

Gene: ENSG00000198457 - Sequence: ENST00000448314
 Transcript: ENST00000448314 - Protein: ENSP00000398594
 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: 501 | End: 824 | Length: 323
 BE AWARE: Flanking intron is shared with the following exon

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. . . . .
ttgttcttcaggcgattcaggaaggcctattaggcagggactgccattttctctctgaga

. . . . .
caaacatcatgccagtaaaactggcccacggtggggtggcagagggagagggccaggtgg

. . . . .
gggcggacactattgcctgcacagttgatgtggaaccagaaagctgactctggatgcagg

. . . . .
aaaaaggtcagggttgcatttcccttccttgcttcttgatgggtgatcaatTTTTTTgaa

. . . . .
atacggacgtcccaaggccaatgagactggtgtcattccagaaaagggccactctgtggg

      |-119      |-109      |-99      |-89      |-79      |-69
GGGTCGGTGGGAGGGTACCTGAAGGTGGGGTCAAGGGAGGCCCAAAACAGTCTACACAG

      |-59      |-49      |-39      |-29      |-19      |-9
CAGGAGGGATGGCTGGGGCTCTTGAGCTATAAGTGGCACCTCAGGGCCCTGACGGGCGTC

      |1      |11      |21      |31      |41      |51
TCGCCATGCTGCTCCTGGGCCTGCTGCTGCTGCCCTGCTGGCTGGCGCCCGCCTGCTGT
      M L L L G L L L L P L L A G A R L L W
      |1      |11

      |61      |71      |81      |91      |101      |111
GGAAGTGGTGAAGCTCCGGAGCCTCCACCTCCCGCCTCTTGCCCCGGGCTTCTTGCACT
      N W W K L R S L H L P P L A P G F L H L
      |21      |31

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121	131	141	151	161	171
TGCTGCAGCCCGACCTCCCAATCTATCTGCTTGGCCTGACTCAGAAATTCGGGGCCATCT					
L	Q	P	D	L	P
I	Y	L	L	G	L
T	Q	K	F	G	P
I	Y				
41			51		

181	191
ACAGGCTCCACCTTGGGCTGCAAGgtgagaggctgatctcgctctggccctcaccatagg							
R	L	H	L	G	L	Q	D
61							

.	.
agggggcgagg	

Exon 2 | Start: 922 | End: 1011 | Length: 89
BE AWARE: Flanking intron is shared with both adjacent exons

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. . . . . |201 |211
tgacggagagggcctctctccgctgacgctgctttggctgtctccaATGTGGTGGTGC
                                         V V V L
                                         |71

      |221      |231      |241      |251      |261      |271
TGA ACTCCAAGAGGACCAT TGAGGAAGCCATGGTCAAAAAGTGGGCAGACTTTGCTGGCA
  N S K R T I E E A M V K K W A D F A G R
                        |81 |91

      |281
GACCTGAGCCACTTACCTgtaagggtggtgggcattttttctttcttaaacaatttttt
  P E P L T Y

. . . . .
tttaagagatgggttcttgctatgttgcccaggctggtcttaaattcctagtctcaaatg

. . . . .
atcctcccacctcagcctcaagtgtgagccacctttgg
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Exon 3 | Start: 1294 | End: 1448 | Length: 154
BE AWARE: Flanking intron is shared with both adjacent exons

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. . . . .
ggcatccccaatccaggtccctggaagctcttgggggggcataatctggtggggagaaaagc

. . . . .
aggggttggggaggccgaagaaggtcaggccctcagctgccttcacagttcccaccctc

. . . . . |291      |301      |311      |321
cagcccccaactcctcctgcaACAAGCTGGTGTCTAAGAACTACCCGGACCTGTCCTTGG
                        K  L  V  S  K  N  Y  P  D  L  S  L  G
                        |101

|331      |341      |351      |361      |371      |381
GAGACTACTCCCTGCTCTGGAAAGCCCACAAGAAGCTCACCCGCTCAGCCCTGCTGCTGG
D  Y  S  L  L  W  K  A  H  K  K  L  T  R  S  A  L  L  L  G
|111                        |121

|391      |401      |411      |421      |431      |441
GCATCCGTGACTCCATGGAGCCAGTGGTGGAGCAGCTGACCCAGGAGTTCTGTGAGgtaa
I  R  D  S  M  E  P  V  V  E  Q  L  T  Q  E  F  C  E
|131                        |141

. . . . .
ggctgggctcctgaggccacctcgggtcagcctcgctctcacagtag
```

Exon 4 | Start: 1556 | End: 1657 | Length: 101
BE AWARE: Flanking intron is shared with both adjacent exons

.
cccccgccctgcccgctgcacagcggcctgctgaactcacactgtttctccacaCGCATG
R M

|451 |461 |471 |481 |491 |501
AGAGCCCAGCCCGGCACCCCTGTGGCCATTGAGGAGGAATTCTCTCTCCTCACCTGCAGC
R A Q P G T P V A I E E E F S L L T C S
|151 |161

|511 |521 |531 |541
ATCATCTGTTACCTCACCTTCGGAGACAAGATCAAGgtgcctcacagcccctcaggccca
I I C Y L T F G D K I K
|171 |181

. . . .
ccccagcccctccctga

Exon 5 | Start: 1746 | End: 1847 | Length: 101
BE AWARE: Flanking intron is shared with both adjacent exons

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. . . . . |551 |561
gcctctccttgtcctgaactgaaagtactccctccttttctggcaGACGACAACTTAATG
                                     D D N L M

      |571      |581      |591      |601      |611      |621
CCTGCCTATTACAAATGTATCCAGGAGGTGTTAAAAACCTGGAGCCACTGGTCCATCCAA
P  A  Y  Y  K  C  I  Q  E  V  L  K  T  W  S  H  W  S  I  Q
      |191                                |201

      |631      |641      . . . . .
ATTGTGGACGTGATTCCCTTCTCAGGgtgaggacctggagcctagacaccctgggttg
I  V  D  V  I  P  F  L  R
      |211

. . . . .
taggggagaggctgggg
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Exon 6 | Start: 1949 | End: 2035 | Length: 86
BE AWARE: Flanking intron is shared with both adjacent exons

. |651
tggagggagagggctccttcccacagctgcattctcatgcttcctgccgcaTTCTTCCCCA
F F P N

|661 |671 |681 |691 |701 |711
ATCCAGGTCTCCGGAGGCTGAAGCAGGCCATAGAGAAGAGGGATCACATCGTGGAGATGC
P G L R R L K Q A I E K R D H I V E M Q
|221 |231

|721 |731
AGCTGAGGCAGCACAAGgtggggactgtacgtggacggcctcccctcggcccacagccag
L R Q H K
|241

.
tgatgctaccggcctcagcattgctatgaggcgggttcttt

Exon 7 | Start: 2205 | End: 2405 | Length: 200
 BE AWARE: Flanking intron is shared with both adjacent exons

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. . . . .
tgcataccccagttatgggcctgttgccactctgtactcctctccccaggccagccgctc

. . . . . |741 |751 |761 |771
agcccgctcctttcaccctctgcaGAGAGCCTCGTGGCAGGCCAGTGGAGGGACATGATG
                        E S L V A G Q W R D M M
                        |251

|781 |791 |801 |811 |821 |831
GACTACATGCTCCAAGGGGTGGCGCAGCCGAGCATGGAAGAGGGCTCTGGACAGCTCCTG
D Y M L Q G V A Q P S M E E G S G Q L L
|261 |271

|841 |851 |861 |871 |881 |891
GAAGGGCACGTGCACATGGCTGCAGTGGACCTCCTGATCGGTGGCACTGAGACCACAGCA
E G H V H M A A V D L L I G G T E T T A
|281 |291

|901 |911 |921 |931 . . .
AACACCCCTCTCCTGGGCCGTGGTTTTTTTGCTTCACCACCCTGAGgtgcgtcctggggac
N T L S W A V V F L L H H P E
|301 |311

. . . . .
aagcaaaaggctccttcccagcaacctggccagggcggtgggcaccctcactcagctctg

. . . .
agcactgtgcggctggggctgtg

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Exon 8 | Start: 2606 | End: 2784 | Length: 178
 BE AWARE: Flanking intron is shared with both adjacent exons

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. . . . .
cttgcctcaccggcactcaggctcactgggttgctgagggagcggctggaggctgggcag

. . . . . |941 |951
ctgtgggctgctggggcaggactccacccgatcattccccaATTGAGCAGCGACTGCAGG
                                I Q Q R L Q E

|961 |971 |981 |991 |1001 |1011
AGGAGCTAGACCACGAACTGGGCCCTGGTGCCTCCAGCTCCCGGGTCCCCTACAAGGACC
E L D H E L G P G A S S S R V P Y K D R
|321 |331

|1021 |1031 |1041 |1051 |1061 |1071
GTGCACGGCTGCCCTTGCTCAATGCCACCATCGCCGAGGTGCTGCGCCTGCGGCCCCGTTG
A R L P L L N A T I A E V L R L R P V V
|341 |351

|1081 |1091 |1101 |1111 . . . .
TGCCCTTAGCCTTGCCCCACCGCACCACACGGCCCAGCAGgtgactcccgagggttgggg
P L A L P H R T T R P S S
|361 |371

. . . .
atgagtgaggaaagcccag

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Exon 9 | Start: 2868 | End: 2971 | Length: 103
 BE AWARE: Flanking intron is shared with the previous exon

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. . . . . |1121 |1131
cccaggagggtcctggccagcctctaactccagcccccttcaCATCTCCGGCTACGACAT
                                     I S G Y D I

      |1141      |1151      |1161      |1171      |1181      |1191
CCCTGAGGGGCACAGTCATCATTCCGAACCTCCAAGGCGCCACCTGGATGAGACGGTCTG
P E G T V I I P N L Q G A H L D E T V W
      |381                                |391

      |1201      |1211      . . . . .
GGAGAGGCCACATGAGTTCTGGCCTGgtatgtggggggccgggggcctgccgtgaaaatg
E R P H E F W P D
      |401

. . . . .
tgggtggaggctggtccccgctgccgctgaacgcctccccacccacctgtccacccgcccg

. . . . .
cagatcgcttcctggagccaggcaagaactccagagctctggccttcggctgcggtgccc

. . . . .
gcgtgtgcctgggcgagccgctggcgcgctggagctcttcgtggtgctgacccgactgc

. . . . .
tgcaggccttcacgctgctgccctccggggacgccctgccctccctgcagcccctgcccc

. . . . .
actgcagtgtcatcctcaagatgcag

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Exon 10 | Start: 3069 | End: 3838 | Length: 769
 BE AWARE: Flanking intron is shared with the previous exon

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. . . . . |1221 |1231
ccccgctgccgctgaacgcctccccacccacctgtccacccgcccgcaATCGCTTCCTGG
                                         R F L E
                                         |411

      |1241      |1251      |1261      |1271      |1281      |1291
AGCCAGGCAAGAACTCCAGAGCTCTGGCCTTCGGCTGCGGTGCCCCGCGTGTGCCTGGGCG
  P G K N S R A L A F G C G A R V C L G E
                        |421                        |431

      |1301      |1311      |1321      |1331      |1341      |1351
AGCCGCTGGCGCGCCTGGAGCTCTTCGTGGTGCTGACCCGACTGCTGCAGGCCTTCACGC
  P L A R L E L F V V L T R L L Q A F T L
                        |441                        |451

      |1361      |1371      |1381      |1391      |1401      |1411
TGCTGCCCTCCGGGGACGCCCTGCCCTCCCTGCAGCCCCCTGCCCCACTGCAGTGTTCATCC
  L P S G D A L P S L Q P L P H C S V I L
                        |461                        |471

      |1421      |1431      |1441      |1451      |1461      |1471
TCAAGATGCAGCCTTTCCAAGTGCGGCTGCAGCCCCGGGGGATGGGGGCCACAGCCCAG
  K M Q P F Q V R L Q P R G M G A H S P G
                        |481                        |491

      |1481|*1      |*11      |*21      |*31      |*41
GCCAGAACCAGTGATGGGGCAGGACCGATGCCAGCCGGGTACCTCAGTTTCTCCTTTATT
  Q N Q *

      |*51      |*61      |*71      |*81      |*91      |*101
GCTCCCGTACGAACCCCTCCCCTCCCCCTGTAAACACAGTGCTGCGAGATCGCTGGCAG

      |*111      |*121      |*131      |*141      |*151      |*161
AGAAGGCTTCCTCCAGCGGCTGGGTGGTGAAGGACCCTGGCTCTTCTCTCGGGGCGACCC

      |*171      |*181      |*191      |*201      |*211      |*221
CTCAGTGCTCGGCAGTCATACTGGGGTGCGAGAGAGGTGGGCAGCAGCTCAGCCTCCCCC

      |*231      |*241      |*251      |*261      |*271      |*281
CGCTGGGGAGCGAAAGTTTCTTGGTCTCAGCTTCATTTCCGTGAAGGGCACCGAGAACTC

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|*291 |*301 |*311 |*321 |*331 |*341
 GAAGCCCTTCCAGTGGTACCAGCTCACTCCCTGGGAAAGGGGTTGTCAAGAGAGAGTCAA

 |*351 |*361 |*371 |*381 |*391 |*401
 AGCCGGATGTCCCATCTGCTCTTCCCGTTCCCTTAAGGAGGTAGCTCCCAGCACTCAAC

 |*411 |*421 |*431 |*441 |*451 |*461
 CAACCTCCCCGAGAGCTCCCTTCCTGACCCTCCGCTGCAGAGGATTGAGGCTTAATTCT

 |*471 |*481 |*491 |*501
 GAGCTGGCCCTTTCCAGCCAATAAATCAACTCCAGCTCcctctgcgaggctggcatgatt

 gttccatttcacccagccgctcagtccttgcctgttacctgtggggctgaaacctagg

 caggccgagccccagccaccccagctctgagccgcctccccacccctcacctgatggtcc

 actgtgctcccgtagagcccgttgaggttggcgtagtggcagttcctgtaccaccaggcc

 cctcggtaggagacagcgcaggagatgagcaagctgttgggggtcccgatcacgggcagag

 aagacactgccgctgtggtagctcatggagtcccctgg

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