

Gene: SLC25A13 - Sequence: NG\_012247.1  
 Transcript: NM\_014251.2 - Protein: NP\_055066.1  
 Date : February 26, 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

. . . . .
gtgacgggcttgacacagggccaagttacctctgcccacgccgagaggccgctgctg

. . . . .
gacagcgcagcggggtcacgtgtccctgtccagccactgccccaggatctgagccctccc

. . . . .
cggctgcagggcagaggccgctggatcacgcgcaccccgggcgggccaaggctggggccagc

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

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

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

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

. . . . .
tgccggtatcagcctgggggtgcgagaggctcgggcgctcacgtgtctgggttggtgccac

. . . . .
aggcggagcaaatgggggtgccgggcagccgttccaccttgccgctgctggaggaggagg

```

. . . . .  
gccactcctcccgggcagcacaggcggtcggtgcagagaccgaggcccgacctcgagcg  
. . . . .  
gaaactttgtgggccggacactcccacttggccggggagggaagttggtgcctgacgggg  
. . . . .  
ctgagccagggccgacgaggaggagc

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

```
. . . . .
aaaatgacaggtgtttgcaatgtttcaccttttcttttaagaaatgatatacattcaac
. . . . .
attagatattaagagattttgctactgggttccttgcataaaatcttatcattttagaagc
. . . . .
tgcataaggagtgtaatttaaaatgggaaggggcatagtttatgcactggggcaacatg
. . . . .
gtatttaagtacaaaaatgaaatgcactaatttttaaatctgagatttatcttcttagt
. . . . .
cggctagcttgatttctcagctttgttgtaataacataaacatcttttatgtttattata
      |21      |31      |41      |51      |61      .
GTGGCTTTAACCAAGAGAGCAGATCCAGCTGAGCTTAGAACAATATTTTGAAGgtcagt
V  A  L  T  K  R  A  D  P  A  E  L  R  T  I  F  L  K
      |11                      |21
. . . . .
ataaaggaatctattttgtttctgtttgatttgtgttctataacttttaaatattattt
. . . . .
tcaaacagctctttatttcgttggttatattttgaaaaagtcctaaagtgtagcccgga
. . . . .
cgggtggctcatgcctgtaatccagcactttgggaggctgaggtgggtggatcccctgag
. . . . .
gtcaggagtttgagaccagcctggccaacatggtgaaacccatctttactgaaagtacc
. . . . .
aaaactagccaggtgtggtggcacgtgcctgtaatcctggctattcaggaggct
```

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

```
. . . . .
ctggcctctacctgctggagccagtagcacccttcctcacatgttatgacaatcttaatt

. . . . .
gccatcagatattgcccaagtgtccctgtgggaaaacattgctcccagttaagttataaa

. . . . .
cataaaacttgacttggtttttctttgaactgttgggagataatggtcattatgaagcgt

. . . . .
acatgatcatatctgatattttatcgtttgatattttcaactttttcacatatcactct

. . . . .
catgattatatactgttgaaaatcaaataataaaaatactgtttttctttttttaaaca

|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     . . . . .
GTGGTGGATCAGACCAAAGATGGgtatgtttatttttaattattttatttaattcctttt
V V D Q T K D G
                        |71

. . . . .
gtttgaaacagctttcttttgctctaataccaggaagtcattaaatgtataccttcca

. . . . .
tctcttaggaccattttctgtacttactactattattattacattatttgagtaataatg

. . . . .
tagttccttcaatcattgtttcaagttggggattcaatatgtatatatttgactctttta

. . . . .
aagaatgctttatcgatatcagagaactattattgtcaacaaggtggtgatgtgcagatgt

. . . . .
atgagagtatttatatgcctttg
```

Exon 4 | Start: 92231 | End: 92346 | Length: 115

. . . . .  
ccaaaaacatgcctgaaggaagactgaaacatatatgatgctggataaaacagttatatt

. . . . .  
tgaaattataatgtatttagtgtgtaacttcatgggtcaatgctgtcatgtccttttagatt

. . . . .  
ctgtgcattattatttcagtcactatatgtaaaagtaaagctaaatgtttataaatgt

. . . . .  
tttcagtttgaattttatgttttggcagtgatatcttattttttccctcagtgatatgttt

. . . . .  
gtatatgcttctgttttccacaagctactgacttttgctttatgtttgttttatgttgca

221	231	241	251	261	271
ATTAATATCTTTTCAAGAATTTGTTGCCTTTGAATCTGTCCTGTGTGCCCTGATGCTTT					
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
GTTTATGGTAGCCTTTCAGCTGTTTGACAAAGCTGGCAAAGGAGAAGTAACTTTTGtaa				
F	M	V	A	F
Q	L	F	D	K
A	G	K	G	E
V	T	F	E	
		101		

. . . . .  
gaagcatgcttctgtgcacgtctgtgtaagtgtggacttgtgtgagcattttcttttttt

. . . . .  
tttttctaggaacaaaataaagtgtatggaaattctttatagaagatagtatatcccttt

. . . . .  
tagtaaagatacttcaatgttttggagctccagttcaaacatcttcctattcgaaaaaa

. . . . .  
caagtttctggcttagtagttttgatactctgatttagttttaaggtagaggagatgtgg

. . . . .  
catgactgttttcatagagtcatagcatgtgagagccagacaggacttctgtttct

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

```
. . . . .
caaatgagatgataatagtttttagctttctacttgggtttcctacagcctttaggtgtg

. . . . .
tgtgtgtgagagacttcttgattttccccctttattttatttgggtgggattcttgacc

. . . . .
ttgatttcttaactgattccttgagggttcttttatttctgccatagtttgtaatctgta

. . . . .
tgtctattttaaattgaattacttttttaataagaaaaagcgatatccattcaataaca

. . . . .
gattttatcagaaagaacttcatttactttgttttaataaacctgtcttttctcttttca

|331      |341      |351      |361      |371      |381
AGGATGTTAAGCAAGTTTTTGGACAGACCACAATTCATCAACATATTCCATTTAACTGGG
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
ATTCAGAATTTGTGCAACTACATTTTGAAAAAGAAAGAAAAAGACACCTGACATATGCGG
S E F V Q L H F G K E R K R H L T Y A E
|131                                |141

|451      |461
AATTTACTCAGTTTTTATTGgtgggtctagctcttaaaaagttaaaagggtatggccggg
F T Q F L L
|151

. . . . .
cacggtggctcacacctataatcctagcactttgggaggccgaggtgggcggatgacgag

. . . . .
gtcaggagatcgagaccatcctggccaacatggtgaaacccgtctctactaaaaataca

. . . . .
aaaaattagccaggcgtggtggccggcgcctgtggtcccagctacttgggaggctgaggc

. . . . .
aggagaatggtgtgaacttgggaggtggagcttgagtgagctgagatcgaccactgca

. . . . .
gtccggcctgggtgaaagag
```

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

```
. . . . .
cacttcattccggttttcattttctttgtgttcattgactctagagagcacggaagtaat

. . . . .
tgggtttcattgggtagcagagtaaggtagagtgcactttctcaagacatgttcattggg

. . . . .
ctgagcaaagtctgtctcttggaaggagacagtagcttggtgggtgctattaggcagatg

. . . . .
agggcttgtagatcaagatgtaaactctctttcttcagtctgattttaaatgaactgt

. . . . .
catttgtagtaattattaaatgagaagcataaattaagtattttactttctttatttgta

|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
AGAGTCACAGCCATCGACTTCCGAGACATCATGGTCACCATCCGCCCCATGTCTTGACT
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      . . . . .
CCTTTTGTAGAAGAATGTCTAGTAGCTgtaagttgtaacttgccctaataagtggttta
P F V E E C L V A
|201

. . . . .
actcagtaaataaggaattataaaaacaattcttataagttatgtagtgttttttctg

. . . . .
ttgcaaacactaacattttaatgtgatcacatacattactaaactcaacaaagagtgtg

. . . . .
ttttggctagtttttagtgggtaggttaatttggtgtctgggtaatacgggaatatcttga

. . . . .
gaaatacattagctttttaagataaaaggaaaaacagatttgatctcatgttaattt

. . . . .
tgtttttagattttgaagtatatcaagg
```

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

```
. . . . .
catttccaattgaaatttgaactctaaattactactagtccctgttataatcctacacatg

. . . . .
tatttatattagttttctaggagtgtgtatggagagtacaagttctggcatgattagtt

. . . . .
gcagttgcttctaaaaatcatttatagcttttttttaatgcaaagaaaaataacttatag

. . . . .
tcatttcttctgtacatgcatttttgattactttcaaactagtttttagatgtatgtact

. . . . .
tctgaatcatctataaatcttttgtatatatttttcttctctctgttttcaatttatttga

      |621      |631      |641      |651      |661      |671
GCTGCTGGAGGTACCACATCCCATCAAGTTAGTTTCTCCTATTTAATGGATTAAATTCG
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
CTCCTTAACAACATGGAACCTATTAGAAAGATCTATAGCACTCTGGCTGGCACCAGGAAA
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

. . . . .
aacaaaacaatcttcaccaagttttaacaaaactgggcccagcttggtgactcacacctg

. . . . .
aatcccagcagtttgggaggccgaggtgggtggatcacctgagatcaggagttcgagac

. . . . .
cagcctggccaagagggtg
```



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

```

. . . . .
ctgggacctatgcatatgaagtttcatatatgattccttattttctctgattggcaattt
. . . . .
gaataacatcagatgaccccaagcttttaaactgtatattcaagaacatattctgcacac
. . . . .
gctttccaaatggatgcaagaactgcaaaaattcttaccttatttgattgacttgaatt
. . . . .
ctacaactgatgataataataaagttttgtgatgttcactcattccagtccttgagtta
. . . . .
atgctgaatttagtggctatttcttcttgaatctttcccttttaacttttcttccataca

      |761      |771      |781      |791      |801      |811
AGGAGTTTGTCTGGCAGCTCAGAAATTTGGTCAGGTTACACCCATGGAAGTTGACATCT
  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

. . . . .
ttggaattagcttaataaaaaggaggttaggaggagggcagcaatcaggagaaaaagaagc

. . . .
caaactgaaggcta

```

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

```

. . . . .
tactgaaatatgagaaatgaaaaagggatgtttttaattttataatgtaaattgtaat

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

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

|931 . . . . .
AGGCAGgtgagcaaagaggagcaaacttgcattctggggtttccttagcaggatacccca
R Q
|311

. . . . .
accctgttccacttgacgttgcctttgcggcattgggtatctgacttaaccatttttctgt

. . . . .
tttgcacaaaactctcgataaattttaaggctgtaaagagaaattgcttagaaatggt

. . . . .
gtgataaacaggtatttggcatttgctcatttcttgctaagggtggagtgatttcctaact

. . . . .
gcaccagaagcctaatagtccgaaaacatgaaggaggtcttttctcctttcttgaagtct

. . .
catggt

```

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

```
. . . . .
gtgatgtttttaactgtgcatttttttaatgagaatttgaaaataggcttataacatca
. . . . .
tttttgtcaaaggatggttttgattatattgaaatttggttttaacattaatagcacta
. . . . .
aaatctttttattaataagaacttatgaaatccattcctactttttccttgacatttta
. . . . .
gttgatcatggatttagaacccaatgagtaatgtttaagttaatgtatggtcattttt
. . . . .
gttttatcttcttttctgtcacctgttttcctgctctctgattcctgtgcatgcaaagca
      |941      |951      |961      |971      |981      |991
AAGGCCTCAGGTGATTGAGCTCGACCAGTTCTTCTACAAGTTGCAGAGTCGGCCTACAGG
K A S G D S A R P V L L Q V A E S A Y R
                        |321                        |331

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

. . . . .
ttattcctattctcttttctctgattttccatcttgaaatgttaaaggacaaagtcttct
. . . . .
ctagtctttcccaatgccagttatcagtttctacctgtgaggtagggaagagtaaaggg
. . . . .
aactcttttcccatacctgcatgtgatcagggacttacaagtggcattttcctctatcta
. . . . .
cagatcaggccttccaaagacaggttactaatgaggttgggctgtagcaatgtgccttaa
. . . . .
tttaaaatatctatgacatcctttt
```

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

. . . . .  
tacaagtggcattttcctctatctacagatcaggccttccaaagacaggttactaatgag  
. . . . .  
gttgggctgtagcaatgtgccttaatttaaaatatctatgacatccttttattaggggat  
. . . . .  
acttgaatgaaaacctcaattgtttattaaagatataagaaaggtagtactgttttaa  
. . . . .  
atctgtatttaaaattcaaaacacatttcttttctcagtcctgtgccttgactttaactac  
. . . . .  
cttcttcttcttctgacctatcttactttcttctgagataatgatattgtttaatttta  
  
|1021 |1031 |1041 |1051 |1061 |1071  
CTGTTGGAGCCACTGCTGTGTATCCTATCGATCTTGTA AAACTCGAATGCAGAACCAAC  
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  
  
. . . . .  
cctgttaggtgaaataacactcaaagggttggtttctcatcttagtgcctgacatgaatt  
. . . . .  
agcaagactgcgttaaaatggaaatacagcagattttcaattatccatgctaattgggagt  
. . . . .  
aggctggaggcatctagggttggttaatacaaacagcccataatctgaaaattatttat  
. . . . .  
agtgcctgtgtgtatgttttatttcttaaaataataaattattttataagcatcttcg  
. . . . .  
tgaatttttgaaaggctcacattgtttgcatccttgaag

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

. . . . .  
gaaggcagaagggaaggagcagactttgggggaataatgagatttcctatatcactaata

. . . . .  
ctgttgatttacccaatgtgcaaacgatccagaatgtgagatggaatgtacttgggactg

. . . . .  
atgcggtgttagaagttagtacttaaacattccatttcaaataattctcattaagggtaa

. . . . .  
tagcagaaaattgcagcagagattaagcagtttttaagtggtaatcccctctgtagaact

. . . . .  
gttgatttgtttagttaatgctaatttatttaataatacttctactttttgttgattttta

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

. . . . .  
ttttgagttggtgcttcttattaagggtttgtacataactcaagaatctagcaggtatt

. . . . .  
cttataggaaacagcaggtttgttttctcagaggtgctagctaatttgaaagcaggggaa

. . . . .  
ctcaccacacacattttttcattattctccatttagcagtgacatccaaaaatgtgata

. . . . .  
ataaacttgaggacttttggtggtgctgccatcattgttgcatcacagcagtagcagc

. . . . .  
taaattcccttcagtatctaccatattaggcacatgacgtatattacgtttta

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

```
. . . . .
tattgtcagtggtgtctcccttcttaagctatttctaattgcagaacatatttttggtttc

. . . . .
atztatcttttaacttttctcctgaaacatcagtgatgcataatatatttttcatttttta

. . . . .
aaactattatactattaagagtacttctcaaactattttaagatctgttaaattgtgcaa

. . . . .
tttatgattactccaatagactgatagaaaaaatagcatatatattgacagatgcataatt

. . . . .
ataagtgtttaatgactttctttatatatttttcacagcttctataaatatttttcttcaa

|1231      |1241      |1251      |1261      |1271      |1281
GTGAACGATTTTGTGAGGGATAAATTTATGCACAAAGATGGTTCGGTCCCACTTGACAGCA
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

. . . . .
cctttgaaaaatctttgctaa
```

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

. . . . .  
gttgtttgtgcttcaaggagcatttgaatttcagtgtgaggacttggggaggtcaacag

. . . . .  
aaatctctggtccttgcctccttcagcctccacattaaggacatagggttcttcctag

. . . . .  
gcagtaactgtatcttgattcatacttgtaatccctaaccgttttgattgatgggacaag

. . . . .  
ggctgaaactacattctcttgaggatagaatatcttgccttgagaataatcattcatc

. . . . .  
tcttcctgaccaccttaccatttcttggcttgcctaaatctttttgttcttttta

          |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  
GTGGCAGGAGAAATCACCACCTGGTCCTCGAGTCAGTGCTCTGTCTGTCGTGCGGGACCTG  
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

. . . . .  
tctcatctcctaatagcagaaactattcttcaacatgaaatccattttttttgcatctt

. . . . .  
acacctggagaagatgttctacccaagctgcagaggcaccttcatgccaattagaaggag

. . . . .  
gtgccccttctgccatctgccaaaaactatcctcatcaatacacaaatctaggatgac

. . . . .  
tatgataggaatcgtgccctctagagttgttcagtgcagctctttgcagcaacactggct

. . . . .  
agaattttatttgatttggtt

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

. . . . .  
ttaaaaggatgtcacaggcagagttccaaaaagggtttaattgaagtcttagcattagaa

. . . . .  
taaattgagaactgccctcctcccatcacagcacacaacaaaactagtttttgttttaaaa

. . . . .  
acaaagtttagatttattacttgacagaatttgggttatttttttctgttcccctttcaat

. . . . .  
gtgtgagtacaaatttggcttctccaggaatattttgttctgaggtgagttccagagaa

. . . . .  
gcttgatctcctcaatgggatatgtgtttcatgtggtttttcttcctttgcaaatgtgca

          |1461      |1471      |1481      |1491      |1501      |1511  
GGTGCCAAAGCATGCTTTCTGCGGGACATTCTTTCTCGGCCATCTACTTTCCGTGCTAT  
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.      .      .      .      .      .  
TTAGCTGGTGCCATAGCTGgtgagtggtcccccattggagttctcttacttccttgaccctt  
L  A  G  A  I  A  G  
                  |531

. . . . .  
gctgcccccttccctagctgcatgtactacattgaaattttttgctgatagttcttg

. . . . .  
acagggtatgtgtgagtttgctagagattgaagagctactgaaaaaaagggaataatt

. . . . .  
aatcttcattctggcagatgagattattagacaagcttaatttcttaactctagaaaagg

. . . . .  
ctcattaaatattggcttattttgtggtattgtggaactccaattcaagttccatatcat

. . . . .  
ttcatagtctcacttcaca



Exon 16 | Start: 205151 | End: 205309 | Length: 158  
 BE AWARE: Flanking intron is shared with the following exon

```

. . . . .
gaagtagtggttattgataaacaatttcttaaaaaatatttaaagatttaggaatagaaaca

. . . . .
aaggaaatgaaggctgagtgatttgggcaagggaaggtgtgagtcctggtggggaggggtg

. . . . .
taggagcagacatcttgcacacaggagaggcatggttgcatctccatagaggacagcaa

. . . . .
aataggcattttattgaatcggatttttaactttctttccacggccagcagttcaaagca

. . . . .
cagttatttttatatagtgagaatgtgaccagactgagatgggtgtgtgtctctcctgca

      |1601      |1611      |1621      |1631      |1641      |1651
GTATGCCTGCAGCATCTTTAGTGACCCCTGCTGATGTTATCAAGACGAGATTACAGGTGG
  M P A A S L V T P A D V I K T R L Q V A
                        |541                                |551

      |1661      |1671      |1681      |1691      |1701      |1711
CTGCCCCGGGCTGGCCAAACCACTTACAGCGGAGTGATAGACTGCTTTAGAAAGATACTGC
  A R A G Q T T Y S G V I D C F R K I L R
                        |561                                |571

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

. . . . .
aactaactctttggtatcaggtaaa

```

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

```

. . . . .|1751      |1761
tttttaaaatatctaattatatctgtgatttctccatttttttaaaCTCGTGATTTCGA
                                     R V F R

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

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

. . . . .
aatgctctgttgctgtagggaaaggtaacctcagtctagtctctataataaaatagaagag

. . . . .
atttagggaaatggtgaaattgtgttaacaaggtcatttccaaggggaacatacaactga

. . . . .
tgagaatgtatcaactcctttacattaaatatttgaacaggttttgaatgttatttttat

. . . . .
caagttaaacaacaaaaatacagcttcttactctgttttctcttcttttagaaaacc

. . . . .
catgggatcagagccag

```

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

```

. . . . .
atttccaaggggaacatacaactgatgagaatgtatcaactcctttacattaaatatttg

. . . . .
aacaggttttgaatgttatttttatcaagttaaaacaaacaaaaatacagcttcttact

. . . . . |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
CCTGCCTGCCCCGAATCCTGATCAGTTGGGGGCTACAACTGGCAGTTGCTACATTTGC
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 P *
          |671

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

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

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

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

          |*281 |*291 |*301 |*311 |*321 |*331
AAAGGAGCAAGTCAGAATAGAAATTAGTACTCCCCTCCTTGAAGTAGGATTGTAGTCCCA

          |*341 |*351 |*361 |*371 |*381 |*391
AAGAGGCTACTGTAAGCAATCATGGTGCTCAGAGCAGTGTTCGTGTGTGTTTTAACT

```

*401	*411	*421	*431	*441	*451
GGTAGGAAACTAGGTGCATATTTATAAAAAATAAAAAACACTGGGAGAAATGAAAAATAT					
*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
GAAGTACTTTTGGCATTTTACTGTACAGACAGAAAAAAGTGAAGATGTAGCCCCTCTCCTG					
*701	*711	*721	*731	*741	*751
GAAGTGCTAATTTTGAAAAAGTGTCTCATATGATGTACATGTACTGATTACTGCCTATTTT					
*761	*771	*781	*791	*801	*811
AATAAACACTCTTGAAAAATGCATGTTGCCCTGTTGCTGCCTGCCCTATTCTCCTCATCT					
*821	*831	*841	*851	*861	*871
CCCCATCATTGGTACCACTTGCTTTTAAATCCACTTTATCTTGAATAATGTAAGACAA					
*881	*891	*901	*911	*921	*931
ATATGTTCTGACATAAGTATTTAATTCATGTTGCCTTGCATAATGGTCAGAGGCGCATGA					
*941	*951	*961	*971.	.	.
ATTTGTGAAGGTGGAATAAACTATTTGTAAAGTGAttgagttgtttgggtcttgtgcatg					
.	.	.	.	.	.
ttgcatgggtgagtggtgatacagtaattgtcattattggcctacatgttttgtttgagggg					
.	.	.	.	.	.
aattttattcatattggcatggatatatcctcagtatctttcaaggtaatgtgttcctc					
.	.	.	.	.	.
attttcatgggtcaaagtgagtacaagaaactgtatttaaccatagataagcaaagaact					
.	.	.	.	.	.
agcaaactgacaccttagttctagtcattccagcactttcggcagtttattgtgatgatg					
.	.	.	.	.	.
aaggtcacatagtttttagtagaacttagccatgaaa					

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