037063070|ref|XP_017142052.1| PREDICTED: neurofibromin isoform X3 [Drosophila miranda]
>gi|1037063036|ref|XP_017142049.1| PREDICTED: neurofibromin isoform X1 [Drosophila miranda]
>gi|1037063052|ref|XP_017142051.1| PREDICTED: neurofibromin isoform X2 [Drosophila miranda]
>gi|1037104380|ref|XP_017145206.1| PREDICTED: enhancer of split mdelta protein [Drosophila mira
>gi|1037104584|ref|XP_017145234.1| PREDICTED: enhancer of split mgamma protein [Drosophila mira
>gi|1037103882|ref|XP_017145140.1| PREDICTED: enhancer of split mbeta protein [Drosophila mirar
>gi|1037031455|ref|XP_017139981.1| PREDICTED: enhancer of split malpha protein [Drosophila mira
>gi|1037032348|ref|XP_017140030.1| PREDICTED: enhancer of split M1 protein [Drosophila miranda]
>gi|1037031528|ref|XP_017139986.1| PREDICTED: enhancer of split M2 protein [Drosophila miranda]
>gi|1037104850|ref|XP_017145272.1| PREDICTED: enhancer of split m3 protein [Drosophila miranda]
>gi|1037031418|ref|XP_017139979.1| PREDICTED: enhancer of split m4 protein [Drosophila miranda]
>gi|1037030924|ref|XP_017139948.1| PREDICTED: enhancer of split m5 protein [Drosophila miranda]
>gi|1037031475|ref|XP_017139982.1| PREDICTED: enhancer of split m6 protein [Drosophila miranda]
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>gi|1037103953|ref|XP_017145149.1| PREDICTED: enhancer of split m7 protein [Drosophila miranda]
>gi|1037103403|ref|XP_017145078.1| PREDICTED: enhancer of split m8 protein [Drosophila miranda]
>gi|1037103385|ref|XP_017145077.1| PREDICTED: actin-binding protein IPP-like [Drosophila mirand
>gi|1037049648|ref|XP_017141132.1| PREDICTED: protein groucho isoform X1 [Drosophila miranda]
>gi|1037049664|ref|XP_017141133.1| PREDICTED: protein groucho isoform X1 [Drosophila miranda]
>gi|1037049680|ref|XP_017141134.1| PREDICTED: protein groucho isoform X1 [Drosophila miranda]
>gi|1037049696|ref|XP_017141135.1| PREDICTED: protein groucho isoform X2 [Drosophila miranda]
>gi|1037049713|ref|XP_017141136.1| PREDICTED: exocyst complex component 8 [Drosophila miranda]
>gi|1037049747|ref|XP_017141138.1| PREDICTED: charged multivesicular body protein 2a [Drosophil
>gi|1037049779|ref|XP_017141141.1| PREDICTED: UMP-CMP kinase [Drosophila miranda]
>gi|1037049794|ref|XP_017141142.1| PREDICTED: DNA replication complex GINS protein SLD5 [Drosop
>gi|1037049810|ref|XP_017141143.1| PREDICTED: pre-rRNA-processing protein TSR2 homolog [Drosoph
>gi|1037049860|ref|XP_017141146.1| PREDICTED: phosphatidylinositol N-acetylglucosaminyltransfe
>gi|1037049844|ref|XP_017141145.1| PREDICTED: uncharacterized protein LOC108155069 [Drosophila
>gi|1037049828|ref|XP_017141144.1| PREDICTED: 60S ribosomal protein L34 [Drosophila miranda]
>gi|1037049729|ref|XP_017141137.1| PREDICTED: tRNA (adenine(58)-N(1))-methyltransferase cataly
>gi|1037049763|ref|XP_017141139.1| PREDICTED: uncharacterized protein LOC108155064 [Drosophila
>gi|1037104120|ref|XP_017145170.1| PREDICTED: neuronal PAS domain-containing protein 4 [Drosoph
>gi|1037102894|ref|XP_017145022.1| PREDICTED: uncharacterized protein LOC108157460 isoform X2
>gi|1037102885|ref|XP_017145021.1| PREDICTED: uncharacterized protein LOC108157460 isoform X1
>gi|1037102876|ref|XP_017145020.1| PREDICTED: uncharacterized protein LOC108157459 [Drosophila
>gi|1037106644|ref|XP_017145525.1| PREDICTED: glucose dehydrogenase [FAD, quinone] [Drosophila
>gi|1037106540|ref|XP_017145511.1| PREDICTED: GPI ethanolamine phosphate transferase 1-like [Dr
>gi|1037106657|ref|XP_017145527.1| PREDICTED: suppressor protein SRP40-like isoform X2 [Drosoph
>gi|1037106650|ref|XP_017145526.1| PREDICTED: clumping factor A-like isoform X1 [Drosophila mir
>gi|1037085656|ref|XP_017143631.1| PREDICTED: uncharacterized protein LOC108156599 [Drosophila
>gi|1037085600|ref|XP_017143627.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085614|ref|XP_017143628.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085628|ref|XP_017143629.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085642|ref|XP_017143630.1| PREDICTED: uncharacterized protein LOC108156597 [Drosophila
>gi|1037085670|ref|XP_017143632.1| PREDICTED: natterin-3 [Drosophila miranda]
>gi|1037085573|ref|XP_017143626.1| PREDICTED: protein HGH1 homolog [Drosophila miranda]
>gi|1037038365|ref|XP_017140382.1| PREDICTED: LOW QUALITY PROTEIN: proteasome subunit beta type
>gi|1037033508|ref|XP_017140101.1| PREDICTED: neurotrypsin-like [Drosophila miranda]
>gi|1037101313|ref|XP_017144867.1| PREDICTED: uncharacterized protein LOC108157356 [Drosophila
>gi|1037037714|ref|XP_017140343.1| PREDICTED: uncharacterized protein LOC108154551 [Drosophila
>gi|1037023427|ref|XP_017139486.1| PREDICTED: uncharacterized protein LOC108153899 [Drosophila
>gi|1037083391|ref|XP_017143460.1| PREDICTED: proteasome subunit beta type-7-like [Drosophila m
>gi|1037083375|ref|XP_017143459.1| PREDICTED: fructose-bisphosphate aldolase isoform X3 [Drosop
>gi|1037083333|ref|XP_017143456.1| PREDICTED: fructose-bisphosphate aldolase isoform X1 [Drosop
>gi|1037083363|ref|XP_017143458.1| PREDICTED: fructose-bisphosphate aldolase isoform X1 [Drosop
>gi|1037083349|ref|XP_017143457.1| PREDICTED: fructose-bisphosphate aldolase isoform X2 [Drosop
>gi|1037083319|ref|XP_017143455.1| PREDICTED: dipeptidase 1 [Drosophila miranda]
>gi|1037098293|ref|XP_017144601.1| PREDICTED: ABC transporter G family member 20 [Drosophila mi
>gi|1037023441|ref|XP_017139487.1| PREDICTED: uncharacterized protein LOC108153900 [Drosophila
>gi|1037106346|ref|XP_017145483.1| PREDICTED: fructose-bisphosphate aldolase-like [Drosophila m
>gi|1037036816|ref|XP_017140289.1| PREDICTED: uncharacterized protein LOC108154500 [Drosophila
>gi|1037038333|ref|XP_017140380.1| PREDICTED: uncharacterized protein LOC108154587 [Drosophila
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>gi|1037103069|ref|XP_017145042.1| PREDICTED: helix-loop-helix protein delilah [Drosophila mira
>gi|1037033142|ref|XP_017140081.1| PREDICTED: hexokinase type 1 [Drosophila miranda]
>gi|1037106025|ref|XP_017145440.1| PREDICTED: hexokinase type 2 [Drosophila miranda]
>gi|1037043816|ref|XP_017140744.1| PREDICTED: probable uridine-cytidine kinase isoform X1 [Dros
>gi|1037043830|ref|XP_017140745.1| PREDICTED: probable uridine-cytidine kinase isoform X2 [Dros
>gi|1037043784|ref|XP_017140742.1| PREDICTED: zinc metalloproteinase nas-1 isoform X2 [Drosophi
>gi|1037043768|ref|XP_017140741.1| PREDICTED: uncharacterized protein LOC108154836 isoform X1
>gi|1037043800|ref|XP_017140743.1| PREDICTED: uncharacterized protein LOC108154838 [Drosophila
>gi|1037043846|ref|XP_017140746.1| PREDICTED: uncharacterized protein LOC108154840 [Drosophila
>gi|1037043737|ref|XP_017140739.1| PREDICTED: protein crumbs isoform X1 [Drosophila miranda]
>gi|1037043755|ref|XP_017140740.1| PREDICTED: protein crumbs isoform X2 [Drosophila miranda]
>gi|1037044023|ref|XP_017140753.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037043992|ref|XP_017140751.1| PREDICTED: zinc finger CCCH domain-containing protein 14 [D
>gi|1037044008|ref|XP_017140752.1| PREDICTED: uncharacterized protein LOC108154847 [Drosophila
>gi|1037043950|ref|XP_017140748.1| PREDICTED: bromodomain and WD repeat-containing protein 3 is
>gi|1037043962|ref|XP_017140749.1| PREDICTED: bromodomain and WD repeat-containing protein 3 is
>gi|1037043976|ref|XP_017140750.1| PREDICTED: protein RRP5 homolog [Drosophila miranda]
>gi|1037089702|ref|XP_017143913.1| PREDICTED: exosome complex exonuclease RRP44 [Drosophila mi
>gi|1037105468|ref|XP_017145363.1| PREDICTED: acyl-CoA synthetase short-chain family member 3,
>gi|1037106681|ref|XP_017145530.1| PREDICTED: dromyosuppressin [Drosophila miranda]
>gi|1037064318|ref|XP_017142137.1| PREDICTED: RNA-binding protein FUS isoform X4 [Drosophila mi
>gi|1037064288|ref|XP_017142135.1| PREDICTED: RNA-binding protein FUS isoform X2 [Drosophila mi
>gi|1037064302|ref|XP_017142136.1| PREDICTED: RNA-binding protein FUS isoform X3 [Drosophila mi
>gi|1037064272|ref|XP_017142134.1| PREDICTED: RNA-binding protein FUS isoform X1 [Drosophila mi
>gi|1037064210|ref|XP_017142130.1| PREDICTED: C2 domain-containing protein 5 isoform X1 [Drosop
>gi|1037064226|ref|XP_017142131.1| PREDICTED: C2 domain-containing protein 5 isoform X2 [Drosop
>gi|1037064242|ref|XP_017142132.1| PREDICTED: golgin-84 [Drosophila miranda]
>gi|1037064258|ref|XP_017142133.1| PREDICTED: golgin-84 [Drosophila miranda]
>gi|1037091762|ref|XP_017144074.1| PREDICTED: tRNA modification GTPase GTPBP3, mitochondrial [
>gi|1037091898|ref|XP_017144083.1| PREDICTED: uncharacterized protein LOC108156866 [Drosophila
>gi|1037091794|ref|XP_017144076.1| PREDICTED: uncharacterized protein LOC108156860 isoform X1
>gi|1037091806|ref|XP_017144077.1| PREDICTED: uncharacterized protein LOC108156860 isoform X2
>gi|1037091826|ref|XP_017144078.1| PREDICTED: uncharacterized protein LOC108156861 [Drosophila
>gi|1037091840|ref|XP_017144079.1| PREDICTED: uncharacterized protein LOC108156862 [Drosophila
>gi|1037091856|ref|XP_017144080.1| PREDICTED: uncharacterized protein LOC108156863 [Drosophila
>gi|1037091884|ref|XP_017144082.1| PREDICTED: uncharacterized protein LOC108156865 [Drosophila
>gi|1037091870|ref|XP_017144081.1| PREDICTED: uncharacterized protein LOC108156864 [Drosophila
>gi|1037091780|ref|XP_017144075.1| PREDICTED: uncharacterized protein LOC108156859 [Drosophila
>gi|1037035853|ref|XP_017140233.1| PREDICTED: uncharacterized protein LOC108154472 [Drosophila
>gi|1037023461|ref|XP_017139488.1| PREDICTED: uncharacterized protein LOC108153901 [Drosophila
>gi|1037035770|ref|XP_017140228.1| PREDICTED: uncharacterized protein LOC108154468 [Drosophila
>gi|1037023479|ref|XP_017139489.1| PREDICTED: uncharacterized protein LOC108153902 [Drosophila
>gi|1037034559|ref|XP_017140167.1| PREDICTED: LOW QUALITY PROTEIN: uncharacterized protein LOC
>gi|1037023499|ref|XP_017139490.1| PREDICTED: uncharacterized protein LOC108153904 [Drosophila
>gi|1037023517|ref|XP_017139491.1| PREDICTED: uncharacterized protein LOC108153905 [Drosophila
>gi|1037035700|ref|XP_017140223.1| PREDICTED: uncharacterized protein LOC108154464 [Drosophila
>gi|1037074554|ref|XP_017142830.1| PREDICTED: acetylcholine receptor subunit alpha-like 1 [Dros
>gi|1037074616|ref|XP_017142834.1| PREDICTED: uncharacterized protein LOC108156062 [Drosophila
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>gi|1037074632|ref|XP_017142835.1| PREDICTED: uncharacterized protein LOC108156063 [Drosophila
>gi|1037074568|ref|XP_017142831.1| PREDICTED: acetylcholine receptor subunit alpha-like 2 [Dros
>gi|1037074584|ref|XP_017142832.1| PREDICTED: acetylcholine receptor subunit beta-like 2 [Droso
>gi|1037074507|ref|XP_017142826.1| PREDICTED: zinc finger protein DZIP1L isoform X1 [Drosophila
>gi|1037074522|ref|XP_017142827.1| PREDICTED: zinc finger protein DZIP1L isoform X2 [Drosophila
>gi|1037074538|ref|XP_017142828.1| PREDICTED: zinc finger protein DZIP1L isoform X3 [Drosophila
>gi|1037074600|ref|XP_017142833.1| PREDICTED: serine/threonine-protein phosphatase alpha-1 isof
>gi|1037035667|ref|XP_017140222.1| PREDICTED: uncharacterized protein LOC108154461 [Drosophila
>gi|1037105834|ref|XP_017145413.1| PREDICTED: protein takeout [Drosophila miranda]
>gi|1037097586|ref|XP_017144529.1| PREDICTED: protein phosphatase 1 regulatory subunit 21 [Dros
>gi|1037097597|ref|XP_017144530.1| PREDICTED: iron-sulfur cluster assembly 2 homolog, mitochond
>gi|1037056760|ref|XP_017141613.1| PREDICTED: uncharacterized protein LOC108155362 [Drosophila
>gi|1037056776|ref|XP_017141614.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037056792|ref|XP_017141615.1| PREDICTED: probable multidrug resistance-associated protein
>gi|1037065176|ref|XP_017142186.1| PREDICTED: proton-associated sugar transporter A isoform X2
>gi|1037065161|ref|XP_017142185.1| PREDICTED: proton-associated sugar transporter A isoform X1
>gi|1037065191|ref|XP_017142188.1| PREDICTED: proton-associated sugar transporter A isoform X3
>gi|1037065250|ref|XP_017142192.1| PREDICTED: uncharacterized protein LOC108155709 isoform X2
>gi|1037065219|ref|XP_017142190.1| PREDICTED: uncharacterized protein LOC108155709 isoform X1
>gi|1037065234|ref|XP_017142191.1| PREDICTED: uncharacterized protein LOC108155709 isoform X1
>gi|1037065206|ref|XP_017142189.1| PREDICTED: survival of motor neuron-related-splicing factor
>gi|1037091596|ref|XP_017144062.1| PREDICTED: synaptosomal-associated protein 25 isoform X2 [Dr
>gi|1037091583|ref|XP_017144061.1| PREDICTED: synaptosomal-associated protein 25 isoform X1 [Dr
>gi|1037091610|ref|XP_017144063.1| PREDICTED: synaptosomal-associated protein 25 isoform X3 [Dr
>gi|1037091637|ref|XP_017144064.1| PREDICTED: synaptosomal-associated protein 25 isoform X4 [Dr
>gi|1037092359|ref|XP_017144113.1| PREDICTED: 28S ribosomal protein S5, mitochondrial [Drosophi
>gi|1037045773|ref|XP_017140873.1| PREDICTED: casein kinase II subunit alpha [Drosophila mirand
>gi|1037045791|ref|XP_017140874.1| PREDICTED: casein kinase II subunit alpha [Drosophila mirand
>gi|1037045744|ref|XP_017140870.1| PREDICTED: unconventional myosin heavy chain 6 isoform X1 [
>gi|1037045759|ref|XP_017140872.1| PREDICTED: myosin-I heavy chain isoform X2 [Drosophila miran
>gi|1037094869|ref|XP_017144293.1| PREDICTED: DEAD-box helicase Dbp80 [Drosophila miranda]
>gi|1037023535|ref|XP_017139493.1| PREDICTED: LOW QUALITY PROTEIN: probable E3 ubiquitin-prote
>gi|1037100223|ref|XP_017144763.1| PREDICTED: LOW QUALITY PROTEIN: 60S ribosomal protein L15 [
>gi|1037096957|ref|XP_017144470.1| PREDICTED: uncharacterized protein LOC108157108 [Drosophila
>gi|1037096968|ref|XP_017144472.1| PREDICTED: NADH dehydrogenase [ubiquinone] 1 beta subcomplex
>gi|1037098733|ref|XP_017144642.1| PREDICTED: uncharacterized protein LOC108157205 [Drosophila
>gi|1037023553|ref|XP_017139494.1| PREDICTED: cholesterol 7-desaturase, partial [Drosophila mi
>gi|1037089209|ref|XP_017143874.1| PREDICTED: uncharacterized protein LOC108156730 isoform X1
>gi|1037089223|ref|XP_017143875.1| PREDICTED: uncharacterized protein LOC108156730 isoform X2
```