

Gene: SLC25A13 - Sequence: NG\_012247.1  
 Transcript: NM\_014251.2 - Protein: NP\_055066.1  
 Date : March 2, 2015

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

Exon 1 | Start: 5001 | End: 5206 | Length: 205

```

. . . . .
caacttccccaggccgtgcaccatctgtcgttacttgcccctcccatcggtccgagggcc

. . . . .
cagtagcccctggtatcagcagaagccagcagcttcctggtggggccacctgccctcag

. . . . .
gtgacgggcttgacacggccaagttacctctgcccacgccgcagaggccgccgtgctg

. . . . .
gacagcgcagcggggtcacgtgtccctgtccagccactgccccaggatctgagccctccc

. . . . .
cggctgcagggcagaggccgctggatcacgcgcaccccggcgccaaggctgggcccagc

|-189      |-179      |-169      |-159      |-149      |-139
AATGGGCGGGCAGCATCCACATGACCCGCGCCGGCGGGAGGGCGTGGGGAGGCAGGCCAG

|-129      |-119      |-109      |-99       |-89       |-79
GAACGCACGCTGCTTGCCGTATCGCCGCCCCACCGCCGCCGCCGGGACTAGAAAGT

|-69       |-59       |-49       |-39       |-29       |-19
GAGCCGCCCCGGGTCCCAAACGCCAGCCAGCCAGTCAGTGGGTCCCGCAGTCGCCCCGAAC

|-9        |1         |11        . . . . .
CGGGGCGAATCATGGCGGCCGCAAGgtaaccgcgggcccaggccggggggagcgcgcc
      M  A  A  A  K
      |1

. . . . .
tgccggatcagcctgggggtgcgagaggctcgggcgctcacgtgtctgggttggtgccac

. . . . .
aggcggagcaaaatggggtgccgggcagccgttccaccttgccggtgctggaggaggagg

```

. . . . .  
gccactcctcccgggcagcacaggcggtcggcgcagagaccgaggcccgacctcgacgcg  
. . . . .  
gaaactttgtggccggacactcccacttggccggggagggaagttggtgcctgacgggg  
. . . . .  
ctgagccagggccgacgaggaggagc

Exon 2 | Start: 30197 | End: 30250 | Length: 53

```
. . . . .
aaaatgacaggtgtttgcaatgtttcaccttttctttttaagaaatgatatacattcaac

. . . . .
attagatattaagagattttgctactggttccttgcataaaattttatcattttagaagc

. . . . .
tgcataaggagtctaatttaaaatgggaaggggcatagtttatgcactggggcaacatg

. . . . .
gtatttaagtacaaaaatgaaatgcactaatttttaatctgagatttattttcctttagt

. . . . .
cggctagcttgatttctcagctttgttgtaataacataaacatttttatgtttatttata

      |21      |31      |41      |51      |61      .
GTGGCTTTAACCAAGAGAGCAGATCCAGCTGAGCTTAGAACAATATTTTTGAAGgtcagt
V  A  L  T  K  R  A  D  P  A  E  L  R  T  I  F  L  K
                |11                                |21

. . . . .
ataaaggaatctattttgtttcttgtttgatttggttctataacttttaaatattattt

. . . . .
tcaaacagctctttatttcgttgttatattttgaaaaagtcccaaaggtgcagcccggca

. . . . .
cggtggtcatgcctgtaatcccagcactttgggaggctgaggtgggtggatcccctgag

. . . . .
gtcaggagtttgagaccagcctggccaacatggtgaaaccccatctttactgaaagtacc

. . . . .
aaaactagccaggtgtggtggcacgtgcctgtaatcctggctattcaggaggct
```

Exon 3 | Start: 49810 | End: 49952 | Length: 142

```
. . . . .
ctggcctctacctgctggagccagtagcacccttcctcacatggttatgacaatcttaatt

. . . . .
gccatcagatattgccaaagtgtccctgtgggaaaacattgctcccagttaagtttataaa

. . . . .
cataaaacttgacttggtttttctttgaactggtgggagataatggtcattatgaagcgt

. . . . .
acatgatcatatctgatattttatcgtttggaatattttcaactttttcacatatcactct

. . . . .
catgattatatactgttgaaaatcaaatatataaaaatactgtttttcttttttttaaca

|71      |81      |91      |101     |111     |121
TATGCAAGCATTGAGAAAAACGGTGAATTTTTCATGTCCCCCAATGACTTTGTCACTCGA
Y A S I E K N G E F F M S P N D F V T R
|31                      |41

|131     |141     |151     |161     |171     |181
TACTTGAACATTTTGGAGAAAGCCAGCCTAATCCAAAGACTGTGGAACTTTAAAGTGGA
Y L N I F G E S Q P N P K T V E L L S G
|51                      |61

|191     |201     |211     . . . . .
GTGGTGGATCAGACCAAAGATGGgtatgtttatttttaattattttatttaattcttttt
V V D Q T K D G
|71

. . . . .
gtttgaaacagctttcttttgctctaataccaggaagtcattaaatgtataccttccca

. . . . .
tctcttaggaccattttctgtacttactactattattattacattatttgagtaataatg

. . . . .
tagttccttcaatcattgtttcaagttggggattcaatatgtatataattgactctttta

. . . . .
aagaatgctttatcgtatcagagaactattattgtcaacaagggtggtgatgtgcagatgt

. . . . .
atgagagtattttatatatgcctttg
```

ccaaaaacatgcctgaaggaagactgaaacatatatgatgctggataaaaacagttatatt  
.  
tgaaattataaatgtatatttagtgtgtaacttcattggtcaatgctgtcatgtcttttagatt  
.  
ctgtgcattattatttcagtcactatatgtaaaagtaaagctaaatgtttataaatgt  
.  
tttcagtttgaaattttatgttttggcagtgatatcttattttttccctcagtgtagttt  
.  
gtatatgcttctgtttttccacaagctactgacttttgctttatgtttgttttatgttgca  
|221 |231 |241 |251 |261 |271  
ATTAATATCTTTTCAAGAATTTGTTGCCTTTGAATCTGTCTGTGTGCCCTGATGCTTT  
L I S F Q E F V A F E S V L C A P D A L  
|81 |91  
|281 |291 |301 |311 |321  
GTTTATGGTAGCCTTTTCTGCTGTTTGACAAAGCTGGCAAAGGAGAAGTAACTTTTGgtaa  
F M V A F Q L F D K A G K G E V T F E  
|101

5

Exon 5 | Start: 118171 | End: 118310 | Length: 139

```
. . . . .
caaatgagatgataatagtttttagcttttctacttgggtttcctacagcctttaggttggtg

. . . . .
tgtgtgtgagagacttcttgattttccccccctttattttatttgggtgggattcttgacc

. . . . .
ttgattttcttaactgattccttgagggttcttttattttctgccatagtttgtaatctgta

. . . . .
tgtctattttaaaattgaattactttttttaataagaaaaagcgatatccattcaataaca

. . . . .
gattttatcagaaagaacttcatttactttgttttaataaacctgtcttttctcttttca

|331      |341      |351      |361      |371      |381
AGGATGTTAAGCAAGTTTTTGGACAGACCACAATTCATCAACATATTCCATTAACTGGG
D V K Q V F G Q T T I H Q H I P F N W D
|111                                |121

|391      |401      |411      |421      |431      |441
ATTCAGAATTTGTGCAACTACATTTTGGAAAAGAAAAGAAAAGACACCTGACATATGCGG
S E F V Q L H F G K E R K R H L T Y A E
|131                                |141

|451      |461      . . . . .
AATTTACTCAGTTTTTATTGgtgggtctagctcttaaaaagttaaagggtatggccggg
F T Q F L L
|151

. . . . .
cacggtgggtcacacctataatcctagcactttgggaggccgaggtgggcggatgacgag

. . . . .
gtcaggagatcgagaccatcctggccaacatggtgaaaccccgctcttactaaaaataca

. . . . .
aaaaattagccaggcgtggtggccggcgctgtggtcccagctacttgggagggtgaggg

. . . . .
aggagaatggtgtgaacttgggaggtggagcttgagtgagctgagatcgaccactgca

. . . . .
gtccggcctgggtgaaagag
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Exon 6 | Start: 133965 | End: 134111 | Length: 146

```
. . . . .
catttcattccggttttcattttctttgtgttcattggactctagagagcacggaagtaat

. . . . .
tgggtttcattgggtagcagagtaaggtagagtgcactttctcaagacatgttcattggt

. . . . .
ctgagcaaagtctgtctcttggaggagacagtagcttggtggtgctattaggcagatg

. . . . .
agggcttgtagatcaagatgtaaactctctttcttcagtctgattttaaaatgaactgt

. . . . .
catttgtagtaattattaaatgagaagcataaaattaagtattttactttctttatttgta

|471      |481      |491      |501      |511      |521
GAAATACAACTGGAGCACGCAAAGCAAGCCTTTGTGCAACGGGACAATGCTAGGACTGGG
E I Q L E H A K Q A F V Q R D N A R T G
|161                                |171

|531      |541      |551      |561      |571      |581
AGAGTCACAGCCATCGACTTCCGAGACATCATGGTCACCATCCGCCCCCATGTCTTGACT
R V T A I D F R D I M V T I R P H V L T
|181                                |191

|591      |601      |611      . . . . .
CCTTTTGTAGAAGAAATGTCTAGTAGCTgtaagttgtaacttgcctaataagaagtggttta
P F V E E C L V A
|201

. . . . .
actcagtaaataaggaattataaaaaacaattcttataagttatgtagtggtttttttctg

. . . . .
ttgcaaacactaacatttttaatgtgatcacatacattactaaactcaaacaagagtggtg

. . . . .
ttttggctagtttttagtgggtaggttaatttggtgtctgggtaatacgggaatatcttcga

. . . . .
gaaatacattagcttttttaagataaaaggaaaaacagatttgtgatctcatgttaattt

. . . . .
tgtttttagattttgaagtatatcaagg
```

Exon 7 | Start: 135901 | End: 136039 | Length: 138

```
. . . . .
catttccaattgaaatttgaactctaaattactactagtcctgttataatcctacacatg

. . . . .
tattttatttagttttctaggagttgtgtatggagagtacaagttctggcatgattagtt

. . . . .
gcagttgcttctaaaaatcatttatagcttttttttaatgcaaaagaaaaatacttatag

. . . . .
tcatttcttctgtacatgcatttttgattactttcaaactagtttttagatgtatgtact

. . . . .
tctgaatcatctataaatcttttgatatattttcttctctcttgttttcaatttatttga

      |621      |631      |641      |651      |661      |671
GCTGCTGGAGGTACCACATCCCATCAAGTTAGTTTCTCCTATTTTAATGGATTTAATTTCG
A  A  G  G  T  T  S  H  Q  V  S  F  S  Y  F  N  G  F  N  S
                |211                        |221

      |681      |691      |701      |711      |721      |731
CTCCTTAACAACATGGAACTCATTAGAAAGATCTATAGCACTCTGGCTGGCACCAGGAAA
L  L  N  N  M  E  L  I  R  K  I  Y  S  T  L  A  G  T  R  K
                |231                        |241

      |741      |751      . . . . .
GATGTTGAAGTGACTAAGGgtgagtgagaatatatctgaattctagcctgcaagtattgt
D  V  E  V  T  K  E
                |251

. . . . .
ggtgtattattattaggggaagaaggcaactagtactttattatttttctatcccattg

. . . . .
acacatattaactatatgataacgtgtgtttttcatcctttgatcagacaaacaaaagca

. . . . .
aacaaaaaatcttcaccaagttttaacaaaactgggccagcttggtgactcacacctg

. . . . .
aaatcccagcagtttgggaggccgaggtgggtggatcacctgagatcaggagttcgagac

. . . . .
cagcctggccaagagggtg
```



Exon 8 | Start: 137474 | End: 137567 | Length: 93  
BE AWARE: Flanking intron is shared with the following exon

```
. . . . .
ctgggacctatgcatatgaagtttcatatatgattccttattttctctgattggcaattt

. . . . .
gaataacatcagatgacccaagcttttaaaactgtatattcaagaacatattctgcacac

. . . . .
gctttccaaatggatgcaagaactgcaaaaattcttaccttatttggattgacttgaatt

. . . . .
ctacaactgatgataataataaagttttgtgatgttcactcattccagtgcccttgagtta

. . . . .
atgctgaatttagtggctatttcttcttgaatctttcccttttaacttttcttccataca

      |761      |771      |781      |791      |801      |811
AGGAGTTTGGTCTGGCAGCTCAGAAATTTGGTCAGGTTACACCCATGGAAGTTGACATCT
  E  F  V  L  A  A  Q  K  F  G  Q  V  T  P  M  E  V  D  I  L
                        |261                                |271

      |821      |831      |841      . . . . .
TGTTTCAGTTAGCAGATTTATATGAGCCAAGGGGgtaagttggctttttttttttttttt
  F  Q  L  A  D  L  Y  E  P  R  G
                        |281

. . . . .
ttggaattagcttaataaaaaggaggtaggaggaggcagcaatcaggagaaaaagaagc

. . .
caaactgaaggcta
```

Exon 9 | Start: 137770 | End: 137854 | Length: 84  
 BE AWARE: Flanking intron is shared with the previous exon

```

. . . . .
tactgaaatatgagaaatgaaaaagggatgtttttaaatTTTataatgtaaattgtaat

. . . . . |851 |861
aaattggtatatTTTgttgcttggtgtttgtttttcccctacaACGTATGACCTTAGCAGAC
                                     R M T L A D

|871 |881 |891 |901 |911 |921
ATTGAACGGATTGCTCCTCTGGAAGAGGGAACCTCTGCCCTTTAACTTGGCTGAGGCCAG
I E R I A P L E E G T L P F N L A E A Q
|291 |301

|931 . . . . .
AGGCAGgtgagcaaagaggagcaaacttgcaattctggggtttccttagcaggatacccca
R Q
|311

. . . . .
accctgttccacttgcaagttgcctttgcggcattggtatctgacttaaccatttttctgt

. . . . .
tttgcacaaaaactctcgataaatTTTaaaggctgtaaagagaaattgcttagaaatggt

. . . . .
gtgataaacaggatTTTggcatttgctcatttcttgctaagggtggagtgatttcctaact

. . . . .
gcaccagaagcctaattgctccgaaaacatgaaggaggtcttttctcctttcttgaagtct

. .
catggt

```

Exon 10 | Start: 142137 | End: 142221 | Length: 84

. . . . .  
gtgatgtttttaactgtgcatttttttaatagagaatttgaaaataggcttataacatca

. . . . .  
tttttgtcaaaggatggttttgattatattgaaatttggttttaaacattaatagcacta

. . . . .  
aaatctttttattaataagaacttatgaaatccattcctacttttttccttgacatttta

. . . . .  
gttgatcatggatttagaacccaatgagtaatgtttaaagttaatgtatggtcattttt

. . . . .  
gttttatcttcttttctgtcacctgttttcctgctctctgattcctgtgcatgcaaagca

          |941      |951      |961      |971      |981      |991  
AAGGCCTCAGGTGATTCAGCTCGACCAGTTCTTCTACAAGTTGCAGAGTCGGCCTACAGG  
K  A  S  G  D  S  A  R  P  V  L  L  Q  V  A  E  S  A  Y  R  
                  |321                                  |331

          |1001      |1011      .      .      .      .      .  
TTTGGTCTGGGTTCGTGCTGGAGgttagtcatggatgagagaaaaattttcacttttg  
F  G  L  G  S  V  A  G  A

. . . . .  
ttattcctattctcttttctctgattttccatcttgaaatgttaaaggacaaagtcttct

. . . . .  
ctagtctttcccaatgccagttatcagtttctacctgtgaggtagggaagagtaaaggg

. . . . .  
aactcttttcccatacctgcatgtgatcagggaacttacaagtggcattttcctctatcta

. . . . .  
cagatcaggccttccaaagacaggttactaatgaggttgggctgtagcaatgtgccttaa

. . . . .  
tttaaaatatctatgacatcctttt

Exon 11 | Start: 142713 | End: 142871 | Length: 158

```
. . . . .
tacaagtggcattttcctctatctacagatcaggccttccaaagacaggttactaatgag

. . . . .
gttgggctgtagcaatgtgccttaatttaaaatatctatgacatccttttattaggggat

. . . . .
acttgaatgaaaaccctcaattgtttattaaagatataagaaaggtagtagtctgttttaa

. . . . .
atctgtatttaaaattcaaaacacatttcttttctcagtccttgctccttgactttaactac

. . . . .
cttcttctttcttctgacctatcttactttcttctgagataatgatattgtttaatttta

|1021      |1031      |1041      |1051      |1061      |1071
CTGTTGGAGCCACTGCTGTGTATCCTATCGATCTTGTA AAAACTCGAATGCAGAACCAAC
V G A T A V Y P I D L V K T R M Q N Q R
|341                      |351

|1081      |1091      |1101      |1111      |1121      |1131
GATCAACTGGCTCTTTTGTGGGAGAACTCATGTATA AAAACAGCTTTGACTGTTTTAAGA
S T G S F V G E L M Y K N S F D C F K K
|361                      |371

|1141      |1151      |1161      |1171      . . . . .
AAGTGCTACGCTATGAAGGCTTCTTTGGACTGTATAGAGgttagtgccacatgctcaata
V L R Y E G F F G L Y R G
|381                      |391

. . . . .
cctgttaggtgaaataacactcaaagggttggtttctcatcttagtgacctgacatgaatt

. . . . .
agcaagactgcgttaaaatggaaatacagcagattttcaattatccatgctaatagggagt

. . . . .
aggctggaggcatctagggcttggttaatacaaaacagcccataatctgaaaattatttat

. . . . .
agtgcctctgttgtagtgttttatttcttaaaataataaattatttttataagcatcttccg

. . . . .
tgaatttttgaaaggctcacattgtttgcatccttgaag
```

Exon 12 | Start: 155624 | End: 155676 | Length: 52

```
. . . . .
gaaggcagaaggggaaggagcagactttgggggaataatgagatttcctatatcactaata

. . . . .
ctggttgatttacccaatgtgcaaacgatccagaatgtgagatggaatgtacttgggactg

. . . . .
atgctggtggttagaagttagtacttaaacattccatttcaaataattctcattaagggtaa

. . . . .
tagcagaaaaattgcagcagagattaagcagtttttaagtggtaatcccctctgtagaact

. . . . .
gttgatttggttagttaatgctaatttatttaataatacttctactttttgttgattttta

      |1181      |1191      |1201      |1211      |1221      .
GTCTGTTGCCACAGTTATTGGGAGTTGCCCCAGAGAAGGCCATAAACTTACAgtaagtc
      L  L  P  Q  L  L  G  V  A  P  E  K  A  I  K  L  T
                        |401

. . . . .
ttttgagttggtgcttcttattaagggattttgtacataactcaagaatctagcaggtatt

. . . . .
cttataggaaacagcaggtttgttttctcagaggtgctagctaatttgaaagcaggggaa

. . . . .
ctcaccacacacattttttcattattctccatttagcagtgagacatccaaaaatgtgata

. . . . .
ataaacttgaggacttttggtggttgctgccatcattggtgtcatcacagcagtagcagc

. . . . .
taaattcccttcagtatctaccatattaggcacatgacgtatattacgtttta
```

Exon 13 | Start: 157023 | End: 157103 | Length: 80

```
. . . . .
tattgtcagtggtgtctcccttcttaagctatttctaattgcagaacatatttttggctttc

. . . . .
atztatcttttaacttttctcctgaaacatcagtgatgcatatatatttttcatttttta

. . . . .
aaactattatactattaagagtacttctcaaactatttttaagatctgttaaattgtgcaa

. . . . .
tttatgattactccaatagactgatagaaaaaaatagcatatatattgacagatgcatattt

. . . . .
ataagtgtttaatgactttctttataatttttcatcagcttctataaaatatttttcttcaa

|1231      |1241      |1251      |1261      |1271      |1281
GTGAACGATTTTGTGAGGGATAAAATTTATGCACAAAGATGGTTCGGTCCCACTTGCAGCA
V  N  D  F  V  R  D  K  F  M  H  K  D  G  S  V  P  L  A  A
|411                                |421

|1291      |1301      |1311. . . . .
GAAATTCTTGCTGGAGGCTGCgtaagtaccttttgaagctctcttcattgaaaagacttg
E  I  L  A  G  G  C
|431

. . . . .
tttcacatatatatcactaccatggtcaacaggtgtggactaaggcttctgtttaaccac

. . . . .
agatcctgcacaagggcagagtctactagggcacctcggcttttcctagacaggcgaacc

. . . . .
attccagaggtggaagggtaaatgtttctgaccttactgcctgttaagtcatgcctgtta

. . . . .
aattcgtcattgcttatggtttgttctgcaatggaagtcatttgtggcagaagtgtttgg

. . . . .
cctttgaaaaatcttttgctaa
```

Exon 14 | Start: 180452 | End: 180592 | Length: 140

. . . . .  
gttgtttggtgcttcaaggagcatttgtaatttcagtgtaggacttggggaggtcaacag

. . . . .  
aaatctctggtctttgccctccttcagcctccacattaaggacatagggcttcttcctag

. . . . .  
gcagtaactgtattttgattcatacttgtaatccctaaccgttttgattgatgggacaag

. . . . .  
ggctgaaactacattctcttgaggatagaatatttttatccttgagaataatcattcatc

. . . . .  
tcttcctgaccaccttaccatttctttgctttgatcctaaatcttttttgttcttttta

          |1321      |1331      |1341      |1351      |1361      |1371  
GCTGGAGGCTCCCAGGTGATTTTCACAAATCCTTTAGAAATCGTCAAGATCCGTTTGCAA  
A  G  G  S  Q  V  I  F  T  N  P  L  E  I  V  K  I  R  L  Q  
          |441                          |451

          |1381      |1391      |1401      |1411      |1421      |1431  
GTGGCAGGAGAAATCACCACCTGGTCCTCGAGTCAGTGCTCTGTCTGTCTGCGGGACCTG  
V  A  G  E  I  T  T  G  P  R  V  S  A  L  S  V  V  R  D  L  
          |461                          |471

          |1441      |1451  .      .      .      .      .      .  
GGGTTTTTTGGGATCTACAAGgtaacttttttttttttagtcatttatttgattactt  
G  F  F  G  I  Y  K  
          |481

. . . . .  
tctcatctcctaatagcagaaactattcttcaacatgaaatccatttttttttgcatcttct

. . . . .  
acacctggagaagatgttctaccaagctgcagaggcaccttcatgccaattagaaggag

. . . . .  
gtgccccttcctgccatctgccccaaaaactatcctcatcaatacaciaaatctaggatgac

. . . . .  
tatgataggaatcgtgccctctagagttgttcagtgagctctttgcagcaaacactggct

. . . . .  
agaattttatttgatttggt

Exon 15 | Start: 195267 | End: 195405 | Length: 138

. . . . .  
ttaaaaggatgtcacaggcagagttccaaaaaggtttaattgaagtcttagcattagaa

. . . . .  
taaatagagaactgccctcctcccatcacagcacacaacaaaactagtttttggttttaaaa

. . . . .  
acaaagtttagatttattacttgcagaatttgggttatttttttctgttccccctttcaat

. . . . .  
gtgtgagtacaaaatttggcttctccaggaatattttgttctgaggtaggtccagagaa

. . . . .  
gcttgatctcctcaatgggatatgtgtttcatgtggtttttcttcctttgcaaagtgtga

          |1461      |1471      |1481      |1491      |1501      |1511  
GGTGCCAAAGCATGCTTTCTGCGGGACATTCCTTTCTCGGCCATCTACTTTCCGTGCTAT  
G  A  K  A  C  F      L  R  D  I  P  F  S  A  I  Y  F  P  C  Y  
                  |491                                  |501

          |1521      |1531      |1541      |1551      |1561      |1571  
GCTCATGTGAAGGCTTCCTTTGCAAATGAAGATGGGCAGGTTAGCCCAGGAAGCCTGCTC  
A  H  V  K  A  S      F  A  N  E  D  G  Q  V  S  P  G  S  L  L  
                  |511                                  |521

          |1581      |1591.      .      .      .      .      .  
TTAGCTGGTGCCATAGCTGgtgagtggtcccccattggagttctcttacttccctgaccctt  
L  A  G  A  I  A  G  
                  |531

. . . . .  
gctgcccccttccctagctgcatgctactacattgaaattttttgctgatagttcttg

. . . . .  
acagggtatgtgtgagtttgctagagattgaagagctactgaaaaaaaaaggaaaaaatt

. . . . .  
aatcttcattctggcagatgagattattagacaagcttaatttcttaactctagaaaagg

. . . . .  
ctcattaaatattggcttattttgtggtattgtggaactccaattcaagttccatatcat

. . . . .  
ttcatagtctcacttcaca



gaagtagtggttattgataaacaatttcttaaaaaatatttaaagatttaggaatagaaaca  
aaggaaatgaaggctgagtgatttgggcaaggaaggtgtgagtctggtggggaggggtg  
taggagcagacatcttgcacacaggagaggcatggttgcatttcccatagaggacagcaa  
aataggcattttattgaatcggatttttaactttctttccacggccagcagttcaaagca  
cagttatttttatatagtgagaatgtgaccagactgagatggtgttgtctctcctgca

|1721      |1731      |1741      .      .      .      .  
 GTGAAGAAGGACCAAAAGCTCTGTGGAAGGGAGCTGGTGgtatggaaataatgtgttctt  
 E E G P K A L W K G A G A  
                                  |581

Exon 17 | Start: 205403 | End: 205493 | Length: 90  
 BE AWARE: Flanking intron is shared with the previous exon

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. . . . . |1751 |1761
tttttaaaatatctaattatatctgtgatttctccatttttttaaaCTCGTGTATTTCGA
                                     R V F R

      |1771      |1781      |1791      |1801      |1811      |1821
TCCTCACCCCAGTTTGGTGTAAC TTTGCTGACTTACGAATTGCTACAGCGATGGTTCTAC
S S P Q F G V T L L T Y E L L Q R W F Y
      |591                               |601

      |1831      |1841. . . . .
ATTGATTTTGGAGGAGTgtaagtatcatgctaaatctgctgctaaattttggctgctgct
I D F G G V
      |611

. . . . .
aatgctctgttgtcgtagggaaaggtaacctcagtctagtctctataataaaaatagaagag

. . . . .
atttagggaaatggtgaaattgtgttaacaaggtcatttccaaggggaacatacaactga

. . . . .
tgagaatgtatcaactcctttacattaaatatttgaacaggttttgaaatggtatttttat

. . . . .
caagttaaaacaaacaaaaatacagcttcttactctgttttctcttcttttagaaaacc

. . . . .
catgggatcagagccag

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Exon 18 | Start: 205771 | End: 206928 | Length: 1157  
BE AWARE: Flanking intron is shared with the previous exon

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. . . . .
atttccaaggggaacatacaactgatgagaatgtatcaactcctttacattaaatatttg

. . . . .
aacagggttttgaaatgttatttttatcaagttaaaacaaacaaaaatacagcttcttact

. . . . . |1851 |1861 |1871 |1881
ctgttttctcttcttttaAAAACCCATGGGATCAGAGCCAGTTCCTAAATCCAGGATCAA
          K P M G S E P V P K S R I N
                      |621

|1891 |1901 |1911 |1921 |1931 |1941
CCTGCCTGCCCCGAATCCTGATCACGTTGGGGGCTACAAACTGGCAGTTGCTACATTTGC
  L P A P N P D H V G G Y K L A V A T F A
      |631                      |641

|1951 |1961 |1971 |1981 |1991 |2001
AGGGATTGAAAACAAATTTGGACTTTACCTACCTCTCTTCAAGCCATCAGTATCTACCTC
  G I E N K F G L Y L P L F K P S V S T S
      |651                      |661

|2011 |2021 |*1 |*11 |*21 |*31
AAAGGCTATTGGTGGAGGCCCATAGGAAGATCAGCCCTGGGATAGTGCTGTCTTTTTGTG
  K A I G G G P *
      |671

|*41 |*51 |*61 |*71 |*81 |*91
GGTACTGCAGTAAAGAACATCCCTCCTGGGAATGAAGCAATGCTTCATCCCTTTTACGTC

|*101 |*111 |*121 |*131 |*141 |*151
CATCTCTTGTTTAAATTCAAGTCCAGGCTTTTTTATCATGTGAAATCATTCATTTTCTGG

|*161 |*171 |*181 |*191 |*201 |*211
GTGTTTTCTTAACCAGATCATTGTGAAATTATTCATAATTATTATTTGGCCCTCTGCCCA

|*221 |*231 |*241 |*251 |*261 |*271
GAAACCTTTGTTTGCACTGAAAATTGATGGGATTTGGTCAACACTAACATGATTGGGG

|*281 |*291 |*301 |*311 |*321 |*331
AAAGGAGCAAGTCAGAATAGAAATTAGTACTCCCCTCCTTGAAGTAGGATTGTAGTCCCA
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*341	*351	*361	*371	*381	*391
AAGAGGCTACTGTAAGGCAATCATGGTGCTCAGAGCAGTGTTCGTGTGTGTTTAAACT					
*401	*411	*421	*431	*441	*451
GGTAGGAACTAGGTGCATATTTATAAAAAATAAAAAACACTGGGAGAAATGAAAAATAT					
*461	*471	*481	*491	*501	*511
ATATCAAATATATTCAGCCTGGCTTCAAATTGTAAGCATGCACAAATTCTGTCTCTGGAT					
*521	*531	*541	*551	*561	*571
TATATTATGAAGCTTTTATGTGAAACATGTTTCTTTGTAATGAAAACCACATTGGAGATG					
*581	*591	*601	*611	*621	*631
TTTAGTAATCATATTGTTACTGGTACCAAGACTACTAGGGAAATGCCTTTGTACTTTAGG					
*641	*651	*661	*671	*681	*691
GAAGTACTTTTGGCATTTTACTGTACAGACAGAAAAAACTGAGATGTAGCCCCTCTCCTG					
*701	*711	*721	*731	*741	*751
GAAGTGCTAATTTTGAAAACTGCTCATATGATGTACATGTACTGATTACTGCCTATTTT					
*761	*771	*781	*791	*801	*811
AATAAACACTCTTGAAAAATGCATGTTGCCCTGTTGCTGCCTGCCCTATTCTCCTCATCT					
*821	*831	*841	*851	*861	*871
CCCCATCATTTGGTACCCACTTGCTTTTAAAAATCCACTTTATCTTGAATAATGTAAGACAA					
*881	*891	*901	*911	*921	*931
ATATGTTCTGACATAAGTATTTAATTCATGTTGCCTTGCATAATGGTCAGAGGCGCATGA					
*941	*951	*961	*971.	.	.
ATTTGTGAAGGTGGAAATAAACTATTTGTAAAGTGAttgagttgtttggtcttgatg					
.	.	.	.	.	.
ttgcatggtgagtggtacagtaattgtcattattggcctacatgttttgagggg					
.	.	.	.	.	.
aatatttattcatattggcatggatatacctcagtatctttcaaggtaatgtgttcctc					
.	.	.	.	.	.
atattcatggttcaaagtgagtacaagaaactgtatttaaccatagataagcaaagaact					
.	.	.	.	.	.
agcaaactgacaccttagttctagtcattccagcactttcggcagtttattgtgatgatg					
.	.	.	.	.	.

aaggtcacatagttttagtagaacttagccatgaaa

GBK Parser: Version: 1.1, Version Date: 11/02/2015  
Reader: Version: 1, Version Date: 11/02/2015  
Writer: Version: 1, Version Date: 11/02/2015  
Control: Version: 1, Version Date: 11/02/2015