

Gene: ENSG00000165671 - Sequence: ENST00000439151
 Transcript: ENST00000439151 - Protein: ENSP00000395929
 Date : March 2, 2015

1st line: Base numbering. Full stops for intronic +/- 5, 10, 15...
 2nd line: Base sequence. lower case Introns, upper case Exons
 3rd line: Amino acid sequence. Printed on FIRST base of codon
 4th line: Amino acid numbering. Numbered on 1st and increments of 10

Exon 1 | Start: 1401 | End: 1428 | Length: 27

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. . . . . . . . . . . . . . . . .
aacccccgaagagtgagagaagggaaacgcgcgcgcgtcgggtgggggaaggggtgcgcgcgca

. . . . . . . . . . . . . . . . .
ctcggggcccagccgcacgcgggccggcgcgaggcgctcggtcgcacgcgcggccgcggg

. . . . . . . . . . . . . . . . .
ggcgcgcgcggtgggggtgtgaggaggaggaggcgccggcggaataggccggggcaggtc

. . . . . . . . . . . . . . . . .
gcgctcgctgccttctcccctgaagagagacgcggggggagggggggtgcggcgagcggcc

. . . . . . . . . . . . . . . . .
ccgctctctccccaccgctccgctcgacccccagtgtaatgaggggtcacccctccccc

      | -39      | -29      | -19      . . . . .
GCTGCCCCGGGAGGGGGCGCGGGGCACGgtaactagtgcgctgggggtgggcggcgggcag

. . . . . . . . . . . . . . . . .
gcgcgaggagaagggaggaggagggtggccggcggggaagatggtggtggccgtaagg

. . . . . . . . . . . . . . . . .
tgaggggctcgggggagggccaggcgcgatgcgggggttggtggccggcggcgctgcagcc

. . . . . . . . . . . . . . . . .
gccggcctcctccccctccccctcctccatcactaccagccggggtcaggcctagctggc

. . . . . . . . . . . . . . . . .
cgggctgccgcgaacttcctcccggcgcggcccgtgccccgccggccgcctgcgaacacc

. . . . . .
tcggcctccgcctcccctcaggtagcag

```

Exon 2 | Start: 2563 | End: 3506 | Length: 943

.
gggaattttgacgggcagaggggttttaatttttagttcatcccaagtgtccaccagtcta
.
cagaggaggaaaaagagacgggctgtttctatgtagcaggatcggccagcttcgggaaa
.
atggagttttcagaggctcatcgaggccattttttcatctccagtcgggggaactttttc
.
tgcccatggaagtgcagcagaaaggcatagaggccactaggccttgaagtggctgccatt
.
ttaaagagtcgagtcagatggcctattaactcagattaattgctgtgcttttgattcca

 |-9 |1 |11 |21 |31 |41
GTTGATGCCGCCCAGGATGGATCAGACCTGTGAACTACCCAGAAGAAATTGTCTGCTGC
 M D Q T C E L P R R N C L L P
 |1 |11

 |51 |61 |71 |81 |91 |101
CCTTTTCCAATCCAGTGAATTTAGATGCCCCTGAAGACAAGGACAGCCCTTTCGGTAATG
 F S N P V N L D A P E D K D S P F G N G
 |21 |31

 |111 |121 |131 |141 |151 |161
GTCAATCCAATTTTCTGAGCCACTTAATGGGTGTACTATGCAGTTATCGACTGTCAGTG
 Q S N F S E P L N G C T M Q L S T V S G
 |41 |51

 |171 |181 |191 |201 |211 |221
GAACATCCCAAAATGCTTATGGACAAGATTCTCCATCTTGTTACATTCCACTGCGGAGAC
 T S Q N A Y G Q D S P S C Y I P L R R L
 |61 |71

 |231 |241 |251 |261 |271 |281
TACAGGATTTGGCCTCCATGATCAATGTAGAGTATTTAAATGGGTCTGCTGATGGATCAG
 Q D L A S M I N V E Y L N G S A D G S E
 |81 |91

Exon 3 | Start: 59360 | End: 59495 | Length: 135

```
. . . . .
ttttctttattgtgtgtgttttagttctttaaaaaataaagccagcttaattttattttc

. . . . .
aaaatacggatagattcattatataaaaacaattataagttgagaactatgtataaagggc

. . . . .
tatgttatgaacaattaagatggaataatttagttgtacttattttgtaattcttttag

. . . . .
gctagagtgttttcattctcaatttttcatacattgctttttcagaaggctaataaggaat

. . . . .
gacaataatgtttcaaaaatattttgattcttattgatgccccatgttttgtctgtctaaa

      |931      |941      |951      |961      |971      |981
TGTC AACCTAAGAAAAAGTCTACGCCACTGAAGTATGAAGTTGGAGATCTCATCTGGGCA
C  Q  P  K  K  K  S  T  P  L  K  Y  E  V  G  D  L  I  W  A
      |311                        |321

      |991      |1001     |1011     |1021     |1031     |1041
AAATTC AAGAGACGCCCATGGTGGCCCTGCAGGATTGTCTGATCCGTTGATTAACACA
K  F  K  R  R  P  W  W  P  C  R  I  C  S  D  P  L  I  N  T
      |331                        |341

      |1051     |1061     . . . . .
CATTCAAAAATGAAAGGtaataacttgcagtgattatacatgttaaaggcagttgccttta
H  S  K  M  K  V
      |351

. . . . .
gaataactcaattttttgttatcttaataataataatatttttttccttggaacattttgtg

. . . . .
aatgaattggtgttacataattttatcttttggggcttttaaaagttcgttttactctatg

. . . . .
atttaaaccttctctcagtttttttttttgtttgtttgttttgagacggagctctcgct

. . . . .
cttggtgcccaggctggactgtagtggtgcaatctcagctcactgcaacctctgacttgt

. . . . .
gagttcaagcgattct
```

Exon 4 | Start: 71596 | End: 71768 | Length: 172

```
. . . . .
ttacctgtctactatcttaatttcttctttgtagttaatctacatttagctggttttgaa

. . . . .
gctataccacgtcatcagactagactactgattctttcataacctcctgtgtattttctt

. . . . .
gccacacatatataataacttaaaattttccttgtgtatgatctattctagggtgtctagt

. . . . .
tcagtgggcatgttagttgtttccagacagtccttctttggcggcaatgatgtggctgttc

. . . . .
tcttaatgatgagaagtaatttctttgatctaattgattctggttctcttacccttaccta

      |1071      |1081      |1091      |1101      |1111      |1121
TTTCCAACCGGAGGCCCTATCGGCAGTACTACGTGGAGGCTTTTGGAGATCCTTCTGAGA
  S  N  R  R  P  Y  R  Q  Y  Y  V  E  A  F  G  D  P  S  E  R
           |361                        |371

      |1131      |1141      |1151      |1161      |1171      |1181
GAGCCTGGGTGGCTGGAAAAGCAATCGTCATGTTTGAAGGCAGACATCAATTCGAAGAGC
  A  W  V  A  G  K  A  I  V  M  F  E  G  R  H  Q  F  E  E  L
           |381                        |391

      |1191      |1201      |1211      |1221      |1231      .
TACCTGTCCCTTAGGAGAAGAGGGAAACAGAAAGAAAAAGGATATAGGCATAAGgtaggaa
  P  V  L  R  R  R  G  K  Q  K  E  K  G  Y  R  H  K
           |401                        |411

. . . . .
acgaaaaaggctttttattgagtgacagaagcaagtaagaaaaagaaagaaaatggcctc

. . . . .
ttattttattttcgagacagaactttgctctgtttccagattggagtacagtgggtgcaat

. . . . .
ctttgttcactgcaacctccgcgtcccaggttcaagtgattctcgtgcttcagcctcttg

. . . . .
agtagctggtattataagcgagcaccaccacgcctggctaacttttgtatttagtagaga

. . . . .
caggatttcgccatgttgccaggttggtcttgaactcctgagctcaagcagc
```

Exon 5 | Start: 77112 | End: 79671 | Length: 2559

.
ctggccaatatggtgaaacccctattctactgaaaatacaaaaaattagctgggtgtggt
.
ggcggtgcctgtaatcccagctactcgggaggctgaggaaggagaatcacttgaaccg
.
ggaggcagaggttggtgagcgaagattgtgccactgcactccagcctgggcgacagag
.
caagactctgtctcaaaaaaaaaaaaaaaaaaggaataaaaaaaaaagcttctgatttcac
.
tcccttttccccaccatttctttgataagtgataattctttttctcctttaaatata

|1241 |1251 |1261 |1271 |1281 |1291
GTTCTCAGAAAAATTTGAGTAAATGGGAAGCCAGTGTGGACTTGCAGAACAGTATGAT
V P Q K I L S K W E A S V G L A E Q Y D
|421 |431

|1301 |1311 |1321 |1331 |1341 |1351
GTTCCAAGGGGTCAAAGAACCGAAAAATGTATTCCTGGTTCAATCAAGTTGGACAGTGAA
V P K G S K N R K C I P G S I K L D S E
|441 |451

|1361 |1371 |1381 |1391 |1401 |1411
GAAGATATGCCATTTGAAGACTGCACAAATGATCCTGAGTCAGAACATGACCTGTTGCTT
E D M P F E D C T N D P E S E H D L L L
|461 |471

|1421 |1431 |1441 |1451 |1461 |1471
AATGGCTGTTTGAAATCACTGGCTTTTGATTCTGAACATTCTGCAGATGAGAAGGAAAAG
N G C L K S L A F D S E H S A D E K E K
|481 |491

|1481 |1491 |1501 |1511 |1521 |1531
CCTTGCCTAAATCTCGAGCCAGAAAGAGCTCTGATAATCCAAAAAGGACTAGTGTGAAA
P C A K S R A R K S S D N P K R T S V K
|501 |511

1541	1551	1561	1571	1581	1591
AAGGGCCACATACAATTTGAAGCACATAAAGATGAACGGAGGGGAAAGATTCCAGAGAAC					
K	G	H	I	Q	F
			A	H	K
			D	E	R
			R	G	K
			I	P	E
			N		
		521			531
1601	1611	1621	1631	1641	1651
CTTGGCCTAAACTTTATCTCTGGGGATATATCTGATACGCAGGCCTCTAATGAACTTTCC					
L	G	L	N	F	I
			S	G	D
			I	S	D
			T	Q	A
			S	N	E
			L	S	
		541			551
1661	1671	1681	1691	1701	1711
AGGATAGCAAATAGCCTCACAGGGTCCAACACTGCCCCAGGAAGTTTTCTGTTTTCTTCC					
R	I	A	N	S	L
			T	G	S
			N	T	A
			P	G	S
			F	L	F
			S	S	
		561			571
1721	1731	1741	1751	1761	1771
TGTGAAAAAACACTGCAAAGAAAGAATTTGAGACTTCAAATGGTGACTCTTTATTGGGC					
C	G	K	N	T	A
			K	E	F
			E	T	S
			N	G	D
			S	L	L
			G		
		581			591
1781	1791	1801	1811	1821	1831
TTGCCTGAGGGTGCTTTGATCTCAAAGTGTTCTCGAGAGAAGAATAAACCCCAACGAAGC					
L	P	E	G	A	L
			I	S	K
			C	S	R
			E	K	N
			K	P	Q
			R	S	
		601			611
1841	1851	1861	1871	1881	1891
CTGGTGTGTGGTTCAAAAGTGAAGCTCTGCTATATTGGAGCAGGTGATGAGGAAAAGCGA					
L	V	C	G	S	K
			V	K	L
			C	Y	I
			G	A	G
			D	E	E
			K	R	
		621			631
1901	1911	1921	1931	1941	1951
AGTGATTCCATTAGTATCTGTACCATTCTGATGATGGAAGCAGTGACCTGGATCCCATA					
S	D	S	I	S	I
			C	T	T
			S	D	D
			G	S	S
			D	L	D
			P	I	
		641			651
1961	1971	1981	1991	2001	2011
GAACACAGCTCAGAGTCTGATAACAGTGTCTTGAAATTCCAGATGCTTTGATAGAACA					
E	H	S	S	E	S
			D	N	S
			V	L	E
			I	P	D
			A	F	D
			R	T	
		661			671
2021	2031	2041	2051	2061	2071
GAGAACATGTTATCTATGCAGAAAAATGAAAAGATAAAGTATTCTAGGTTTGCTGCCACA					
E	N	M	L	S	M
			Q	K	N
			E	K	I
			K	Y	S
			R	F	A
			A	A	T
		681			691

2621	2631	2641	2651	2661	2671
GGTGAGGATGTCAGTGACTCTGGAACATCAAAGCCATCAAACCATTACTTTTCTCTTCT					
G	E	D	V	S	D
S	G	T	S	K	P
S	K	P	L	L	F
S	S				
		881			891
2681	2691	2701	2711	2721	2731
GCTTCTAGTCAGAATCACATACCTATTGAACCAGACTACAAATTCAGTACATTGCTAATG					
A	S	S	Q	N	H
I	P	I	E	P	D
Y	K	F	S	T	L
L	M				
		901			911
2741	2751	2761	2771	2781	2791
ATGTTGAAAGATATGCATGATAGTAAGACGAAGGAGCAGCGGTTGATGACTGCTCAAAC					
M	L	K	D	M	H
D	S	K	T	K	E
Q	R	L	M	T	A
Q	N				
		921			931
2801	2811	2821	2831	2841	2851
CTGGTCTCTTACCGGAGTCCTGGTCGTGGGGACTGTTCTACTAATAGTCCTGTAGGAGTC					
L	V	S	Y	R	S
P	G	R	G	D	C
S	T	N	S	P	V
G	V				
		941			951
2861	2871	2881	2891	2901	2911
TCTAAGGTTTTGGTTTCAGGAGGCTCCACACACAATTCAGAGAAAAAGGGAGATGGCACT					
S	K	V	L	V	S
G	G	S	T	H	N
S	E	K	K	G	D
G	T				
		961			971
2921	2931	2941	2951	2961	2971
CAGAACTCCGCCAATCCTAGCCCTAGTGGGGGTGACTCTGCATTATCTGGCGAGTTGTCT					
Q	N	S	A	N	P
S	P	S	G	G	D
S	A	L	S	G	E
L	S				
		981			991
2981	2991	3001	3011	3021	3031
GCTTCCCTACCTGGCTTACTGTCCGACAAGAGAGACCTCCCTGCTTCTGGTAAAAGTCGT					
A	S	L	P	G	L
L	S	D	K	R	D
L	P	A	S	G	K
S	R				
		1001			1011
3041	3051	3061	3071	3081	3091
TCAGACTGTGTTACTAGGCGCAACTGTGGACGATCAAAGCCTTCATCCAAATTGCGAGAT					
S	D	C	V	T	R
R	N	C	G	R	S
K	P	S	S	K	L
R	D				
		1021			1031
3101	3111	3121	3131	3141	3151
GCTTTTTTCAGCCCAAATGGTAAAGAACACAGTGAACCGTAAAGCCTTAAAGACCGAGCGC					
A	F	S	A	Q	M
V	K	N	T	V	N
R	K	A	L	K	T
E	R				
		1041			1051

|3701 |3711 |3721 |3731 |3741 |3751
 CGGTTAAATGTTTGTGATAAATCCAGTGCCAGCATTGGTGACATGGAAAAGGAGCCAGGA
 R L N V C D K S S A S I G D M E K E P G
 |1241 |1251

|3761 |3771 |3781 |3791
 ATTCCCAGTTTGACACCACAGGCTGAGCTCCCTGAACCAGgtaaggacatctaaatgtga
 I P S L T P Q A E L P E P A
 |1261

.
 taaaaaaaaaaaaattggagaaagtgctgataaacattgtccttctgaaattggtctggt

.
 gtaatggtaaagtgaagtataaactagcggggaagaatcatcacttcaatgaagaaggag

.
 ttttatgtagaaaggctctgaattttgaagatgagatttccatatatattcaaaagttaaa

.
 ctttaccagaagcccttttccccaaccttttcaagtaaattctctataacatctttctttc

.
 tcactttctctctctctcttttctctctctctctctttttttt

Exon 6 | Start: 103297 | End: 103421 | Length: 124

```
. . . . .
tacaggtgtcagtgactataagacctggctttaaatTTAACCTTTTCCTTAATTTGGAT

. . . . .
gggccacatatattctattatttGTGTATATAAGGTAACACAAGTCCTTTGTCTCCCC

. . . . .
cagctccttttgcatatgagaaaacataagccattttgtgcctcttgcacctaagccat

. . . . .
agtctattttactatgtggtttcccatctggttacttttgggagtatcagatggctcat

. . . . .
aaaaacagtgggttttcctgaagctttttgattaatggtgaatttgtttatcatctttta

      |3801      |3811      |3821      |3831      |3841      |3851
CTGTGCGGTCAGAGAAGAAACGCCTTAGGAAGCCAAGCAAGTGGCTTTTGGAAATATACAG
  V  R  S  E  K  K  R  L  R  K  P  S  K  W  L  L  E  Y  T  E
                |1271                                |1281

      |3861      |3871      |3881      |3891      |3901      |3911
AAGAATATGATCAGATATTTGCTCCTAAGAAAAAACAAAAGAAGGTACAGGAGCAGGTGC
  E  Y  D  Q  I  F  A  P  K  K  K  Q  K  K  V  Q  E  Q  V  H
                |1291                                |1301

|3921. . . . .
ACAAGgtatgttgcaaaatttcagcaaactttcactggtccttaggaaactgcaatttta
K

. . . . .
tcttcaatgtcatactttatcttcatgaacaataatttccttaactgagatcttgtttt

. . . . .
ttattctcagcttctagcacagtacttaggatttagtggttattcaggaaatatataaat

. . . . .
gaatctacatatattgtattttatttaigtatatatTTTTCCCTCTTGCCTTCAAAGAGTAA

. . . . .
tgacttaggaatattgttatgatttaatcttaagtaaaaatataggctataaagtgaaat

.
atata
```

Exon 7 | Start: 105713 | End: 105983 | Length: 270

.
aaattaaaaatcaatacccatataaccaataactcccttttctgctcagcccctggcaatt

.
gccatttctattttctgtctcaagaattttcttattctagatacctcatatacttgaaatc

.
atacgatatttggtctttgtgtcttaagtaatttcccttagcatacataatgtcttcaag

.
gttcatccactttttgtagccttggcagaatttcattccttttaagtggttattctt

.
tttgacacttaaattacaacaattttggcctgtggactctatttttattttttgttctta

 |3931 |3941 |3951 |3961 |3971 |3981
GTAAGTTCCCGCTGTGAAGAGGAAAAGCCTTCTAGCCCGAGGTCGATCTAGTGCTCAGAAC
V S S R C E E E S L L A R G R S S A Q N
 |1311 |1321

 |3991 |4001 |4011 |4021 |4031 |4041
AAGCAGGTGGACGAGAATTCTTTGATTTC AACCAAAGAAGAGCCTCCAGTTCTTGAAAGG
K Q V D E N S L I S T K E E P P V L E R
 |1331 |1341

 |4051 |4061 |4071 |4081 |4091 |4101
GAGGCTCCGTTTTTGGAGGGCCCCCTTGGCTCAGTCAGAACTTGAGGTGGACATGCTGAG
E A P F L E G P L A Q S E L G G G H A E
 |1351 |1361

 |4111 |4121 |4131 |4141 |4151 |4161
TTGCCGCAGCTGACCTTGTCTGTGCCTGTGGCTCCGGAAGTCTCTCCACGGCCTGCCCTT
L P Q L T L S V P V A P E V S P R P A L
 |1371 |1381

 |4171 |4181 |4191
GAGTCTGAGGAATTGCTAGTTAAAACGCCAGgtaaggtggggttggggtctcagtatttg
E S E E L L V K T P G
 |1391

.
agcagatatgattagaggaagcaggagatttttagtatgttttgatgtaaagccaacattg

.
tatctatatacaataaaactaccccccttttgtcctgggaaataacttaaaatgatggttaat
.
tagatatagttactaaccatgaactgtggcacactatcaagaagatgtatttttaataac
.
tatgctccttgctccagtggtgctcttcatgattggtgatcagccactgtgtaaattaac
.
aaccaggataacagggttcaagaggagtac

Exon 8 | Start: 107232 | End: 107341 | Length: 109

```
. . . . .
caccaacaatcagttgttataaaggcagttccgtccttctgacttctcccttagtgattt

. . . . .
tgtgagttaaagtcaccacacacacatccctcacattagttggtgcttg cattgaaaact

. . . . .
gtttacttgttttgtttactaatatttttataataagcaaattaccatcctgcctcttccc

. . . . .
ataagatgacggggaaaacatcaaaaacattgagattcattttgtgtggtatacagattt

. . . . .
tttaaaaattaacttgtgcccagtttctaaatcatctaataagatacatgcatttca

      |4201      |4211      |4221      |4231      |4241      |4251
GAAATTATGAAAGTAAACGTCAAAGAAAACCAACTAAGAAACTTCTTGAATCCAATGATT
  N  Y  E  S  K  R  Q  R  K  P  T  K  K  L  L  E  S  N  D  L
      |1401                                |1411

      |4261      |4271      |4281      |4291      |4301 . . .
TAGACCCCTGGATTTATGCCCAAGAAGGGGGACCTTGGCCTTTCTAAAAAGgtatgttatt
  D  P  G  F  M  P  K  K  G  D  L  G  L  S  K  K
      |1421                                |1431

. . . . .
tttgtaagttctaaaagaaataaaactcaggaaatgagaaattttaaaaatgacattttga

. . . . .
gtagcagttataacattgatgtacatacatatagaaattagtggtgtgtgcagcagtcatt

. . . . .
acatgatctgagatttcctacatgaaaggctgttttgagttgtgtgaccatgtatgtgt

. . . . .
ataaactgggtttcaggtttcctttccaagtgaaaaaaatttttaaatgctttgaagatta

. . . . .
agtttgtgtttgttataattttcttttctgggttcaaatacctcttacctgt
```


Exon 9 | Start: 111671 | End: 111746 | Length: 75

```
. . . . .
tgttggtcaggctggtctttaactctcaacctcaggatccacccgcctcagcctccta

. . . . .
acgtgctaggattacagggtgtgagccatcatgccctgtctggtggagcattttaaaatct

. . . . .
gattcctttccccctgaagtttccgttcaaccctttactgtggtcaggttgatttcttta

. . . . .
attgctaaaacaagtcaaaattcaatatccatggcagctgacaattcagactttggcata

. . . . .
taaagtaaagggtttattttttcatcctctgtaaattggtgttgttttcacttatttata

      |4311      |4321      |4331      |4341      |4351      |4361
TGCTATGAAGCTGGTCACCTGGAGAATGGCATAACTGAATCTTGTGCCACATCTTATTCA
C Y E A G H L E N G I T E S C A T S Y S
      |1441                                |1451

      |4371
AAAGATTTTGGTGGAGgtgagtatttttgagatttaaaaaacgtaatgcagtagtaagtt
K D F G G G

. . . . .
tgaagtgctttgtctgttaaccacaaaaattgttacatgtgtaagcccgaccagtgaggt

. . . . .
acaagttttaaaactggtttgcccaaatctgcttcctgaaaaagaaaaaatattgataa

. . . . .
attgaatatctactctaccaggacttttagtaggaatgctacctgtaatctctctaaagc

. . . . .
gttacagagaggttaggttattatctatagattcctgttttatagaagaaatgtgaagat

. . . . .
gaacaaggtagcaagg
```

Exon 10 | Start: 114154 | End: 114272 | Length: 118

```
. . . . .
gatttaattcctttgaagaactacaagatttgtcatttcaatcatataaattaaaatttc

. . . . .
tgtaataacctatacacttactatttaatctatatgagggagcagttacatatatgtagc

. . . . .
catttcccttcttggtatgtcatcatgagttagctcaattattatttcccccgttttcc

. . . . .
taatccacaaagctggagaattaaatgagttttaaggttggtttttattctaataatagg

. . . . .
ataagagattttggacatgtgtgtagtagccagcagttaacacctattttcctgtcata

|4381      |4391      |4401      |4411      |4421      |4431
GCACTACCAAGATATTTGACAAGCCAAGGAAGCGAAAACGACAGAGGCATGCTGCAGCCA
T T K I F D K P R K R K R Q R H A A A K
|1461                                |1471

|4441      |4451      |4461      |4471      |4481      |4491
AGATGCAGTGTAAAAAAGTGAAAAATGATGACTCGTCAAAAGAGATTCCAGGCTCAGAGg
M Q C K K V K N D D S S K E I P G S E
|1481                                |1491

. . . . .
tattactcagttcctgatcttttcaccttctaagagaagctacttttcacgccagaggc

. . . . .
cacatccctctttcccttgagttttcttaagaatactattctactgaaatggctgatggc

. . . . .
cataaggaatcttgaggaaaagtagagccttatcctaataataaggcaaggcatgtatt

. . . . .
gcagctgtcaatatggtctttttaaaagatcatcacatcctatgctgtaggtgtccaatc

. . . . .
agggtatataagtgtttcatcgttaagtaaacttgtaaaaccggagagtacaatatcaag
```

Exon 11 | Start: 115657 | End: 115800 | Length: 143

```
. . . . .
aaataagtaatcaaataaaacttaagaatgttttctctcccttagtgaaaccttaaaatgg

. . . . .
aacagctcagaaaagttccagtggaaacaaacagcctcagagcagttagtggcagggcatga

. . . . .
ggcgcccactacccgcccacacagcaggggttagaactaacattgcatgcagtcgcccc

. . . . .
gagtgattggctgaacatctgtaagtgcctaatggctagacaaatagcagcccagagggga

. . . . .
gggggtcaaaggaagagacatcaataatacagatgtgggacattatTTTTTctttgcaa

      |4501      |4511      |4521      |4531      |4541      |4551
GGAGAACTAATGCCTCACAGGACGGCCACAAGCCCCAAGGAGACTGTTGAGGAAGGTGTA
G E L M P H R T A T S P K E T V E E G V
      |1501                                |1511

      |4561      |4571      |4581      |4591      |4601      |4611
GAACACGATCCCGGGATGCCTGCCTCTAAAAAATGCAGGGTGAACGCGGTGGAGGAGCT
E H D P G M P A S K K M Q G E R G G G A
      |1521                                |1531

      |4621      |4631      |4641. . . . .
GCACTCAAGGAGAATGTCTGTCAgtagagaaatgtttgccacttggtgttttcattgca
A L K E N V C Q
      |1541

. . . . .
tgttcatctttaagggaaaccactccatctctttatgatggtttcttggtagaataac

. . . . .
aatgcttttggatgattccagtggagtcactttctttaaggatttgacccttttttct

. . . . .
ccccacaaagaaagatacttggagaaatagtttagtggtggtgaaccacgtgcccttaag

. . . . .
gcccatgaaggaatatatgccgcgaacaagtctgagatggaatatgtgaatctctggtt

. . . . .
gcttattcaaataataattctccag
```

Exon 12 | Start: 119206 | End: 119329 | Length: 123

.
agtagctttctggtttatctctatgctcttttctgacctttctactcttgaactcgtttt
.
ccatgtagcagtcagaacagtccttttaaaaaatgaatacctgatttttagctttccctttgt
.
taaagctctccaatggtacacctttgtaactagactaaaatctacaactacgggcccttg
.
cctctttctcacctccttttctgccacttttaacctttgtttactttaaccactgacac
.
tggttttactcttgattctcaaacatggaaaaacagatagatgttttctttctttaaca

 |4651 |4661 |4671 |4681 |4691 |4701
AATTGTGAAAAATTGGGTGAGCTGCTGTTATGTGAGGCTCAGTGCTGTGGGGCTTTCCAC
N C E K L G E L L L C E A Q C C G A F H
 |1551 |1561

 |4711 |4721 |4731 |4741 |4751 |4761
CTGGAGTGCCTTGGATTGACTGAGATGCCAAGAGGAAAAATTTATCTGCAATGAATGTCGC
L E C L G L T E M P R G K F I C N E C R
 |1571 |1581

.
ACAGGtaagtagatatcgaacggtcttcctccaaagaaagtttgaattttaagtttttc
T G

.
aggaaagacatttattggagacactattttgtggcaacactgggctagttgttacagata

.
tagaaatggtgctgtggggtcactgctttcagaaaagctaccaagtgaataataaatggg

.
tagcataatacagtgcaaaaagcattgtgttggtgatacatttagaatgtaatgtaagca

.
aagatgagctgatgccatagaggaagatatctgtgttttagctttaagagagaaagagaaa

.
ttga

Exon 13 | Start: 124427 | End: 124627 | Length: 200

```
. . . . .
cctgtaaacttgtagttagtccaggggcttgattaggtttaatttcagttttttctttt

. . . . .
ttggcgggggtgccagcggcaagcattttctgtagacaatatgtactttctgttgatca

. . . . .
cgtcagggagtatctattaccagattgttcatttataataataactaagattgttacagt

. . . . .
ggttcagacgatgtcaaaccgatcagtcattataaaaatttcttatgaacttttcaccta

. . . . .
atggttttagcatttggtagattcttgaattcttactaatttatcttcttttggttctca

      |4771      |4781      |4791      |4801      |4811      |4821
GAATCCATACCTGTTTTGTATGTAAGCAGAGTGGGGAAGATGTTAAAAGGTGCCTTCTAC
  I  H  T  C  F  V  C  K  Q  S  G  E  D  V  K  R  C  L  L  P
      |1591                        |1601

      |4831      |4841      |4851      |4861      |4871      |4881
CCTTGTGTGGAAAGTTTTACCATGAAGAGTGTGTCCAGAAGTACCCACCCACTGTTATGC
  L  C  G  K  F  Y  H  E  E  C  V  Q  K  Y  P  P  T  V  M  Q
      |1611                        |1621

      |4891      |4901      |4911      |4921      |4931      |4941
AGAACAAAGGGCTTCCGGTGCTCCCTCCACATCTGTATAACCTGTCATGCTGCTAATCCAG
  N  K  G  F  R  C  S  L  H  I  C  I  T  C  H  A  A  N  P  A
      |1631                        |1641

      |4951      |4961      . . . . .
CCAATGTTTCTGCATCTAAAGgtatggatttcttatgtggaccagtctaattgtaaaacc
  N  V  S  A  S  K  G
      |1651

. . . . .
tcagtttaattggcaacagatattttcttttttctttcttttttttttttttttttttga

. . . . .
gacagagtcctcgctgtgttgccgaggctggagtgacagtggcgtgatcttgcccactgca
```

.
agctccgcctcctgggttcacgccattctcctgcttcagcctcctgagtagctgggacta

.
caggcgctgccaccatgcctggctaattttttgtatttttagtagagacggggtttca

.
ccgtgtagccaggatggtct

Exon 14 | Start: 127465 | End: 127644 | Length: 179

```
. . . . .
tctccctatttattctggggctgatttcctagttgtcaggacagtggtttgagttgggct

. . . . .
ttcccttggtgaagcaggctcactgacagttacgggatttgattccttttgagacttat

. . . . .
cacatagctcccatctgagttttcttgatggatcgagtgtttaagatccatcatcttagt

. . . . .
ggtcattccttctatttgtgacattttttcagtgctaacaaatagacttgaataactcac

. . . . .
attcctttttatatttaatatattttgtttttcctttgcttgctccctgattttcctgcttta

      |4971      |4981      |4991      |5001      |5011      |5021
GTCGGTTGATGCGCTGTGTCCGCTGTCCTGTGGCATACCACGCCAATGACTTTTGCCTGG
  R  L  M  R  C  V  R  C  P  V  A  Y  H  A  N  D  F  C  L  A
                |1661                                |1671

      |5031      |5041      |5051      |5061      |5071      |5081
CTGCTGGGTCAAAGATCCTTGCATCTAATAGTATCATCTGCCCTAATCACTTTACCCCTA
  A  G  S  K  I  L  A  S  N  S  I  I  C  P  N  H  F  T  P  R
                |1681                                |1691

      |5091      |5101      |5111      |5121      |5131      |5141
GGCGGGGCTGCCGAAATCATGAGCATGTTAATGTTAGCTGGTGCTTTGTGTGCTCAGAAG
  R  G  C  R  N  H  E  H  V  N  V  S  W  C  F  V  C  S  E  G
                |1701                                |1711

. . . . .
gtaagaaatcatttcttcctctatttgtagtctaaaaagggattaaatcaatgttttaat

. . . . .
tggaaacaaaaatacttttcatcatattgccactggaaaaaatattagaaatgatactatt

. . . . .
catctgccattcaggaaatgatgtcccgaattaatgattacatagcagaactttttttttt

. . . . .
ttttttttttttttttgagacaacgtttcactctgttgcccaggctggagtgctatggca

. . . . .
caatctcagctaactgcagcctccacctcccaggttcaagcagttctcctgcctcagcct
```

Exon 15 | Start: 135038 | End: 135194 | Length: 156

```
. . . . .
acagatgtggttattgtttattagagaaaactccttaccgctgaggaaattatagctcag

. . . . .
caaggaaagtgaatttcccaagggcctgtggtcctagagctagaattccaacctgaacct

. . . . .
gacgagaggttaaataaagcttatgctgtttatactctggtcattatgtgtcactgataat

. . . . .
gtttttatttagtatatcttttagtgaagagaaaagaaaatatatatatatatgtgtatgg

. . . . .
atgtacacatacatgacttgcagtccttgatctgaatgccacatttttttattcccaca

      |5151      |5161      |5171      |5181      |5191      |5201
GAGGCAGCCTTCTGTGCTGTGATTCTTGCCCTGCTGCTTTTCATCGTGAATGCCTGAACA
  G  S  L  L  C  C  D  S  C  P  A  A  F  H  R  E  C  L  N  I
                |1721                                |1731

      |5211      |5221      |5231      |5241      |5251      |5261
TTGATATCCCTGAAGGAAACTGGTATTGCAATGACTGTAAAGCAGGCAAAAAGCCACACT
  D  I  P  E  G  N  W  Y  C  N  D  C  K  A  G  K  K  P  H  Y
                |1741                                |1751

      |5271      |5281      |5291      |5301      .      .      .      .
ACAGGGAGATTGTCTGGGTAAAAAGTTGGACGATACAGgtaagcctgaagaatagcactca
  R  E  I  V  W  V  K  V  G  R  Y  R
                |1761

. . . . .
tctcttttaccatcctctgtttcttgagacctctcagatacaatgcttaacgtatttcta

. . . . .
atgatctacttaattactcatggtactcctcccctcttcttcttgatttttttccttata

. . . . .
ggaagagaaacctaactttatgatttacaattataggaataatacatgttctctgtcgaa

. . . . .
attttttttcttttgtgtattttgtgaaaagtttttttttttttttttaaatgaaaaga

. . . . .
cagtaataatctcacaaggcaggcgcatgtttaacat
```


Exon 16 | Start: 137078 | End: 137283 | Length: 205

.
tcagtgaatgttattgtcataactctctgttcctatatcattatttcctttaagtagaat

.
ccttgaagtagaataattgagtctaagggagtgcgcgctgtgtgggaatgtgggcagat

.
gtttccagcttctagcacatacgacttgtttggttctagttaggttgtaagaatgccg

.
taagatggactttaatgtggacagacagacattgctaatccttacttttatatgagtagg

.
ttattttcctaatagccttgagccttctagaggttttccttctccttttcacctttccca

 |5311 |5321 |5331 |5341 |5351 |5361
GTGGTGGCCAGCTGAGATCTGCCATCCTCGAGCTGTTCCCTTCCAACATTGATAAGATGAG
W W P A E I C H P R A V P S N I D K M R
 |1771 |1781

 |5371 |5381 |5391 |5401 |5411 |5421
ACATGATGTGGGAGAGTTCCCAGTCCTCTTTTTTGGATCTAATGACTATTTGTGGACTCA
H D V G E F P V L F F G S N D Y L W T H
 |1791 |1801

 |5431 |5441 |5451 |5461 |5471 |5481
CCAGGCCCAGTCTTCCCTTACATGGAGGGTGACGTGAGCAGCAAGGATAAGATGGGCAA
Q A R V F P Y M E G D V S S K D K M G K
 |1811 |1821

 |5491 |5501
AGGAGTGATGGGACATATAAAAAAGgtaactttatcctttttgtttctcaggcaaacac
G V D G T Y K K A
 |1831

.
agacctctgttacctgagtgctgatctgttttagaattcacatatgctccattttgaaa

.
ctgcctttgtcctctcagggcattatctggctgcaaatacagtattttgcaaggaagttg

.
acccatgtaactcattatTTTTgagccttaacctttacttaatttgaatttccttgagct

.
TTTTTTgacttatcagttgtttgtcagcattccatcaaataatggagttctgaaacta

.
tttcatagaagaaacacttcagtttg

Exon 17 | Start: 141148 | End: 141260 | Length: 112

```
. . . . .
taagcaaaaaagttctcgtattcagatggatggatgccagtgatctaaagggatagtc

. . . . .
tgtattgtgccatttagttctggattatgttttgtcttaatactatgtctgaataccctt

. . . . .
tggactacattacctgtttgaaaatagctgttatgtgttttcttttcatatagcattgg

. . . . .
tcgatttttgtgtaaaacatggagtttctccaacttaaaggggaaaaagtaaaaaataag

. . . . .
tgaataaataagtaattccacccagagattttgaagtgacttgtgctgtctgttttcata

|5511      |5521      |5531      |5541      |5551      |5561
CTCTTCAGGAAGCTGCAGCAAGGTTTGAGGAATTAAAGGCCCAAAAAGAGCTAAGACAGC
  L  Q  E  A  A  A  R  F  E  E  L  K  A  Q  K  E  L  R  Q  L
                |1841                        |1851

|5571      |5581      |5591      |5601      |5611      |5621 .
TGCAGGAAGACCGAAAGAATGACAAGAAGCCACCACCTTATAAACATATAAAAGgtgagga
  Q  E  D  R  K  N  D  K  K  P  P  P  Y  K  H  I  K
                |1861                        |1871

. . . . .
gaaaatcttgggggaccttctctagaagagaaatggaatagctggctcttcccactctgt

. . . . .
tcatgacaagaacggaagcacaagcatagttcgttttaggtagaggatgagcttaaaag

. . . . .
aagatcagaaatagccgggcacggtggctcacgcttgtaatccagcactttgggaggcc

. . . . .
gaggcgggcggatcacctgaagtcaggagtttgagaccagcctggccaggcaggtgcctg

. . . . .
taattccagctacttgggaggctgaggctggagaattgcttgaaccgggagg
```

Exon 18 | Start: 148041 | End: 148310 | Length: 269

.
accgtacctggctcgtattttatttttttgagatggaatctcaacatatcggccaggctg

.
gagtgcagtggtgaaatcttgtctcactgcaacctccgcctcctgggttcaagcaattct

.
tgtggagttcataattttgtatgtcaagtgaggctctgtttttatatgaaactaagtgttg

.
atagttcaaaatcatgggaaatgtggctgcaacttcaaggaaaaaagtttgcctttttc

.
aggacgtgaattgtcttctgctgacttgttttatgcggtgtactttgtgttacttttcca

 |5631 |5641 |5651 |5661 |5671 |5681
GTAAACCGTCCTATTGGCAGGGTACAGATCTTCACTGCAGACTTATCTGAAATACCCCGT
V N R P I G Q I F T A D L S E I P R
 |1881 |1891

 |5691 |5701 |5711 |5721 |5731 |5741
TGCAACTGTAAAGCTACTGATGAGAACCCCTGTGGGATAGACTCTGAATGCATCAACCGC
C N C K A T D E N P C G I D S E C I N R
 |1901 |1911

 |5751 |5761 |5771 |5781 |5791 |5801
ATGCTGCTCTATGAGTGCCACCCACAGTGTGTCCTGCCGAGGGCGCTGTCAAAACCAG
M L L Y E C H P T V C P A G G R C Q N Q
 |1921 |1931

 |5811 |5821 |5831 |5841 |5851 |5861
TGCTTTTCCAAGCGCCAATATCCAGAGGTTGAAATTTTCCGCACATTACAGCGGGGTTGG
C F S K R Q Y P E V E I F R T L Q R G W
 |1941 |1951

 |5871 |5881 |5891
GGTCTACGGACAAAAACAGATATTAAAAAGgttagaaaaagctaaattaccatatacttt
G L R T K T D I K K
 |1961

.
ctcctctttgcagttgcttgatatcattgatccttgacattagaaaattcatcatagaag

aaaataacacagttaataattaaccttattgttggtccttgccatatccttctaccgttt
agaggcttacgaatggatggttttgatctccaagtcctttgtattgatcatggtcactg
tgtaggaggttggaaaaaatgggacattcatagagatgctatggcctttgtttgctttttt
gtgaatatcctacagctggtaattaagaga

Exon 19 | Start: 149941 | End: 150057 | Length: 116

```
. . . . .
agattggccatccatctctctttataaggacgctctagtttacttactaatcttccattc

. . . . .
tggaggatcttttagtatccagtgtatcctaatactttctgctgctgacagtggtaggagta

. . . . .
aggaggtgtttctcattatcaaatagttggggatcgtaaagtaattatgtgtgcctaaaa

. . . . .
ttatactcatgtaatctgctttgtagaatgtgatgttttcattaattttttaaaaaaatg

. . . . .
tatacatTTggataccagtgtcctTTTTtgccattaagtcaggaggatTTtcttgttcta

      |5901      |5911      |5921      |5931      |5941      |5951
GGTGAATTTGTGAATGAGTATGTGGGTGAGCTTATAGATGAAGAAGAATGCAGAGCTCGA
G E F V N E Y V G E L I D E E E C R A R
                        |1971                        |1981

      |5961      |5971      |5981      |5991      |6001
ATTCGCTATGCTCAAGAACATGATATCACTAATTTCTATATGCTCACCTAGACAAAgtA
I R Y A Q E H D I T N F Y M L T L D K
                        |1991                        |2001

. . . . .
agtaatgggaaatgctgttttcactgttacaagattgtaaatttgtgttgtcccagccat

. . . . .
agtatttgtaggcatgtaacgcagttcccaaggtagggcttttcccataccattggga

. . . . .
ttgctggattggggttctggagtaggtaagagtgctcattttctgggtattcatggccag

. . . . .
aggcaatacagcagcttttagcattgtttttgaacaattcaatgaaaattggagcttata

. . . . .
tctttgagtattttttctacaggattacaagtttgaactcagaaaagtacataattc
```

Exon 20 | Start: 151263 | End: 151404 | Length: 141

```
. . . . .
tcaatgtagtcagaaacccctggcagccttggtatatataacgcagaatgctgcacccca

. . . . .
cagtctctgcttttagtgatattgaagcacaacccaggaatttgcattgctaacaaattgc

. . . . .
caaatggtgatggtgctggtccagggatcacactttgagaaccaaagttctttgggatct

. . . . .
tttctctgagaggttcagtcctttacaaatagaaaactccaacttattagagagaatagtca

. . . . .
aattttaatccacagcagaggtctcaggaagtctgatgtgtagcttcttttgaattcta

|6011      |6021      |6031      |6041      |6051      |6061
GACCGAATCATTGATGCTGGTCCCAAAGGAAACTATGCTCGGTTTCATGAATCATTGCTGC
D R I I D A G P K G N Y A R F M N H C C
                        |2011                        |2021

|6071      |6081      |6091      |6101      |6111      |6121
CAGCCCAACTGTGAAACACAGAAGTGGTCTGTGAATGGAGATACCCGTGTAGGCCTTTTT
Q P N C E T Q K W S V N G D T R V G L F
                        |2031                        |2041

|6131      |6141      |6151. . . . .
GCACTAAGTGACATTAAAGCAGgtaagaatcatttcaggattctgcagctgacatctgaa
A L S D I K A G
                        |2051

. . . . .
tttcagggcttttgttttttacaacagcttcctcagattataattttaacatttctat

. . . . .
atgtaccgttctctggaattaagttcatttacatttttgtgcaaccatcaccaccatccg

. . . . .
tctccagaactcttttcatcttcccctgctaaaactctatacctactaaacactaagtcc

. . . . .
ccattcttcttttcttcacagcccctggcaactaccattctacttttctgtctctgaatt

. . . . .
tgactactatatgtacttcata
```

Exon 21 | Start: 156295 | End: 156401 | Length: 106

.
gaattaacataatgttgaatctgaatggttaataaacaatgaaaaacaacttgaggtaat
.
taacacctctggaggaagagttatcttaaatattacttggttaagccatttaagttttctc
.
tgttaaattgtattacctgattattaatggtataagattagaattaattaaaatatagga
.
ttagagctaaattctttttaaaattaatctgttctcttgggagttggtatcctttgtaat
.
taatacagaaataatgtaattaaaaccatagatattaatattttcacggtctcttatgca

 |6161 |6171 |6181 |6191 |6201 |6211
GCACTGAACTTACCTTCAACTACAACCTAGAATGTCTTGGGAATGGAAAGACTGTTTGCA
 T E L T F N Y N L E C L G N G K T V C K
 |2061 |2071

 |6221 |6231 |6241 |6251 . .
AATGTGGAGCCCCGAAGTGCAGTGGCTTCTTGGGTGTAAGGCCAAAGgtaccacccttct
 C G A P N C S G F L G V R P K
 |2081

.
agacttctgctttgggattagtggtgcggtcctccctccctaacagtgttagggcagtct
.
acattttaagaagaaatctctttgaaattaatgacattcagccaggcatggtggctcatg
.
cctgtaatcccagcactttgggaggctgaggtgggcggatcacctgaggtcaggagttcg
.
agaccagcctggccaacatgatgaaaccccatctctaataagaatagaaaaattagctgg
.
gccgtggtggcatgtgcttgtagtcccagctactcaggaggctgagg

Exon 22 | Start: 159430 | End: 159634 | Length: 204

```
. . . . .
gaaaaataaataaattaactaataaaaacttatgaaaggcatgtcatttaacaccag

. . . . .
ccttatggatcagcaaaaacacaaagatttttgtcttctgttttgttcaggtggcaggag

. . . . .
tagttgatacaaaaattattgctacgtatctgtgatgtaccaggtacctttcctgcattt

. . . . .
tatcttcatgacaatctcttttcccagagaagagaatgaggctcagagagggtagttaac

. . . . .
ccggttaagattggtactaatgtgttcacagaatgctgactgttcaatatctgacctgta

|6261      |6271      |6281      |6291      |6301      |6311
AATCAACCCATTGCCACGGAAGAAAAGTCAAAGAAATTCAAGAAGAAGCAACAGGGAAAG
N Q P I A T E E K S K K F K K K Q Q G K
|2091                                |2101

|6321      |6331      |6341      |6351      |6361      |6371
CGCAGGACCCAGGGTGAAATCACAAAGGAGCGAGAAGATGAGTGTTTTAGTTGTGGGGAT
R R T Q G E I T K E R E D E C F S C G D
|2111                                |2121

|6381      |6391      |6401      |6411      |6421      |6431
GCTGGCCAGCTCGTCTCCTGCAAGAAACCAGGCTGCCCAAAAGTTTACCACGCAGACTGT
A G Q L V S C K K P G C P K V Y H A D C
|2131                                |2141

|6441      |6451      |6461      . . . . .
CTCAATCTGACCAAGCGACCAGCAGgttgggtgccaaaatccatttgtaccgctactcgtt
L N L T K R P A G
|2151

. . . . .
ctctccatcatactcaggggtctcatgccatttgcacagccttgaagaaagtgccatttg

. . . . .
gtctttccatgcataattttgaggggtatggtctattttcttattttggaaataatgaaa
```

gcaaaggcttagaattcttagttatgaagaatgggacagcgggtgcacttgctgtgtttca
ggtcctttacttacagtcatgtttatttcattccccagcagcccataatggttggtttt
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ACCTGGCAGAGCAATCAACAGGAATGGCTGCTCAGGCACCCAAAATGTCAGATAAACCTC
L A E Q S T G M A A Q A P K M S D K P P

|6771 |6781 |6791 |6801 |6811 |6821
 CTGCTGACACCAACCAGATGCTGTCGCTCTCCAAAAAGCTCTGGCAGGGACTTGTGAGA
 A D T N Q M L S L S K K A L A G T C Q R
 |2261 |2271

|6831 |6841 |6851 |6861 |6871 |6881
 GGCCATTGCTACCTGAAAGACCTCTTGAGAGAACTGACTCCAGGCCCCAGCCTTTAGATA
 P L L P E R P L E R T D S R P Q P L D K
 |2281 |2291

|6891 |6901 |6911 |6921 |6931 |6941
 AGGTCAGAGACCTCGCTGGGTCAGGGACCAAATCCCAATCCTTGGTTTCCAGCCAGAGGC
 V R D L A G S G T K S Q S L V S S Q R P
 |2301 |2311

|6951 |6961 |6971 |6981 |6991 |7001
 CACTGGACAGGCCACCAGCAGTGGCAGGACCAAGACCCCAGCTAAGCGACAAACCCTCTC
 L D R P P A V A G P R P Q L S D K P S P
 |2321 |2331

|7011 |7021 |7031 |7041 |7051 |7061
 CAGTGACCAGCCCAAGCTCCTCACCTCAGTCAGGTCCCAACCACTGGAAAGACCTCTGG
 V T S P S S S P S V R S Q P L E R P L G
 |2341 |2351

|7071 |7081 |7091 |7101 |7111 |7121
 GGACGGCTGACCCAAGGCTGGATAAATCCATAGGTGCTGCCAGCCCAAGGCCCCAGTCAC
 T A D P R L D K S I G A A S P R P Q S L
 |2361 |2371

|7131 |7141 |7151 |7161 |7171 |7181
 TGGAGAAAACCTCAGTTCCCACTGGCCTGAGACTTCCGCCGCCAGACAGACTGCTCATTA
 E K T S V P T G L R L P P P D R L L I T
 |2381 |2391

|7191 |7201 |7211 |7221 |7231 |7241
 CTAGCAGTCCCAAACCCAGACTTCAGACAGGCCTACTGACAAACCCCATGCCTCTTTGT
 S S P K P Q T S D R P T D K P H A S L S
 |2401 |2411

|7251 |7261 |7271 |7281 |7291 |7301
 CCCAGAGACTCCCACTCCTGAGAAAGTACTATCAGCTGTGGTCCAGACCCTTGTAGCTA
 Q R L P P P E K V L S A V V Q T L V A K
 |2421 |2431

7311	7321	7331	7341	7351	7361
AAGAAAAAGCACTGAGGCCTGTGGACCAGAATACTCAGTCAAAAAATAGAGCTGCTTTGG					
E	K	A	L	R	P
		V	D	Q	N
		T	Q	S	K
		N	R	A	A
		L	V		
	2441			2451	
7371	7381	7391	7401	7411	7421
TGATGGATCTCATAGACCTAACTCCTCGCCAGAAGGAGCGGGCAGCTTCACCTCATCAGG					
M	D	L	I	D	L
		T	P	R	Q
		K	E	R	A
		A	A	S	P
		H	Q	V	
	2461			2471	
7431	7441	7451	7461	7471	7481
TCACACCACAGGCTGATGAGAAGATGCCAGTGTGGAGTCAAGTTCATGGCCTGCCAGCA					
T	P	Q	A	D	E
		K	M	P	V
		L	E	S	S
		S	S	W	P
		A	S	K	
	2481			2491	
7491	7501	7511	7521	7531	7541
AAGGTCTGGGGCATATGCCGAGAGCTGTTGAGAAAGGCTGTGTGTCAGATCCTCTTCAGA					
G	L	G	H	M	P
		R	A	V	E
		K	G	C	V
		S	D	P	L
		Q	T		
	2501			2511	
7551	7561	7571	7581	7591	7601
CATCTGGGAAAGCAGCAGCCCCTTCAGAGACCCCTGGCAAGCTGTTAAATCACTCACCC					
S	G	K	A	A	A
		P	S	E	D
		P	W	Q	A
		V	K	S	L
		T	Q		
	2521			2531	
7611	7621	7631	7641	7651	7661
AGGCCAGACTTCTTTCTCAGCCTCCTGCCAAGGCCTTTTTATATGAGCCAACAACCTCAGG					
A	R	L	L	S	Q
		P	P	A	K
		A	F	L	Y
		E	P	T	T
		Q	A		
	2541			2551	
7671	7681	7691	7701	7711	7721
CCTCAGGAAGAGCTTCTGCAGGGGCTGAGCAGACCCCAGGGCCTCTTAGCCAATCCCCGG					
S	G	R	A	S	A
		G	A	E	Q
		T	P	G	P
		L	S	Q	S
		P	G		
	2561			2571	
7731	7741	7751	7761	7771	7781
GCCTGGTGAAGCAGGCGAAGCAGATGGTCGGAGGCCAGCAACTACCTGCACTTGCCGCCA					
L	V	K	Q	A	K
		Q	M	V	G
		G	Q	Q	L
		P	A	L	A
		A	A	K	
	2581			2591	
7791	7801	7811	7821	7831	7841
AGAGTGGGCAATCTTTTAGGTCTCTCGGGAAGGCCCCAGCCTCCCTCCCCACTGAAGAAA					
S	G	Q	S	F	R
		S	L	G	K
		A	P	A	S
		L	P	T	E
		E	K		
	2601			2611	

*481	*491	*501	*511	*521	*531
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*541	*551	*561	*571	*581	*591
AACCTTTGACCAGGAAGTAACAGGAAGTTCTGAGGGGCCCTGGGGCTTTAGACTCATTTT					
*601	*611	*621	*631	*641	*651
GAAATGTCCCTTTGTGGCACCAGAAAGTGGTTGTGTTGAGGAAGTGTCTCTTGGCTGCGGTG					
*661	*671	*681	*691	*701	*711
TGCATGGGTGCGTGTGCATGCGCGCACACTCACAGAGGTCTCCTCTATAGATGCAAGGGT					
*721	*731	*741	*751	*761	*771
GCTGCATTGAGGCCAGCAAGGCTGTTGGCTGTGGGGTCGCCGCTGCTGCTTTTGTCTGGG					
*781	*791	*801	*811	*821	*831
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*841	*851	*861	*871	*881	*891
GTGCGAATGTAGATTTCCAGCAGTGGAAGAAGGCATTTGGCAAGCTTCTCTTTCTTTGCT					
*901	*911	*921	*931	*941	*951
TTTGTCTTCTACCTATTTTTCTCTTTGTACATGAATCCACCCCATCCCTATTTCCCTAAAA					
*961	*971	*981	*991	*1001	*1011
CACTCAGGTGCTTTTCAGATTTTCAGAGCCTCGGGCAGTGGACATAGGGAATCTCTGGCAAG					
*1021	*1031	*1041	*1051	*1061	*1071
CTCTGAGCTAGACACACCAGCTTCAGGAAGAGTACCAGATCCTGATGGGAAATTTCTTTT					
*1081	*1091	*1101	*1111	*1121	*1131
CCCCATTCCTTTTCCCTCCTGAGTGGAGGGAGTCCTCTTCTTCGCCTCCCTGAGAATTGC					
*1141	*1151	*1161	*1171	*1181	*1191
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*1201	*1211	*1221	*1231	*1241	*1251
CCCTGAAGACTGGCAAGAGGTGGTGTGTTAGAGCAACGTCCAGGCTAAGAGATGACTCCTA					
*1261	*1271	*1281	*1291	*1301	*1311
TTAACTGCTGATTATCTGTTACTGCTGCCCTGAGCTGGGGCCCAAGGGCTGGGAAATCTG					
*1321	*1331	*1341	*1351	*1361	*1371
TTGGTGCTACCCCTGCCCTACCATTCACCCAGCTCACAGACTGCCAACAGGAAGTGCTGTT					

|*1381 |*1391 |*1401 |*1411 |*1421 |*1431
 TGGCTAGTTTCCTCCCCTTGTCTACCCCTCCTTTGTCCTTAGACCAACATGTTTACCTC

|*1441 |*1451 |*1461 |*1471 |*1481 |*1491
 TCTGCTTTGCCAACTTAGCCAGCAGGCCATCCCCGGCCCTAACGTCTCCTGGCCATTATC

|*1501 |*1511 |*1521 |*1531 |*1541 |*1551
 TCTTAGTTATGGCTTTCACGCTCTCAATAGGATTCTGTATTTGGTCCCAATTTCTCAAG

|*1561 |*1571 |*1581 |*1591 |*1601 |*1611
 TTCTTATTGAGGTTACTCCCATCAATTCCACGGAGGGAACAGTAGTTATTATAGAAGCAT

|*1621 |*1631 |*1641 |*1651 |*1661 |*1671
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|*1681 |*1691 |*1701 |*1711 |*1721 |*1731
 AGGTTGAGACACTTGAACCTCAGGCAGAGGGACGAGGCTGGGCAGGGCTGTCCTGAGTTTA

|*1741 |*1751 |*1761 |*1771 |*1781 |*1791
 GGGGCCTATCCCTGCATTTCACTGAGACCTCGGAATCTCCTCTGTGAATTCCACCTGCCT

|*1801 |*1811 |*1821 |*1831 |*1841 |*1851
 AGTTCTCCCTTTTCATCCTCTCTCTCTTCCCACATCATCAAAGAGGAAAAGCTCTTTGTT

|*1861 |*1871 |*1881 |*1891 |*1901 |*1911
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|*1921 |*1931 |*1941 |*1951 |*1961 |*1971
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|*1981 |*1991 |*2001 |*2011 |*2021 |*2031
 ATGTGTGAGAATAATAAATATATAACTGCAGCTAGTAGGTCCCTTTCCCTAATCTTTTAG

|*2041 |*2051 |*2061 |*2071 |*2081 |*2091
 GTCATATGAGTAGGGTTTGCTTGGTGCCAGTCCTGTGCCCTTTTCTCTCCAGTCATCTGT

|*2101 |*2111 |*2121 |*2131 |*2141 |*2151
 AGTTGTGATCAGAAAAAGGTATCTGCACTGCACTGTCAGAGTCTCCTTTCACTATGTTGT

|*2161 |*2171 |*2181 |*2191 |*2201 |*2211
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|*2221 |*2231 |*2241 |*2251 |*2261 |*2271
 GGGAGGGCGTGCAGGCCATGTAAAAATTTCCGTGGAGAAGTTTGATTCTAAAGTAGCTT

|*2281 |*2291 |*2301 |*2311 |*2321 |*2331
 CTCTAAAGTAGGCTTTGGTAGGTAATCAACTTGACAGCAGTCTAGATGTCTCACAGGACA

 |*2341 |*2351 |*2361 |*2371 |*2381 |*2391
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 |*2401 |*2411 |*2421 |*2431 |*2441 |*2451
 CATTCTCCGCCATTCAATTGGAGGCTTCGTTCCAGACCTGCCTGGGAAAACAGCTTCTGAG

 |*2461 |*2471 |*2481 |*2491 |*2501 |*2511
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 |*2521 |*2531 |*2541 |*2551 |*2561 |*2571
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 |*2581 |*2591 |*2601 |*2611 |*2621 |*2631
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 |*2641 |*2651 |*2661 |*2671 |*2681 |*2691
 GAGTCACAACAACTGTTGGGAAATGTGACTCTTGGAGGAAGGTGGGGAGGGAGTGGCCTT

 |*2701 |*2711 |*2721 |*2731 |*2741 |*2751
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 |*2761 |*2771 |*2781 |*2791 |*2801 |*2811
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 |*2821 |*2831 |*2841 |*2851 |*2861 |*2871
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 |*2881 |*2891 |*2901 |*2911 |*2921 |*2931
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 |*2941 |*2951 |*2961 |*2971 |*2981 |*2991
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 |*3001 |*3011 |*3021 |*3031 |*3041 |*3051
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 |*3061 |*3071 |*3081 |*3091 |*3101 |*3111
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 |*3121 |*3131 |*3141 |*3151 |*3161 |*3171
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*3181	*3191	*3201	*3211	*3221	*3231
ATTTTATTCTTTACATCCTTCACCCACACTATGGTCAGGGCATGAAACACCCTGTTGAT					
*3241	*3251	*3261	*3271	*3281	*3291
CCCTTCCCAGGCTCGGCACTGTCTGCTCACTGGAGCCGGAAGTCCAGGTTGTAATTCTAA					
*3301	*3311	*3321	*3331	*3341	*3351
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*3361	*3371	*3381	*3391	*3401	*3411
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*3421	*3431	*3441	*3451	*3461	*3471
TACTAAGGTTTTTTAAATGATACTGTCATCCTCTTGGGGTTTATCAGCCAGGTTAGAGGA					
*3481	*3491	*3501	*3511	*3521	*3531
GCCCAGTGTCTAACCTCTCTCAGATCATGGCAGAGAAGGAGCTGCCTCCAGCCCCCTTTC					
*3541	*3551	*3561	*3571	*3581	*3591
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*3601	*3611	*3621	*3631	*3641	*3651
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*3661	*3671	*3681	*3691	*3701	*3711
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*3721	*3731	*3741	*3751	*3761	*3771
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*3781	*3791	*3801	*3811	*3821	*3831
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*3841	*3851	*3861	*3871	*3881	*3891
TAGAAAAGTTTGAGGCTTTGTTAAAAGTGGGGAGAAGGAAGATCCTCAGTGAAGCCTGCA					
*3901	*3911	*3921	*3931	*3941	*3951
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*3961	*3971	*3981	*3991	*4001	*4011
GGTGGCCATTGAGTTTCTCAGGGCTGGGGCCACCTTGTCATAGCCTCCGTCCACGCTGC					
*4021	*4031	*4041	*4051	*4061	*4071
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*4081	*4091	*4101	*4111	*4121	*4131
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*4141	*4151	*4161	*4171	*4181	*4191
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*4201	*4211	*4221	*4231	*4241	*4251
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*4261	*4271	*4281	*4291	*4301	*4311
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*4321	*4331	*4341	*4351	*4361	*4371
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*4381	*4391	*4401	*4411	*4421	*4431
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*4441	*4451	*4461	*4471	*4481	*4491
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*4501	*4511	*4521	*4531	*4541	*4551
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*4561	*4571	*4581	*4591	*4601	*4611
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*4621	*4631	*4641	*4651	*4661	*4671
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*4681	*4691	*4701	*4711	*4721	*4731
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