

Gene: COL1A1 - Sequence: NG\_007400.1  
Transcript: NM\_000088.3 - Protein: NP\_000079.2  
Date : February 17, 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: 5229 | Length: 228

```

. . . . .
agcagaggggcacccctaccactggttagcccacgccattctgaggaccagctgcacc

. . . . .
cctaccacagcacctctggcccaggctgggctggggggctggggaggcagagctgcgaag

. . . . .
aggggagatgtggggtggactcccttcctcctcctccccctctcattccaactccaa

. . . . .
attgggggcccggccaggcagctctgattggctggggcacgggcggcggctccccctct

. . . . .
ccgaggggaggggttcctccctgctctccatcaggacagtataaaaggggcccggccag

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

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

      |1      |11      |21      |31      |41      |51
CTAGACATGTTTCAGCTTTGTGGACCTCCGGCTCCTGCTCCTCTTAGCGGCCACCGCCCTC
      M F S F V D L R L L L L L A A T A L
      |1      |11

      |61      |71      |81      |91      |101      .
CTGACGCACGGCCAAGAGGAAGCCAAGTCGAGGGCCAAGACGAAGACAgtaagtcccaa
L T H G Q E E G Q V E G Q D E D I
      |21      |31

```

. . . . .  
 acttttgggagtgcaaggatactctatatcgcgccttgcgcttggtcccgggggcccgcgg  
 . . . . .  
 cttaaaacgagacgtggatgatccggagactcgggaatggaaggagatgatgagggctc  
 . . . . .  
 ttcctcggcgccctgagacaggaggagctcacctggggcgaggttggggttgaacgcg  
 . . . . .  
 ccccgggagcgggaggtgagggtggagcgcggcgtgagttggtgcaagagagaatccga  
 . . . . .  
 gcgcgcaaccggggaagtggggatctgggtgcagagtgaggaaagcacg

Exon 2 | Start: 6693 | End: 6887 | Length: 194  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 tttttatttcacttggcttcagcgccaatgggctgaggttggagttggaagccaccgcgg  
 . . . . .  
 actaaagctttgtttaattcctgagaactggaaagagttacagcctccctggccaggcg  
 . . . . .  
 cctcggcgtgtcacccgcgctgatgaggagcaggcgagcttttaaggatttgaggaaag  
 . . . . .  
 aagaacggggggaggggcggggaagtgaaaaatccaagtgtgcctcttagacccgggggaa  
 . . . . .  
 aggtggttaagctgggggttgagtcactactgacaacgcccctcttccgcctgtcccag

111	121	131	141	151	161
TCCACCAATCACCTGCGTACAGAACGGCCTCAGGTACCATGACCGAGACGTGTGGA AAC					
P	P	I	T	C	V
		Q	N	G	L
		R	Y	H	D
		R	D	V	W
		K	P		
	41			51	
171	181	191	201	211	221
CCGAGCCCTGCCGGATCTGCGTCTGCGACAACGGCAAGGTGTTGTGCGATGACGTGATCT					
E	P	C	R	I	C
		V	C	D	N
		G	K	V	L
		C	D	D	V
		I	C		
	61			71	

	231	241	251	261	271	281
GTGACGAGACCAAGAACTGCCCCGGCGCCGAAGTCCCCGAGGGCGAGTGCTGTCCCGTCT						
D E T K N C P G A E V P E G E C C P V C						
		81			91	

291	.	.	.	.	.	.	.	.	.
CCCCGACGGCTCAGgtgcggctgcgctcggggcctggggcctggggcctggggcctggg									
P D G S E									

```

. . . . .
gcctggggctggggctgggggtggtcggcgctcgctggccctccgtgctggaggcctctg
. . . . .
ccgacgggagcagcattagcaaaccttggtctaacgcgctctcttcgtcccctagagt
. . . . .
caccacccaccaagaaccaccggcgctcgaggtaatctcctgccctcgaattttgcccc
. . . . .
tgcgcgcccgctgactcctcacagtcctcccttcttaacctggcctctgtttcttctc
. . . . .
ccccaatcccacagg

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Exon 3 | Start: 7050 | End: 7084 | Length: 34  
BE AWARE: This section overlaps with the following exon

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. . . . .
aaccgagccctgccgatctgcgtctgcgacaacggcaaggtggtgtgcgatgacgtga
. . . . .
tctgtgacgagaccaagaactgccccggcgccgaagtccccgagggcgagtgctgtcccg
. . . . .
tctgccccgacggctcaggtgcggctgcgctcggggcctggggcctggggcctggggcct
. . . . .
ggggcctggggctggggctgggggtggtcggcgctcgctggccctccgtgctggaggcct
. . . . .

```

ctgccgacgggagcagcattagcaaaccttggctctaacgcgcgtctcttcgtcccctag

      |301      |311      |321      |331      .      .      .      .  
AGTCACCCACCGACCAAGAAACCACCGGCGTCGAGgtaatctcctgccctcgaattttgc  
      S   P   T   D   Q   E   T   T   G   V   E  
      |101  |111

      .      .      .      .      .      .      .      .      .  
ccctgcgcggcccgtgactcctcacagtcctcccttctctaacctggcctcttgtttctt

      .      .      .      .      .      .      .      .      .  
ctcccccaatcccacagggacccaaggagacactggccccgaggccaagggttaagcg

      .      .      .      .      .      .      .      .      .  
ttgactctgggctgtggggggctgcaggtgggcatggctctcgccccacgctcacccc

      .      .      .      .      .      .      .      .      .  
ggccccgcctctccccctgcagggacccgcaggccccctggccgagatggcatccctg

      .      .      .      .      .      .  
gacagcctggacttcccggacccccggaccccc

Exon 4 | Start: 7187 | End: 7222 | Length: 35  
BE AWARE: This section overlaps with the following exon

      .      .      .      .      .      .      .      .      .  
ggtgcggctgcgctcggggcctggggcctggggcctggggcctggggcctggggcctgggg

      .      .      .      .      .      .      .      .      .  
ctgggggtggtcggcgctcgctggccctccgtgctggaggcctctgccgacgggagcagc

      .      .      .      .      .      .      .      .      .  
attagcaaaccttggctctaacgcgcgtctcttcgtcccctagagtcacccaccgaccaa

      .      .      .      .      .      .      .      .      .  
gaaaccaccggcgtcgaggtaatctcctgccctcgaattttgccctgcgcggcccgtga

      .      .      .      .      .      .      .      .      .  
ctcctcacagtcctcccttctctaacctggcctcttgtttcttctccccaatcccacag

              |341      |351      |361      .      .      .  
GGACCCAAGGGAGACACTGGCCCCCGAGGCCCAAGGgtaagcgttgactctgggctgtg

G P K G D T G P R G P R  
|121

. . . . .  
gggggctgcaggtgggcatggctctcggtcccgctcaccggtcccgccctctcccc  
. . . . .  
ctgcagggacccgcaggccccctggccgagatggcatccctggacagcctggacttccc  
. . . . .  
ggaccccccgacccccggacctcccgacccccctggcctcgaggagtaagtggagag  
. . . . .  
gccttggtgtccactctcccctgttttgttttgtttttggcagatgacataatttta  
. . . . .  
tactttgaaataatttcaaacttacagaaaagttgc

Exon 5 | Start: 7313 | End: 7414 | Length: 101

. . . . .  
aaaccttggctctaacgcgctcttctcggtcccctagagtcaccacccaccaagaacc  
. . . . .  
accggcgtcgaggtaatctcctgccctcgaattttgcccctgcgcggcccgtagctcctc  
. . . . .  
acagtcctcccttcttaacctggcctcttgtttcttctccccaatcccacagggaccc  
. . . . .  
aaggagacactggcccccgaggccaagggttaagcgttgactctgggctgtggggggc  
. . . . .  
tgcaggtgggcatggctctcggtcccgctcaccggtcccgccctctccccctgcag

|371 |381 |391 |401 |411 |421  
GGACCCGCAGGCCCCCTGGCCGAGATGGCATCCCTGGACAGCCTGGACTTCCCGGACCC  
G P A G P P G R D G I P G Q P G L P G P  
|131 |141

|431 |441 |451 |461 |471 . . .  
CCCGGACCCCCGACCTCCCGACCCCCTGGCCTCGGAGGAgtaagtggagaggccttg

P G P P G P P G P P G L G G  
|151

. . . . .  
tgtgtccactctcccctgttttggttttggttttggttcagatgacataatttatacttt  
. . . . .  
gaaataatttcaaacttacagaaaagttgcaagaatcctacaggaaactctcacataccc  
. . . . .  
ttcacagtttgtagcatgtgctttattagtctctgtttatgtatatgtatctttttttt  
. . . . .  
ctgaactgtttgagcaagttgctaacatcaggctcttttgcgctaaataacttaggtgtg  
. . . . .  
tttttcctaaaaacaagagcattctcttaactgacctacaca

Exon 6 | Start: 8136 | End: 8207 | Length: 71  
BE AWARE: This section overlaps with the following exon

. . . . .  
tgcgtttaatcatcatgtctctagttccacaaatgtagaacgttcctcagacttttcttt  
. . . . .  
gtcttttagtggcactgggagttttgatgagtcagttgttttgagactgtccctcaatt  
. . . . .  
tgggattgtctcattagattagatgcagggatgcatctttgcaggaatgtcttaaaagca  
. . . . .  
atgttattcttctcagcacatcacaccaggaagtgcagtgatgtcagtttcttccatcctc  
. . . . .  
agtgccgtcttctgcctttcaattcactgtcctcactctgacttctctgtttgttctag

	481	491	501	511	521	531
	AACTTTGCTCCCCAGCTGTCTTATGGCTATGATGAGAAATCAACCGGAGGAATTTCCGTG					
	N F A P Q L S Y G Y D E K S T G G I S V					
	161			171		
	541	.	.	.	.	.

CCTGCCCCATGgtgagccagcagggggagcatggatgacagaagagagaatgggtatcc  
P G P M  
|181

. . . . .  
agaggaggtgggcataggcggctggtatagacagcttgggaggtccagatcacctttggg  
. . . . .  
acctcagagtccagaaaggatgcaggacgactgggtggtcccaacaggcatgaatgacta  
. . . . .  
catccacatgctttcctacagagggatcacatgacccccctttcttctccctctatagg  
. . . . .  
gtccctctggtcctcgtggtctccctggccccctggtgcacctgtgagtatccaggacg  
. . . . .  
tcttcatatgcc

Exon 7 | Start: 8435 | End: 8479 | Length: 44  
BE AWARE: This section overlaps with the following exon

. . . . .  
gaactttgctccccagctgtcttatggctatgatgagaaatcaaccggaggaatttcggt  
. . . . .  
gcctggccccatggtgagccagcagggggagcatggatgacagaagagagaatgggtatc  
. . . . .  
cagaggaggtgggcataggcggctggtatagacagcttgggaggtccagatcacctttgg  
. . . . .  
gacctcagagtccagaaaggatgcaggacgactgggtggtcccaacaggcatgaatgact  
. . . . .  
acatccacatgctttcctacagagggatcacatgacccccctttcttctccctctatag  
|551 |561 |571 |581 . . .  
GGTCCCTCTGGTCCTCGTGGTCTCCCTGGCCCCCTGGTGACCTgtgagtatccaggac  
G P S G P R G L P G P P G A P  
|191

. . . . .  
 gtcttcatatgcctccttgggctttggtcttttggagggaagactgggatgagggcagga  
 . . . . .  
 gagatgctcagagatctcttggttaagattggagaagggtgacagggacttgtcttctaac  
 . . . . .  
 ccatctttttccttcttctcaagggtccccaaggcttccaagggtcccctggtgagcctg  
 . . . . .  
 gcgagcctggagcttcagtaagcactctctatacagattcatactccttctacaaacaca  
 . . . . .  
 cagactctcctatagaagaactcccaggcctggggtcttccttac

Exon 8 | Start: 8638 | End: 8691 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 caggacgactgggtggtcccaacaggcatgaatgactacatccacatgctttcctacaga  
 . . . . .  
 gggatcacatgacccccctttcttctccctctatagggtccctctggtcctcgtggtct  
 . . . . .  
 ccctggccccctggtgcacctgtgagtatccaggacgtcttcatatgcctccttgggct  
 . . . . .  
 ttggtcttttggagggaagactgggatgagggcaggagagatgctcagagatctcttggt  
 . . . . .  
 aagattggagaagggtgacagggacttgtcttctaaccatcttttcttcttctcaag  
 |591 |601 |611 |621 |631 |641 .  
 GGTCCCCAAGGCTTCCAAGGTCCCCTGGTGAGCCTGGCGAGCCTGGAGCTTCAgtaagc  
 G P Q G F Q G P P G E P G E P G A S  
 |201 |211  
 . . . . .  
 actctctatacagattcatactccttctacaaacacacagactctcctatagaagaactc  
 . . . . .



ccaggcctggggtcttccttacctcttccttcaatcccagccttccccttctttttttc  
 . . . . .  
 ttatccatattctaaccacctcttctatcttttctaggggtcccatgggtccccgagggtcc  
 . . . . .  
 ccagggtccccctggaagaatggagatgatgtaagtatcccagcaagaagataccatc  
 . . . . .  
 tgaccccatggcctccatgggttgggtcctgcaatttcactccaccacatttg

Exon 9 | Start: 8855 | End: 8908 | Length: 53

. . . . .  
 gagatgctcagagatctcttggttaagattggagaagggtgacagggacttgtcttctaac  
 . . . . .  
 ccatctttttccttcttctcaagggtccccaaggcttccaagggtccccctgggtgagcctg  
 . . . . .  
 gcgagcctggagcttcagtaagcactctctatacagattcatactccttctacaaacaca  
 . . . . .  
 cagactctcctatagaagaactcccaggcctggggtcttccttacctcttccttcaatc  
 . . . . .  
 ccagccttccccttcttttttctatccatattctaaccacctcttctatcttttctag  
 |651 |661 |671 |681 |691 .  
 GGTCCCATGGGTCCCCGAGGTCCCCAGGTCCCCCTGGAAGAATGGAGATGATgtaagt  
 G P M G P R G P P G P P G K N G D D  
 |221 |231  
 . . . . .  
 atccccagcaagaagataccatctgaccccatggcctccatgggttgggtcctgcaattt  
 . . . . .  
 ccactccaccacatttggaacgatactcagaggaaggagggaagtcctctctgatgca  
 . . . . .  
 cggactgacctggaacaatgatcttttcgcttagtgagatgattccatgtccccaacaaa

. . . . .  
gtgactgttctcctcaccacccagccaccttagagcaatccccaaccccatccctttgggga

. . . . .  
aattggtgcgagatggtgaaattaaaatgctggtgacagaagtagacagaaat

Exon 10 | Start: 9407 | End: 9460 | Length: 53  
BE AWARE: This section overlaps with the following exon

. . . . .  
ctcaccacccagccaccttagagcaatccccaaccccatccctttggggaaattggtgcgca

. . . . .  
gatggtgaaattaaaatgctggtgacagaagtagacagaaattcctttagaggcactcag

. . . . .  
atttcaccaaacgaaggtttcactgtagatttaaactgagctctagattcaaagataaga

. . . . .  
ttctgggcccccaaactgacctgcaacaatcaaagaagactgagaccttctccacttt

. . . . .  
tccagcccctaggcgtggtggggaggcagaggcatgatggtcttttctctccctctcag

          |701      |711      |721      |731      |741      .  
GGGGAAGCTGGAAAACCTGGTCGTCTGGTGAGCGTGGGCCTCCTGGGCCTCAGgtgagc  
G E A G K P G R P G E R G P P G P Q  
                          |241

. . . . .  
agggggctgtggctgaacctgggcttcactgcacttgggcttcatttaggagctgggtcc

. . . . .  
acagtgatgtgttctaattggcccttccttgtcttcttcatctctctccagggtgctcgag

. . . . .  
gattgcccggaacagctggcctccctggaatgaaggacacagagtgagtcacctttgag

. . . . .  
tcatttaagctccccaagtccctagcatacccccacccatccagtcccaacctcttccccaaa

. . . . .

gatactgagttgcatcatggtgggtggcagctacagaagtcccaaggacagag

Exon 11 | Start: 9577 | End: 9630 | Length: 53

```
. . . . .
aaagataagattctgggccccaaacctgacctgcaacaatccaaagaagactgagacct
. . . . .
tctccacttttccagcccctaggcgggtggtggggaggcagaggcatgatggtcttttctc
. . . . .
tccctctcagggggaagctggaaaacctggctcgtcctggtgagcgtgggcctcctgggcc
. . . . .
tcagggtgagcagggggctgtggctgaacctgggccttcactgcacttgggccttcatttagg
. . . . .
agctgggtccacagtgatgtgttctaattggcccttccttgtcttcttcattctctccag
|751      |761      |771      |781      |791      |801      .
GGTGCTCGAGGATTGCCCGAACAGCTGGCCTCCCTGGAATGAAGGGACACAGAgtgagt
G A R G L P G T A G L P G M K G H R
|251                                |261
. . . . .
cacctttgagtcatttaagctcccaagtcacctagcatacccccattccagtcaccaacctc
. . . . .
ttccccaaaagatactgagttgcatcatggtgggtggcagctacagaagtcccaaggac
. . . . .
agagagtggacatccaaagcacctccctccatgggaaagcagtcctgattaaacgattg
. . . . .
ggtgagatctagagccagttgggggttagtctagctcagaaacaaagggatggcggtgat
. . . . .
gacctcccaaggctctttctcagatctaggtggatgtcaaggctgttccacccc
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Exon 12 | Start: 9970 | End: 10023 | Length: 53

BE AWARE: This section overlaps with the following exon

```
. . . . .
agcatacccccatccagtcaccaacctcttccccaaaagatactgagttgcatcatggtgg

. . . . .
gtggcagctacagaagtcccaaggacagagagtggacatccaaaagcacctccctccat

. . . . .
gggaaagcagtcctcgattaaacgattgggtgagatctagagccagttggggtttagtcta

. . . . .
gctcagaaacaaggatggcggatgacctccaaggctctttctcagatctaggtgg

. . . . .
atgtcaaggctgttccacccctccacaggttcttaccttctacctctttcctgctttag

      |811      |821      |831      |841      |851      .
GGTTTCAGTGGTTTGGATGGTGCCAAGGGAGATGCTGGTCCTGCTGGTCCTAAGgtaaga
G F S G L D G A K G D A G P A G P K
      |271                        |281

. . . . .
ggctgtctgaacatcatggtcctccacatcccagagtcaccacatgaatgaatttctca

. . . . .
ctcattatctctgatctacagggtgagcctggcagccctggtgaaaatggagctcctgg

. . . . .
tcagatggtgagtggtgccagttccagagggcagggatggggcaggaggcaggggcaaga

. . . . .
tggaggcctgggggaacaaggctgtctcccatctcatctgacttctcttggtttggtgt

. . . . .
cagggcccccgtggcctgcctggtgagagaggtcgccctggagcccctggccct
```

Exon 13 | Start: 10112 | End: 10156 | Length: 44

BE AWARE: This section overlaps with the following exon

. . . . .  
 gattgggtgagatctagagccagttgggggttagtctagctcagaaacaagggatggcg  
 . . . . .  
 gtgatgacctcccaaggctctttctcagatctaggtggatgtcaaggctgttccaccccc  
 . . . . .  
 tccacaggttcttaccttctacctctttcctgctttagggtttcagtggtttgatggtg  
 . . . . .  
 ccaaggagatgctggtcctgctggtcctaaggtaaggagctgtctgaacatcatgggcc  
 . . . . .  
 tccacatccccagagtcccaccatgaatgaatttctcactcattattctctgatctacag  
 . . . . .  
 |861 |871 |881 |891 |901 . . .  
 GGTGAGCCTGGCAGCCCTGGTGAAAATGGAGCTCCTGGTCAGATGgtgagtgtgccagtg  
 G E P G S P G E N G A P G Q M  
 |291 |301  
 . . . . .  
 tccagagggcagggatggggcaggaggcaggggcaagatggaggcctgggggaacaaggc  
 . . . . .  
 tgtctcccatctcatctgacttctcttggtttggttgtcaggggccccgtggcctgcctg  
 . . . . .  
 gtgagagaggtcgccctggagcccctggccctgctgtaagtactcctggccccttggggg  
 . . . . .  
 atccctgagctctggaaggggctccccaggaactctagggactggccagtgctcagtgga  
 . . . . .  
 cttaacggggcttcccctctctcctgcagggtgctcgtggaaatg

Exon 14 | Start: 10273 | End: 10326 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ttcagtggtttgatggtgccaaggagatgctggtcctgctggtcctaaggtaaggaggc  
 . . . . .

tgtctgaacatcatggtcctccacatccccagagtcaccacatgaatgaatttctcactc  
 . . . . .  
 attattctctgatctacagggtagcctggcagccctgggtgaaaatggagctcctgggtca  
 . . . . .  
 gatggtgagtggtgccagttccagagggcagggatggggcaggaggcaggggcaagatgg  
 . . . . .  
 aggcctgggggaacaaggctgtctcccatctcatctgacttctcttggtttggtgtcag  
 . . . . .  
 |911 |921 |931 |941 |951 .  
 GGGCCCCGTGGCCTGCCTGGTGAGAGAGGTGCGCCCTGGAGCCCCCTGGCCCTGCTgtaagt  
 G P R G L P G E R G R P G A P G P A  
 |311  
 . . . . .  
 actcctggccccttgggggatccctgagctctggaaggggctcccaggaactctagga  
 . . . . .  
 ctggccagtgtcagtggaacttaacggggcttcccctctctcctgcaggggtgtctgtga  
 . . . . .  
 aatgatggtgtactggtgtgtccgggccccctgtgagtggtggcctgtaggcctcagggc  
 . . . . .  
 ctgggagtgaggagggtctcagtatctgtcttggggctgacaatgggagcaggttatg  
 . . . . .  
 ttggtctgaacccaggaacttctctgtcccagggtgtgacttgagctgccat

Exon 15 | Start: 10441 | End: 10485 | Length: 44  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 agctcctggtcagatggtgagtggtgccagttccagagggcagggatggggcaggaggca  
 . . . . .  
 ggggcaagatggaggcctgggggaacaaggctgtctcccatctcatctgacttctcttg  
 . . . . .  
 tttggtgtcagggccccctggcctgcctgggtgagagaggtcgccctggagcccctggc



```

. . . . .
agggtgtgacttgacgtgccatctcttccttctcgctgacatctccatttcattcacag

      |1011      |1021      |1031      |1041      |1051      .
GGTCCACCGGCCCCGCTGGTCCTCGCTTCCTGGTGCTGTTGGTGCTAAGgtgaga
G P T G P A G P P G F P G A V G A K
      |341                      |351

. . . . .
ccccccactctcctctaagcatgaccctcatgggccaaggggttcattgtctccctgttcc

. . . . .
ccaaaccaaagggaccagagtggcaagagagcagcccgttcactaacacctttgtcctg

. . . . .
ggttctccgtctctgatcttagagtcctgatcattgctctcctgtccctgtctccccttc

. . . . .
ctcctgccatcccagaggcaaggttgggtttccagggtggcttctgatatgtcctttc

. . . . .
ttctgattcaggggtgaagctggtccccaagggccccgaggctctgaaggtcccc

```

Exon 17 | Start: 10975 | End: 11073 | Length: 98  
BE AWARE: This section overlaps with the following exon

```

. . . . .
ccccgctggtcctcctggcttccctgggtgctgttgggtgctaaggtgagacccccactct

. . . . .
cctctaagcatgaccctcatgggccaaggggttcattgtctccctgttcccaaaccaaag

. . . . .
ggaccagagtggcaagagagcagcccgttcactaacacctttgtcctgggttctccgtc

. . . . .
tctgatcttagagtcctgatcattgctctcctgtccctgtctccccttctcctgccatc

. . . . .
ccgagaggcaaggttgggtttccagggtggcttctgatatgtcctttcttctgattcag

      |1061      |1071      |1081      |1091      |1101      |1111

```



GGTGAAGCTGGTCCCCAAGGGCCCCGAGGCTCTGAAGGTCCCCAGGGTGTGCGTGGTGAG  
 G E A G P Q G P R G S E G P Q G V R G E  
 |361 |371

|1121 |1131 |1141 |1151 . . . .  
 CCTGGCCCCCTGGCCCTGCTGGTGCTGGCCCTGCTgtaagtgtccccgactcagtg  
 P G P P G P A G A A G P A  
 |381

. . . . .  
 tccccctttgccactttctaacctcagagtccttgccctgttgctgacactcctttctctgt  
 . . . . .  
 gccacagggaaccctggctgctgatggacagcctggctgctaaagggtccaatgtaagtat  
 . . . . .  
 cctgccaggcttcagtcctcctgcccgcctgcagcctgcctgcccctttccctctgct  
 . . . . .  
 cctaggctcacgcctggctgtctgcctcccacagggtgctcctgggtattgctgggtgctc  
 . . . . .  
 ctggcttcctgggtgcccgaggcccctctggaccccagg

Exon 18 | Start: 11162 | End: 11206 | Length: 44  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ttagagtcctgatcattgctctcctgtccctgtctccccttcctcctgccatcccagagag  
 . . . . .  
 gcaagggtgggtttccagggtggcttctgatatgtcctttcttctgattcagggtgaag  
 . . . . .  
 ctggtccccaagggtcccagggtctgaagggtcccagggtgtgctggtgagcctggcc  
 . . . . .  
 cccctggccctgctgggtgctgctggccctgctgtaagtgtccccgactcagtgctcccctt  
 . . . . .  
 tgccactttctaacctcagagtccttgccctgttgctgacactcctttctctgtgccacag





|441

|451

. . . . .  
ctccccgccatccttcttgagcccagcccaccctgccctaggagccccctgagggaaat  
.  
ccagaaaggaagaggagcccctagtcttctgggggagtcctgccacacccccaggaacc  
.  
cctgacactggaggcccagcctcagccggctctgaggctggcacaggatggcccctcacc  
.  
acaggccgcctcctcctctcgccctctccagggccctgttggtgttcaaggacccccctg  
.  
gcccctgctggagaggaaggaaagcgaggagctcgagggtgaacccggaccactg

Exon 21 | Start: 11812 | End: 11919 | Length: 107  
BE AWARE: This section overlaps with the following exon

. . . . .  
cagctcatcctcttctcccccttgagggtgaacctggtgctcctggcagcaaaggaga  
.  
cactggtgctaaggagagcctgtaagtctccccgccatccttcttgagcccagcccac  
.  
cctgccctaggagccccctgagggaatccagaaaggaagaggagcccctagtcttctgg  
.  
gggagtcctgccacacccccaggaaccctgacactggaggcccagcctcagccggctc  
.  
tgaggctggcacaggatggcccctcaccacaggccgcctcctcctctcgccctctccag

          |1361      |1371      |1381      |1391      |1401      |1411  
GGCCCTGTTGGTGTTC AAGGACCCCTGGCCCTGCTGGAGAGGAAGGAAAGCGAGGAGCT  
G P V G V Q G P P G P A G E E G K R G A  
                          |461                                  |471

          |1421      |1431      |1441      |1451      |1461.      .  
CGAGGTGAACCCGACCCACTGGCCTGCCCGACCCCTGGCGAGCGTgtaagtgtccct

R G E P G P T G L P G P P G E R  
|481

. . . . .  
gccccccccctcccgtccaccctcattgcctggctggtgcctgtgtgtcgcggagttca  
. . . . .  
ctggcctcctctcctcctgcagggtggacctggtagccgtggtttccctggcgcagatgg  
. . . . .  
tgttgctggtcccaaggtaacctctccttgccggccgggggctgaccctgccgctccctg  
. . . . .  
ggcatcttcttctcttttggcccggtggcaaagagccacaaacttgagaccctaactgtt  
. . . . .  
cctgtgacttcccccaaccagggtcccgtggtgaacgtggttctcct

Exon 22 | Start: 12014 | End: 12067 | Length: 53  
BE AWARE: This section overlaps with the following exon

. . . . .  
ggaaccctgacactggaggcccagcctcagccggctctgaggctggcacaggatggccc  
. . . . .  
ctcaccacaggccgctcctcctctcggccctctccagggccctgttggtgttcaaggac  
. . . . .  
cccctggccctgctggagaggaaggaaagcgaggagctcgaggtgaaccgggaccactg  
. . . . .  
gcctgcccggaccccctggcgagcgtgtaagtgtccctgcccggcccctccgctccacc  
. . . . .  
ctcattgcctggctggtgcctgtgtgtcgcggagttcactggcctcctctcctcctgcag  
|1471 |1481 |1491 |1501 |1511 .  
GGTGGACCTGGTAGCCGTGGTTTCCTGGCGCAGATGGTGTGCTGGTCCCAAGgtaacc  
G G P G S R G F P G A D G V A G P K  
|491 |501  
. . . . .

tctccttgccggccgggggctgaccctgccgctccctgggcatcttcttcctcttttggc  
 . . . . .  
 ccgtggcaaagagccacaaacttgagaccctaactgttcctgtgacttccccaaccagg  
 . . . . .  
 gtcccgtggtgaacgtggttctcctggccctgctggccccaaggatctcctggtgaag  
 . . . . .  
 ctggtcgtcccgtgaagctggtctgcctggtgccaaggtgaggccccaggctttcagcc  
 . . . . .  
 tggcttgccaggccctgaccatcccgtgtagggtctgggatgaggcgttctgg

Exon 23 | Start: 12193 | End: 12291 | Length: 98  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ggccctgcccggaccccctggcgagcgtgtaagtgtccctgcccggcccctcccgtccac  
 . . . . .  
 cctcattgcctggctggtgcctgtgtgtcgcgagttcactggcctcctctcctcctgca  
 . . . . .  
 ggggtggacctggttagccgtggtttccctggcgagatggtgttgctggtcccaaggtaac  
 . . . . .  
 ctctccttgccggccgggggctgaccctgccgctccctgggcatcttcttcctcttttgg  
 . . . . .  
 cccgtggcaaagagccacaaacttgagaccctaactgttcctgtgacttccccaaccag

1521	1531	1541	1551	1561	1571
GGTCCCGCTGGTGAACGTGGTTCTCCTGGCCCTGCTGGCCCCAAAGGATCTCCTGGTGAA					
G P A G E R G S P G P A G P K G S P G E					
	511			521	

  

1581	1591	1601	1611	.	.	.	.
GCTGGTCGTCCCGGTGAAGCTGGTCTGCCTGGTGCCAAGgtgaggccccaggctttcagc							
A G R P G E A G L P G A K							
	531						

. . . . .  
 ctggccttgccaggccctgaccatcccgtgtagggctctgggatgaggcgttctggatcag  
 . . . . .  
 gcccaagggtctgccctctggagtcctccccacctccatcatgcttctccccaagtccc  
 . . . . .  
 actcatacctctctgcctccctagggctctgactggaagccctggcagccctggctctgat  
 . . . . .  
 ggcaaaactggccccctgtaagtatcactccccctgaacccctgccattgtcctgtct  
 . . . . .  
 gcctccctgctgtcctcactgctgtttcgtgcctcca

Exon 24 | Start: 12457 | End: 12510 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 gagaccctaactgttcctgtgacttcccccaaccagggtcccgtggtgaacgtggttct  
 . . . . .  
 cctggccctgctggcccaaaggatctcctggtgaagctggctcgtcccgtgaagctggt  
 . . . . .  
 ctgcctggtgccaaggtagggccccaggctttcagcctggcttggccaggccctgaccat  
 . . . . .  
 cccgtgtagggctctgggatgaggcgttctggatcaggccaagggtctgccctctggagt  
 . . . . .  
 cctccccacctccatcatgcttctcccaagtcccactcatacctctctgcctccctag  
 . . . . .  
 |1621 |1631 |1641 |1651 |1661 .  
 GGTCTGACTGGAAGCCCTGGCAGCCCTGGTCCTGATGGCAAACTGGCCCCCTgtaagt  
 G L T G S P G S P G P D G K T G P P  
 |541 |551  
 . . . . .  
 atcactccccctgaacccctgccattgtcctgtctgcctccctgctgtcctcactgctg  
 . . . . .

ctttcgtgcctcccatccttagggtcccgccggtcaagatggtcgccccggacccccagg  
 . . . . .  
 cccacctgggtgccgtgggtcaggctgggtgtgatgggattccctggacctaaggtgctgc  
 . . . . .  
 tgtgagtattaagtgaggatccatgaagagccagggacaaacacacctgagacttgaagg  
 . . . . .  
 agtcctgggctctgggctcagctgtgccgctgacctgccgtgtggccactcact

Exon 25 | Start: 12599 | End: 12697 | Length: 98

. . . . .  
 cccaggctttcagcctggcttggccaggccctgacctcccgtgtagggtctgggatgag  
 . . . . .  
 gcgttctggatcaggcccaagggtctgccctctggagtcctccccacctccatcatgct  
 . . . . .  
 tctccccaagtcccactcatacctctctgcctccctagggtctgactggaagccctggca  
 . . . . .  
 gccctggctcctgatggcaaaactggccccctgtaagtatcactccccctgaacccctg  
 . . . . .  
 ccattgtcctgtctgcctccctgctgtcctcactgctgcttctcgtgcctcccatccttag

1671	1681	1691	1701	1711	1721
GGTCCCGCCGGTCAAGATGGTCGCCCCGGACCCCAGGCCACCTGGTGCCCGTGGTCAG					
G P A G Q D G R P G P P G P P G A R G Q					
	561			571	

1731	1741	1751	1761	.	.	.
GCTGGTGTGATGGGATTCCCTGGACCTAAAGGTGCTGCTGtgagtgatttaagtgaggatcc						
A G V M G F P G P K G A A						
	581					

. . . . .  
 atgaagagccagggacaaacacacctgagacttgaaggagtcctgggctctgggctcagc  
 . . . . .



tgtgccgctgacctgccgtgtggccactcactctcacttttctggacctcagcctccctat  
 . . . . .  
 ctgtaaaatgaaagacttctcggcggggcacggtgggtcatgcctgtaatcccagcactt  
 . . . . .  
 tgggaggccaaggcgggcagaccatgaggtcaggagtttgagaccagtctgggccaacata  
 . . . . .  
 gtgaaaccacgtctctactaaaaatacaaaagattagct

Exon 26 | Start: 13593 | End: 13646 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 cagtcttgggtgattctttgaggtccaaaaggagatagcagagaagatgaaagccctttg  
 . . . . .  
 cagtgcaggccacagcgggcatctaacagggaaaaggcagaggagcctggaagggtcatct  
 . . . . .  
 tgggaggagtggggtcagaaagggccagcaagaagcacctgcaggggcattccccgggg  
 . . . . .  
 gccaaacagtcttttgaaaagaaagtcccttaaaaagtcccactcagagtaaatgagagg  
 . . . . .  
 ccccaggaggccctggcttctcacttcagccccctcaaccctaactccctttctccacag  
 |1771 |1781 |1791 |1801 |1811 |1821.  
 GGAGAGCCCGCAAGGCTGGAGAGCGAGGTGTTCCCGGACCCCTGGCGCTGTCgtaagt  
 G E P G K A G E R G V P G P P G A V  
 |591 |601

. . . . .  
 atctcctttccatccctacctccttccattgctgccccggcactttctcctccctgcag  
 . . . . .  
 gagggtgctagaggccaggtcctcagctgctcggggcctcctaaccctgagttccctt  
 . . . . .  
 ttgctctctccctgcagggtcctgctggcaaatggagaggctggagctcagggacccc

. . . . .  
ctggccctgctgtgagtggtccctgatggggagatctggggagcagaaaaggggagacacc  
. . . . .  
ctcagcccctcgtctcctcggcctccccgtgactgtagtggttctctctgtgcag

Exon 27 | Start: 13790 | End: 13843 | Length: 53  
BE AWARE: This section overlaps with the following exon

. . . . .  
aaagaaagtccttaaaaagtcctcactcagagtaaagagaggccccaggaggccctggc  
. . . . .  
ttctcacttcagccccctcaaccctaactccctttctccacaggagagcccggaaggc  
. . . . .  
tggagagcgagggtgtcccggaacccctggcgctgtcgtaagtatctcctttccatccct  
. . . . .  
acctccttcccattgctgccccggcactttctcctccctgcaggaggggtgctagaggcc  
. . . . .  
acggtcctcagctgctcggggcctcctaaccctgagttcccctttgctctctccctgcag  
. . . . .  
|1831 |1841 |1851 |1861 |1871 .  
GGTCCTGCTGGCAAAGATGGAGAGGCTGGAGCTCAGGGACCCCCTGGCCCTGCTgtgagt  
G P A G K D G E A G A Q G P P G P A  
|611 |621

. . . . .  
gtccctgatggggagatctggggagcagaaaaggggagacaccctcagcccctcgtctcc  
. . . . .  
tcggcctccccgtgactgtagtggttctctctgtgcagggtcccgtggcgagagaggtga  
. . . . .  
acaaggccctgctggctccccggattccaggtaggcctcatggctgtcaggatgctgg  
. . . . .  
gaggtaggggtaggaaacacctctttgggtctcttcagattctaaaccttcctcccttc

. . . . .  
ttcccccatttccacctacaggtctccctggctcctgctggctcctccaggtga

Exon 28 | Start: 13947 | End: 14000 | Length: 53  
BE AWARE: This section overlaps with the following exon

. . . . .  
gtaaagtatctcctttccatccctacctccttcccattgctgccccggcactttctcctcc

. . . . .  
ctgcaggaggggtgctagaggccacggctcctcagctgctcggggcctcctaaccctgagt

. . . . .  
tcccctttgctctctccctgcagggtcctgctggcaaagatggagaggctggagctcagg

. . . . .  
gacccctggccctgctgtgagtgtccctgatggggagatctggggagcagaaaagggga

. . . . .  
gacaccctcagccctcgtctcctcggcctccccgtgactgtagtggttctctctgtgcag

	1881	1891	1901	1911	1921	.
	GGTCCCGCTGGCGAGAGAGGTGAACAAGGCCCTGCTGGCTCCCCCGGATTCCAGgtgagg					
G	P	A	G	E	R	G
			E	Q	G	P
				A	G	S
					P	G
						F
						Q
		631			641	

. . . . .  
cctcatggctgtcaggatgctgggaggtagggtaggaaacacctctttggtctcttcca

. . . . .  
gattctaaaccttccctcccttcttccccatttccacctacaggtctccctggctcct

. . . . .  
gctggctcctccaggtgaagcaggcaaacctgggtgaacaggtgaagaggagcagccggcca

. . . . .  
gaggggtgggagatgcagggaatccagaggacaggccccgctcctagctaatacagaca

. . . . .  
gccatcaactagagggattgaggttagacgccggaagaacttctcccatgaa

Exon 29 | Start: 14112 | End: 14165 | Length: 53

```
. . . . .
aggctggagctcagggaccccctggccctgctgtgagtgctccctgatggggagatctggg
. . . . .
gagcagaaaaggggagacaccctcagcccctcgtctcctcggcctccccgtgactgtagt
. . . . .
gttctctctgtgcagggtcccgctggcgagagaggtgaacaaggccctgctggctcccc
. . . . .
ggattccaggtgaggcctcatggctgtcaggatgctgggaggtaggggtaggaaacacct
. . . . .
ctttgtctcttcagattctaaaccttcctcccttcttccccatttccacctacag
|1931      |1941      |1951      |1961      |1971      |1981      .
GGTCTCCCTGGTCCTGCTGGTCCTCCAGGTGAAGCAGGCAAACCTGGTGAACAGgtaaga
G  L  P  G  P  A  G  P  P  G  E  A  G  K  P  G  E  Q
                        |651                        |661
. . . . .
gggagcagccggccagaggggtgggagatgcagggaatccagaggacaggcccccgctc
. . . . .
ctagctaatacagacagccatcaactagagggattgaggttagacgccgaaagaacttcc
. . . . .
tcccatgaaggagcagcacagaggaagtgggggctgcatgattgctagtctgggtgac
. . . . .
ttcttttaagagctgctggaatatgctgtgactttccctcaacccttctattgataaatc
. . . . .
ttggtccatagtttggggagggggaagcctttgacacatccctaggaggaaga
```

Exon 30 | Start: 14616 | End: 14660 | Length: 44  
BE AWARE: This section overlaps with the following exon

. . . . .  
 ggaagtgggggctgcatgattgctagctctgggtgacttcttttaagagctgctggaatat  
 . . . . .  
 gctgtgactttccctcaacccttctattgataaatcttgggtccatagtttggggaggggg  
 . . . . .  
 gaagcctttgacacatccctaggaggaagagaggggctgtttgggataatctcaattcag  
 . . . . .  
 tgctgagaaggggttcctctctaatacacggccagacccaggaggaaggaccgtgctttc  
 . . . . .  
 cagcagagtggccccaggtgggttttgctcactgtctgttcctctctccctccccctcag  
 . . . . .  
 |1991 |2001 |2011 |2021 . . .  
 GGTGTTCTGGAGACCTTGGCGCCCTGGCCCTCTGGAGCAAGAgtaagtaggcctctc  
 G V P G D L G A P G P S G A R  
 |671  
 . . . . .  
 tcgctgcatccgtcaaggtgcgtcgtacttggccctatctccagagcagccttcacatgc  
 . . . . .  
 cctgtccttcccttctagggcgagagaggtttccctggcgagcgtggtgtgcaaggtccc  
 . . . . .  
 cctggctcctgctgggtccccgaggggccaacggtgctcccggaacgatggtgctaaggtg  
 . . . . .  
 agggcagcgtggaaggggctctggcaagtggcccagggaccaggtctcaccctcctgca  
 . . . . .  
 gcaggggatggcgggcatgaccaaagccatggagatagggtgtg

Exon 31 | Start: 14754 | End: 14852 | Length: 98  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ctaggaggaagagaggggctgtttgggataatctcaattcagtgctgagaaggggttcct

. . . . .  
 ctctaatacagggccagaccccaggaggaaggaccgtgctttccagcagagtggccccagg  
 . . . . .  
 tgggttttgctcactgtctgttcctctctccctccccctcaggggtgttcctggagacctt  
 . . . . .  
 ggcgcccctggcccctctggagcaagagtaagtaggcctctctcgctgcatccgtcaagg  
 . . . . .  
 tgcgtcgtacttggccctatctccagagcagccttcacatgccctgtccttcccttctag  
 |2031 |2041 |2051 |2061 |2071 |2081  
 GCGAGAGAGGTTTCCCTGGCGAGCGTGCTGTGCAAGTCCCCCTGGTCCTGCTGGTCCC  
 G E R G F P G E R G V Q G P P G P A G P  
 |681 |691  
 |2091 |2101 |2111 |2121 . . . . .  
 CGAGGGGCCAACGGTGCTCCCGGCAACGATGGTGCTAAGgtgagggcagcgtggaagggg  
 R G A N G A P G N D G A K  
 |701  
 . . . . .  
 ctctggcaagtggcccagggaccaggtctcaccctcctgcagcaggggatggcggggcca  
 . . . . .  
 tgaccaaagccatggagatagggtgtggggtggggggaaaagaccagggcagggggccac  
 . . . . .  
 acacagcctggagtctgggctgtgagtcttttcatcttttctcaaggctgtcgttggcc  
 . . . . .  
 ttggaacaagcctgggagataccaagcggggcttagggctgtgaccactcttggggcc  
 . . . . .  
 ccaggcctcactccagtcttcttggtgtcacatagggt

Exon 32 | Start: 15150 | End: 15257 | Length: 107

. . . . .  
 aaggtgagggcagcgtggaaggggctctggcaagtggcccagggaccaggtctcaccctt

. . . . .  
 cctgcagcaggggatggcgggcatgaccaaagccatggagatagggtgtgggggtggggg  
 . . . . .  
 gaaaagaccagggcagggggccacacacagcctggagtctgggctgtgagtcttttcac  
 . . . . .  
 ttttctcaaggcttgcgttggccttggaacaagcctgggagataccaagcggggctta  
 . . . . .  
 gggctgtgaccactcttggggcccaggcctcactccagtcttcttgggtgtcacatag  
 |2131 |2141 |2151 |2161 |2171 |2181  
 GGTGATGCTGGTGGCCCTGGAGCTCCCGGTAGCCAGGGCGCCCTGGCCTTCAGGGAATG  
 G D A G A P G A P G S Q G A P G L Q G M  
 |711 |721  
 |2191 |2201 |2211 |2221 |2231 . .  
 CCTGGTGAACGTGGTGCAGCTGGTCTTCCAGGGCCTAAGGGTGACAGAgtaagttcaacc  
 P G E R G A A G L P G P K G D R  
 |731 |741  
 . . . . .  
 ttccccctcccctgagccctacatggctcccctctctgcctgctttgaatctctcagcat  
 . . . . .  
 ctctccttctctctgggatctgtccctcttctcgctaatacctcccttcttcccctttccc  
 . . . . .  
 ctctggcctttttgctgatgaatcctctccctgtgggtccaggcccatctatccccatggg  
 . . . . .  
 ttaccatggtgatgagaggtgggggcatctccttgggtggaggctcccttattcatcccgc  
 . . . . .  
 tacacaagtcaggggcctcttaacctcagttccacctgagtctccagg

Exon 33 | Start: 15716 | End: 15823 | Length: 107  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ctccctgtgggtccaggcccatctatccccatgggttaccatggtgatgagaggtgggggc

. . . . .  
 atctccttgggtggaggctcccttattcatcccgtacacaagtcaggggcctcttaacct  
 . . . . .  
 cagttccacctgagctctccaggcaggcaccctttttcctgaaagaatctttgagtccttg  
 . . . . .  
 gcccaggtggaggcagggcagagctgcagagggcctctcaggaaaccagacacaagcag  
 . . . . .  
 aacactataggtcacctccttgccccacactggaaatctcaagcttatccatgtctttag  
 . . . . .  
 |2241 |2251 |2261 |2271 |2281 |2291  
 GGTGATGCTGGTCCCAAAGGTGCTGATGGCTCTCCTGGCAAAGATGGCGTCCGTGGTCTG  
 G D A G P K G A D G S P G K D G V R G L  
 |751 |761  
 . . . . .  
 |2301 |2311 |2321 |2331 |2341 . .  
 ACTGGCCCCATTGGTCTCCTGGCCCTGCTGGTGGCCCTGGTGACAAGgtgaggtggcgcg  
 T G P I G P P G P A G A P G D K  
 |771 |781  
 . . . . .  
 cctccccaccttctgccctaacacatagcctcctcagcaggcctgggcacggttccgtgg  
 . . . . .  
 ggttgcggtgggagagcaggtcctgccaaactgagctgtcaacctgggaacctggaggga  
 . . . . .  
 ccagaaggaggggaggctctcctgggggtcatctactaggagtattcaggggaggccctga  
 . . . . .  
 ccctgagcctcttgctcccttgctctcaggggtgaaagtgggtcccagcgccctgctggtcc  
 . . . . .  
 cactggagctcgtggtgccccgtaagtacagaagacctgttaagacc

Exon 34 | Start: 16044 | End: 16097 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .



gctctcctggcaaagatggcgtccgtgggtctgactggccccattggtcctcctggccctg  
 . . . . .  
 ctggtgccccctggtgacaaggtgaggtggccgcctcccccaccttctgccctaacacatag  
 . . . . .  
 cctcctcagcaggcctgggcacggttccgtgggggttgcttgggagagcaggtcctgcc  
 . . . . .  
 aactgagctgtcaacctgggaacctggaggaccagaaggaggggaggctctcctgggggt  
 . . . . .  
 catctactaggagtattcaggggaggccctgaccctgagcctcttgtcccttgctctcag  
 . . . . .  
 |2351 |2361 |2371 |2381 |2391 .  
 GGTGAAAGTGGTCCCAGCGGCCCTGCTGGTCCCCTGGAGCTCGTGGTGCCCCGtaagt  
 G E S G P S G P A G P T G A R G A P  
 |791  
 . . . . .  
 acagaagacctgttaagaccccatacttggcccttcctcccttcacacagcaccctgg  
 . . . . .  
 ccctgtctgtgccttcaccccttgctctccctcaccgcatccccgccttcctcctgt  
 . . . . .  
 cagcgcatctctccaatctgactccttttcttctagggagaccgtggtgagcctggtccc  
 . . . . .  
 cccggccctgtggtgttgctggccccctgtgagtaccaagacccccatcattttcat  
 . . . . .  
 caccgactgggacctgggacctcaggggacggaatgaggacaaaggcgtcagcc

Exon 35 | Start: 16260 | End: 16313 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 aaggaggggaggctctcctgggggtcatctactaggagtattcaggggaggccctgacct  
 . . . . .  
 gagcctcttgtcccttgctctcaggggtgaaagtgggtcccagcgccctgctgggtccact

. . . . .  
 ggagctcgtggtgccccgtaagtacagaagacctgttaagacccatacttggcccttc  
 . . . . .  
 cctcccttcacacagcaccctggccctgtctgtgccttcaccccttgccctctccctca  
 . . . . .  
 ccgcatccccgccttccctcctgtcagcgcacatctctccaatctgactccttttcttctag  
 |2401 |2411 |2421 |2431 |2441 |2451.  
 GGAGACCGTGGTGAGCCTGGTCCCCCGGCCCTGCTGGCTTTGCTGGCCCCCCTgtgagt  
 G D R G E P G P P G P A G F A G P P  
 |801 |811  
 . . . . .  
 accaagaccccatcatTTTTcatcaccgactgggacctgggacctcgaggacggaatg  
 . . . . .  
 aggacaaaggcgtcagccatcctcaggggagaagggtggagacgggattgtttccaccc  
 . . . . .  
 aagcatcttctgcctccattactgctcctccccaggtagtggaaactcctgcctcctt  
 . . . . .  
 ccctccatcaccgccctgcttctccccaggtgctgacggccaacctggtgctaaag  
 . . . . .  
 gcgaacctggtgatgctggtgctaaaggcgatgctggtccccctggccctgccg

Exon 36 | Start: 16532 | End: 16639 | Length: 107  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ctctccaatctgactccttttcttctagggagaccgtggtgagcctggtcccccggcc  
 . . . . .  
 tgctggctttgctggccccctgtgagtaccaagaccccatcatTTTTcatcaccgact  
 . . . . .  
 gggacctgggacctcgaggacggaatgaggacaaaggcgtcagccatcctcaggggaga

. . . . .  
 aggggtggagacgggattgtttcccaccaagcatcttcctgcctccattactgctcctcc  
  
 . . . . .  
 cccaggtagtggaactcctgcctccttcctccattaccgcctgcttcctccccag  
  
 |2461 |2471 |2481 |2491 |2501 |2511  
 GGTGCTGACGGCCAACCTGGTGCTAAAGGCGAACCTGGTGATGCTGGTGCTAAAGGCGAT  
 G A D G Q P G A K G E P G D A G A K G D  
 |821 |831  
  
 |2521 |2531 |2541 |2551 . .  
 GCTGGTCCCCCTGGCCCTGCCGACCCGCTGGACCCCTGGCCCCATTgtgagtggttg  
 A G P P G P A G P A G P P G P I  
 |841 |851  
  
 . . . . .  
 gccctctgtgccacgaggctggtgggctgggacccaggacgggtccaggcttgatgcct  
  
 . . . . .  
 ctgtgctctcctacagggtaatgttggtgctcctggagcaaaggctgcgcggcagcgc  
  
 . . . . .  
 tgggtccccctgtgagtatcacccgcctctctgttgagcctctcccctctccccaggcagc  
  
 . . . . .  
 ggtggcaggtgagggcagctgggtcggatgagttggctgttctccctctgactgttccta  
  
 . . . . .  
 tgttctctccttcagggtgctactggtttccctgggtgctgctggccg

Exon 37 | Start: 16728 | End: 16781 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 tgtttcccaccaagcatcttcctgcctccattactgctcctccccaggtagtggaaac  
  
 . . . . .  
 tcctgcctccttcctccattaccgcctgcttcctccccagggtgctgacggccaac  
  
 . . . . .  
 ctggtgctaaaggcgaacctggtgatgctggtgctaaaggcgatgctggtccccctggcc

. . . . .  
 ctgccggaccgcgtggaccccctggccccattgtgagtggttgccctctgtgccacg  
  
 . . . . .  
 aggctggtgggctgggacccaggacgggtccaggcttgatgcctctgtgctctcctacag  
  
 |2561 |2571 |2581 |2591 |2601 |2611 .  
 GGTAATGTTGGTGTCTCTGGAGCCAAAGGTGCTCGCGGCAGCGCTGGTCCCCCTgtgagt  
 G N V G A P G A K G A R G S A G P P  
 |861 |871  
  
 . . . . .  
 atcaccgcctctctgttgagcctctcccctctccccaggcagcgggtggcaggtgagggc  
  
 . . . . .  
 agctgggtcggatgagttggctgttctccctctgactgttcctatgttctctccttcag  
  
 . . . . .  
 ggtgctactggtttccctggtgctgctggccgagtcggtcctcctggcccctctgtaagt  
  
 . . . . .  
 ctctgcagcagagtccactgctctaggttgggggtgctgggtgggggctgccagaaggat  
  
 . . . . .  
 ggtggggctgactgaggacccaatgatgcaccagagccccctggagtctgacag

Exon 38 | Start: 16908 | End: 16961 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ctgccggaccgcgtggaccccctggccccattgtgagtggttgccctctgtgccacg  
  
 . . . . .  
 aggctggtgggctgggacccaggacgggtccaggcttgatgcctctgtgctctcctacag  
  
 . . . . .  
 ggtaatggttgctcctggagccaaaggtgctcgcggcagcgtggtccccctgtgagt  
  
 . . . . .  
 atcaccgcctctctgttgagcctctcccctctccccaggcagcgggtggcaggtgagggc

. . . . .  
agctgggtcggatgagttggctgttctccctctgactgttcctatgttctctccttcag  
|2621 |2631 |2641 |2651 |2661 .  
GGTGCTACTGGTTTCCCTGGTGCTGGCCGAGTCGGTCCTCCTGGCCCCCTCTgtaagt  
G A T G F P G A A G R V G P P G P S  
|881

. . . . .  
ctctgcagcagagtccactgctctaggttgggggtgctgggtgggggctgccagaaggat  
. . . . .  
ggtggggtgactgaggacccaatgatgcaccagagccccctggagtctgacagcccctc  
. . . . .  
ctatcctcatccaggaaatgctggaccccctggccctcctggtcctgctggcaaagaag  
. . . . .  
gcggaaggtccccgtggtgagactggccctgctggacgtcctggtgaagttggtcccc  
. . . . .  
ctggtccccctggccctgctggcgagaaaggatccccctggtgctgatggtcctg

Exon 39 | Start: 17102 | End: 17263 | Length: 161  
BE AWARE: This section overlaps with the following exon

. . . . .  
tgttgagcctctcccctctccccaggcagcgggtggcaggtgagggcagctgggtcggatg  
. . . . .  
agttggctgttctccctctgactgttcctatgttctctccttcagggtgctactggttt  
. . . . .  
ccctggtgctgctggccgagtcggtcctcctggcccctctgtaagtctctgcagcagagt  
. . . . .  
ccactgctctaggttgggggtgctgggtgggggctgccagaaggatggtggggctgactg  
. . . . .  
aggacccaatgatgcaccagagccccctggagtctgacagcccctcctatcctcatccag  
|2671 |2681 |2691 |2701 |2711 |2721

GGAAATGCTGGACCCCTGGCCCTCCTGGTCCTGCTGGCAAAGAAGGCGGCAAAGGTCCC  
G N A G P P G P P G P A G K E G G K G P  
|891 |901  
|2731 |2741 |2751 |2761 |2771 |2781  
CGTGGTGAGACTGGCCCTGCTGGACGTCCTGGTGAAGTTGGTCCCCCTGGTCCCCCTGGC  
R G E T G P A G R P G E V G P P G P P G  
|911 |921  
|2791 |2801 |2811 |2821 . . .  
CCTGCTGGCGAGAAAGGATCCCCCTGGTGCTGATGGTCCTGCTgtaagtccagctcagat  
P A G E K G S P G A D G P A  
|931 |941  
. . . . .  
ctctgcagctccggaggtgtgcagagctggggaggggtccctgtgctgctctctggcacc  
. . . . .  
tcaccctgtttgcctcccaaagggtgctcctgggtactcccggtcctcaaggtattgctg  
. . . . .  
gacagcgtggtgtggtcggcctgcctgggtcagagaggagagagaggcttccctggtcttc  
. . . . .  
ctggcccctctgtaagtgtccccctcaccttggggggccctgagaaaaaccatcacagga  
. . . . .  
cttgagtggggcggagccaaggagaacagatttggttagaga

Exon 40 | Start: 17365 | End: 17472 | Length: 107  
BE AWARE: This section overlaps with the following exon

. . . . .  
cccctggagctctgacagcccctcctatcctcatccagggaatgctggaccccctggccc  
. . . . .  
tcctggtcctgctggcaaagaaggcggcaaagggtccccgtggtgagactggccctgctgg  
. . . . .  
acgtcctggtgaagttggtccccctggtccccctggccctgctggcgagaaaggatcccc  
. . . . .

tgggtgctgatgggtcctgctgtaagtgccagctcagatctctgcagctccggaggtgtgca  
 . . . . .  
 gagctgggggaggggtccctgtgctgctctctggcacctcaccctgtttgcctcccaaag  
 |2831 |2841 |2851 |2861 |2871 |2881  
 GGTGCTCCTGGTACTCCCGGCCTCAAGGTATTGCTGGACAGCGTGGTGTGGTCGGCCTG  
 G A P G T P G P Q G I A G Q R G V V G L  
 |951 |961  
 |2891 |2901 |2911 |2921 |2931 . .  
 CCTGGTCAGAGAGGAGAGAGAGAGGCTTCCCTGGTCTTCTGGCCCCCTCTgtaagtgcccc  
 P G Q R G E R G F P G L P G P S  
 |971  
 . . . . .  
 ctcaccttggggggccctgagaaaaacatcacaggacttggagtggggcggagccaagg  
 . . . . .  
 agaacagatttggttagagatgactccagcggactcaagggtcctccagaccctatctct  
 . . . . .  
 ggcctgactctttcttctcccttaggggtgaacctggcaacaaggctcctctggagcaag  
 . . . . .  
 tgggtgaacgtgggtccccctgggtcccatggggccccctggattggctggacccccctggtga  
 . . . . .  
 atctggacgtgaggtgagcagtcacccagcccccatgccagtaccctca

Exon 41 | Start: 17630 | End: 17737 | Length: 107  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ctctctggcacctcaccctgtttgcctcccaaagggtgctcctggtactcccgggcctc  
 . . . . .  
 aaggatttgctggacagcgtggtgtggtcggcctgcctggtcagagaggagagagaggct  
 . . . . .  
 tccctgggtcttctggccccctgtaagtggccccctcaccttggggggccctgagaaaa

. . . . .  
 accatcacaggacttggagtggggcggagccaaggagaacagatttggtagagatgactc  
 . . . . .  
 cagcggactcaagggtcctcccagaccctatctctggcctgactctttcttctcccttag  
 |2941 |2951 |2961 |2971 |2981 |2991  
 GGTGAACCTGGCAAACAAGGTCCCTCTGGAGCAAGTGGTGAACGTGGTCCCCCTGGTCCC  
 G E P G K Q G P S G A S G E R G P P G P  
 |981 |991  
 |3001 |3011 |3021 |3031 |3041 . .  
 ATGGGCCCCCTGGATTGGCTGGACCCCCTGGTGAATCTGGACGTGAGgtgagcagtgccc  
 M G P P G L A G P P G E S G R E  
 |1001 |1011  
 . . . . .  
 cagccccatgccagtagccctcagcatggccattgtggccttgccctaagccctcttcccc  
 . . . . .  
 ggctgactctcacttctctctctctctctctgcagggggctcctgggtgccgaagggtccc  
 . . . . .  
 ctggacgagacggttctcctggcgccaaggtaagatggcaacactccatgaccacagcct  
 . . . . .  
 tgtctgctgcttccctgccccatcctggcccttcacccggggctgaccatattcccctg  
 . . . . .  
 ctctccccgccagggtgaccgtggtgagaccggccccgctggaccccc

Exon 42 | Start: 17845 | End: 17898 | Length: 53  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 agaacagatttggttagagatgactccagcggactcaagggtcctcccagaccctatctct  
 . . . . .  
 ggctgactctttcttctcccttaggggtgaacctggcaacaaggtccctctggagcaag  
 . . . . .  
 tgggtgaacgtgggtccccctgggtcccatgggccccctggattggctggacccccctgggtga



. . . . .  
 atctggacgtgaggtgagcagtcacccagcccccatgccagtaccctcagcatggccattg  
 . . . . .  
 tggccttgccctaagccctcttccccggctgactctcacttctctctctctctctgcag  
 |3051 |3061 |3071 |3081 |3091 .  
 GGGGCTCCTGGTGCCGAAGGTTCCCCTGGACGAGACGGTTCTCCTGGCGCCAAGgtaaga  
 G A P G A E G S P G R D G S P G A K  
 |1021 |1031  
 . . . . .  
 tggcaacactccatgaccacagccttgctctgctgcttcctgccccatcctggcccttca  
 . . . . .  
 cccggggctgaccatattcccctgctctccccgccagggtgaccgtggtgagaccggcc  
 . . . . .  
 ccgctggacccccctggctgctcctggctgctcctggctggccctggccccgtggccctgctg  
 . . . . .  
 gcaagagtggatgatcgtgggtgagactgtaagtagctgggctccagttccctgtacctggt  
 . . . . .  
 caggccagggactcttcaggcctccttagaggcctggggatgggtgtcggactt

Exon 43 | Start: 18003 | End: 18110 | Length: 107

. . . . .  
 gattggctggacccccctggatgaatctggacgtgaggtgagcagtcacccagcccccatgcc  
 . . . . .  
 agtaccctcagcatggccattgtggccttgccctaagccctcttccccggctgactctcac  
 . . . . .  
 ttctctctctctctctctgcagggggctcctggctgccgaaggttcccctggacgagacgg  
 . . . . .  
 ttctcctggcgccaaggtgaagatggcaacactccatgaccacagccttgctctgctgttc  
 . . . . .

cctgccccatcctggcccttcacccggggctgacccatattcccctgctctccccgccag

|3101 |3111 |3121 |3131 |3141 |3151  
GGTGACCGTGGTGAGACCGGCCCCGCTGGACCCCTGGTGCTCCTGGTGCTCCTGGTGCC  
G D R G E T G P A G P P G A P G A P G A  
|1041 |1051

|3161 |3171 |3181 |3191 |3201 . .  
CCTGGCCCCGTTGGCCCTGCTGGCAAGAGTGGTGATCGTGGTGAGACTgtaagtagctgg  
P G P V G P A G K S G D R G E T  
|1061

. . . . .  
gctccagttccctgtacctgggtcaggccaggactcttcaggcctccttagaggcctggg

. . . . .  
gatgggtgtcggacttcacccaggcagggggaggaaaggagatcctgcaagatgtcaggg

. . . . .  
ccttaatcaaaaaactgagttaaagctcagccccaagtcccctctcccagacaggaccg

. . . . .  
cctctcccatgagttggccccagctcccgtgaagattgcagtggggagggtttccctggga

. . . . .  
gttgggagagatggccacagtgggaagcagctgaggagagagagatcc

Exon 44 | Start: 18491 | End: 18544 | Length: 53  
BE AWARE: This section overlaps with the following exon

. . . . .  
tcggacttcacccaggcagggggaggaaaggagatcctgcaagatgtcagggccttaatc

. . . . .  
caaaaaactgagttaaagctcagccccaagtcccctctcccagacaggaccgcctctccc

. . . . .  
atgagttggccccagctcccgtgaagattgcagtggggagggtttccctgggagttggggag

. . . . .  
agatggccacagtgggaagcagctgaggagagagagatccagcagagggggaggcctcatc

. . . . .  
ctgcagccccagcctcagccttccctggccaagagctcatgctttccttgctctccccag

          |3211      |3221      |3231      |3241      |3251      |3261.  
GGTCCTGCTGGTCCCGCCGGTCCTGTCTGGCCCTGTTGGCGCCCGTGGCCCCGCCgtaagt  
G  P  A  G  P  A  G  P  V  G  P  V  G  A  R  G  P  A  
          |1071  |1081

. . . . .  
accctgctgtgtccccatgcctttagaactctacagatgcagacagtgccccactcgat

. . . . .  
gccaatggaacttccgcctgacagtttgtccctttctcttcttagggaccccaaggccc

. . . . .  
ccgtggtgacaagggtgagacaggcgaacaggcgacagaggcataaagggtcaccgtgg

. . . . .  
cttctctggcctccagggtccccctggccctcctgtaagtatgctcagccccctcccagt

. . . . .  
cccatgctgtgtgtgggataggagggggagcttcgcctcagtttccccctct

Exon 45 | Start: 18657 | End: 18764 | Length: 107

. . . . .  
ctgggagttgggagagatggccacagtgggaagcagctgaggagagagagatccagcaga

. . . . .  
ggggaggcctcatcctgcagccccagcctcagccttccctggccaagagctcatgcttcc

. . . . .  
cttgctctccccagggtcctgctggtcccgcggtcctgtcggccctgttggcgcccggtg

. . . . .  
gccccgccgtaagtaccctgctgtgtcccccagcctttagaactctacagatgcagaca

. . . . .  
gtgccccactcgatgccaatggaacttccgcctgacagtttgtccctttctcttcttag

          |3271      |3281      |3291      |3301      |3311      |3321  
GGACCCCAAGGCCCGTGGTGACAAGGGTGAGACAGGCGAACAGGGCGACAGAGGCATA



G S P G E Q G P S G A S G P A G P R  
 |1131 |1141

. . . . .  
 catgccttctctctctcttctgagccccaagcccaggctcacctcggggacccttgcc  
 . . . . .  
 agggaccagggcaccctttgcctctctggagaagggttcaggacaggagtgaggcaaag  
 . . . . .  
 agaggaagaatcctgaacaaacaatctgatctagctttggcctctctgctccccaatccg  
 . . . . .  
 tcctccctggctcagcggctgggaggagctatggcatgtcctatggaaaggagctgagg  
 . . . . .  
 ctggctctatgagccgtggggccagagccagcaggagggtggtgggcctctcc

Exon 47 | Start: 19518 | End: 19625 | Length: 107  
 BE AWARE: This section overlaps with the following exon

. . . . .  
 ttgccagggacccaggcaccctttgcctctctggagaagggttcaggacaggagtgagg  
 . . . . .  
 caaagagaggaagaatcctgaacaaacaatctgatctagctttggcctctctgctcccca  
 . . . . .  
 atccgtcctccctggctcagcggctgggaggagctatggcatgtcctatggaaaggagc  
 . . . . .  
 tgaggctggctctatgagccgtggggccagagccagcaggagggtggtgggcctctcct  
 . . . . .  
 ccagagctgggggtgttcgggcttctggcagcctttctcaaaccatttccccactccag

|3431 |3441 |3451 |3461 |3471 |3481  
 GGTCCCCCTGGCTCTGCTGGTGCTCCTGGCAAAGATGGACTCAACGGTCTCCCTGGCCCC  
 G P P G S A G A P G K D G L N G L P G P  
 |1151 |1161

|3491 |3501 |3511 |3521 |3531. .

tcatccccctctgctcatggccctccagcccccaagcacttggatgccggaatccccac  
tctcttccctctctgtgcagggtcccccggccctcctggacctcctgggtccccctggtc  
ctcccagcgctggtttcgacttcagcttctgccccagccacctcaagagaaggctcacg  
atggtggccgctactaccgggctgatgatgccaatgtggttcgtgacctgacctcgagg  
tggacaccaccctcaagagcctgagccagcagatcgagaacatccgga

gtggggccagagccagcagggagggttggtgggcctctcctccagagctggggttgttcgg  
gcttctggcagcctttctcaaaccatttccccactccagggtccccctggctctgctgg  
tgctcctggcaaagatggactcaacggtctccctggccccattggggccccctggctcctg  
cggtcgactggtgatgctggtcctgttgtagtagccctcatcccctctgctcatggc  
cctccagccccaaagcatttggatgccggtaatccccactctcttccctctctgtgcag

46

	3601	3611	3621	3631	3641	3651														
	TTCAGCTTCCTGCCCCAGCCACCTCAAGAGAAGGCTCACGATGGTGGCCGCTACTACCGG																			
	F	S	F	L	P	Q	P	P	Q	E	K	A	H	D	G	G	R	Y	Y	R
		1201					1211													
	3661	3671	3681	3691	3701	3711														
	GCTGATGATGCCAATGTGGTTCGTGACCGTGACCTCGAGGTGGACACCACCTCAAGAGC																			
	A	D	D	A	N	V	V	R	D	R	D	L	E	V	D	T	T	L	K	S
		1221					1231													
	3721	3731	3741	3751	3761	3771														
	CTGAGCCAGCAGATCGAGAACATCCGGAGCCCAGAGGGCAGCCGCAAGAACCCCGCCCGC																			
	L	S	Q	Q	I	E	N	I	R	S	P	E	G	S	R	K	N	P	A	R
		1241					1251													
	3781	3791	3801	3811	.	.	.													
	ACCTGCCGTGACCTCAAGATGTGCCACTCTGACTGGAAGAGTGgtgagggcctgccctag																			
	T	C	R	D	L	K	M	C	H	S	D	W	K	S	G					
		1261					1271													
	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
	cctctccctccctcctactcctgcatgccagggtcccatgcccatatgtgccctacc																			
	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
	atatggtgctggctgctccctttccctgactccatcttgccctgccctaccacaggagag																			
	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
	tactggattgacccaaccaaggctgcaacctggatgccatcaaagtcttctgcaacatg																			
	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
	gagactggtgagacctgcgtgtacccactcagcccagtggtggccagaagaactggtac																			
	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
	atcagcaagaacccaaggacaagaggcatgtctgtggttcggcg																			

Exon 49 | Start: 20133 | End: 20323 | Length: 190  
BE AWARE: This section overlaps with the following exon

.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
accgggctgatgatgccaatgtggttcgtgacctgacctcgaggtggacaccaccctca																				
.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.

agagcctgagccagcagatcgagaacatccggagcccagagggcagccgcaagaaccccg

. . . . .  
cccgcacctgccgtgacctcaagatgtgccactctgactggaagagtggtagggcctgc

. . . . .  
cctagcctctccctccctcctactcctgccaatgccaggggtcccatgcccatatgtgccc

. . . . .  
ctaccatatggtgctggctgctccctttccctgactccatcttgccctgccctaccacag

3821	3831	3841	3851	3861	3871														
GAGAGTACTGGATTGACCCCAACCAAGGCTGCAACCTGGATGCCATCAAAGTCTTCTGCA																			
E	Y	W	I	D	P	N	Q	G	C	N	L	D	A	I	K	V	F	C	N
								1281											1291

3881	3891	3901	3911	3921	3931														
ACATGGAGACTGGTGAGACCTGCGTGTACCCCACTCAGCCCAGTGTGGCCCAGAAGAACT																			
M	E	T	G	E	T	C	V	Y	P	T	Q	P	S	V	A	Q	K	N	W
								1301											1311

3941	3951	3961	3971	3981	3991														
GGTACATCAGCAAGAACCCCAAGGACAAGAGGCATGTCTGGTTCGGCGAGAGCATGACCG																			
Y	I	S	K	N	P	K	D	K	R	H	V	W	F	G	E	S	M	T	D
								1321											1331

4001	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
ATGGATTCCAGgtgcgtgagctggacctcagagccagtgtaggagatgggctagcccag																			
G	F	Q																	

. . . . .  
tgctcagaaggacatgaagtcctggagtaggtctctgctaagggtgatggacagagctg

. . . . .  
ggctgggaggcaggggtctcaggtccctgatagtggttcagacacaggctgccgatgggc

. . . . .  
aggtggtgcctcctctcgatataacggtgcattgggcagctctctgaggaccctggacag

. . . . .  
ggaggccaggcaggactagaggttcccgcatagcggctcactcttccctctctcccctcc

. . .  
cctgcagttcg



Exon 50 | Start: 20620 | End: 20862 | Length: 242  
BE AWARE: This section overlaps with the following exon

```
. . . . .
ccaggTgcgtgagctggacctcagagccagtgttaggagatgggctagcccagtgctcag

. . . . .
aagggacatgaagtctggagtaggtctctgctaagggtgatggacagagctgggctggg

. . . . .
aggcaggggtctcaggtccctgatagtgggttcagacacaggctgccgatgggcaggtggt

. . . . .
gcctcctctcgatataacgggtgcattgggcagctctctgaggaccctggacagggaggcc

. . . . .
aggcaggactagaggttcccgcatagcggctcactcttccctctctcccctcccctgcag

      |4011      |4021      |4031      |4041      |4051      |4061
TTCGAGTATGGCGGCCAGGGCTCCGACCCTGCCGATGTGGCCATCCAGCTGACCTTCCTG
F E Y G G Q G S D P A D V A I Q L T F L
      |1341                        |1351

      |4071      |4081      |4091      |4101      |4111      |4121
CGCCTGATGTCCACCGAGGCCTCCAGAACATCACCTACCACTGCAAGAACAGCGTGGCC
R L M S T E A S Q N I T Y H C K N S V A
      |1361                        |1371

      |4131      |4141      |4151      |4161      |4171      |4181
TACATGACCAGCAGACTGGCAACCTCAAGAAGGCCCTGCTCCTCCAGGGCTCCAACGAG
Y M D Q Q T G N L K K A L L L Q G S N E
      |1381                        |1391

      |4191      |4201      |4211      |4221      |4231      |4241
ATCGAGATCCGCGCCGAGGGCAACAGCCGCTTACCTACAGCGTCACTGTCGATGGCTGC
I E I R A E G N S R F T Y S V T V D G C
      |1401                        |1411

ACGgtgagtgcccagaatccccaggcagggcccccacctctccggccttgggctttttggc
T
```

. . . . .  
 caggccatagtgccctctctccatcactcccacgtggtaatgccccctcccggtgtctcc  
 . . . . .  
 gccccaccccagagtcacaccggagcctggggcaagacagtgattgaatacaaaaaccacc  
 . . . . .  
 aagacctcccgctgcccacatcgcgatgtggcccccttgacgttggtgccccagaccag  
 . . . . .  
 gaattcggcttcgacgttggccctgtctgcttctgtaaactccctccatcccaacctgg  
 .  
 ctc

Exon 51 | Start: 20992 | End: 22544 | Length: 1552

. . . . .  
 accgaggcctcccagaacatcacctaccactgcaagaacagcgtggcctacatggaccag  
 . . . . .  
 cagactggcaacctcaagaaggccctgctcctccagggtccaacgagatcgagatccgc  
 . . . . .  
 gccgagggaacagccgcttcacctacagcgtcactgtcgatggctgcacggtgagtgcc  
 . . . . .  
 cagaatccccaggcagggccccacctctccggccttgggctttttggccaggccatagtg  
 . . . . .  
 ccctctctccatcactcccacgtggtaatgccccctcccggtgtctccgccccaccccag

4251	4261	4271	4281	4291	4301
AGTCACACCGGAGCCTGGGGCAAGACAGTGATTGAATACAAAACCAAGACCTCCCGC					
S H T G A W G K T V I E Y K T T K T S R					
	1421			1431	
4311	4321	4331	4341	4351	4361
CTGCCCATCATCGATGTGGCCCCCTTGGACGTTGGTGCCCCAGACCAGGAATTGGGCTTC					
L P I I D V A P L D V G A P D Q E F G F					
	1441			1451	

|4371        |4381        |4391|+1        |+11        |+21        |+31  
 GACGTTGGCCCTGTCTGCTTCCTGTAACTCCCTCCATCCCAACCTGGCTCCCTCCCACC  
 D V G P V C F L \*  
              |1461  
  
              |+41        |+51        |+61        |+71        |+81        |+91  
 CAACCAACTTTCCCCCAACCCGAAACAGACAAGCAACCCAACTGAACCCCTCAAAA  
  
              |+101        |+111        |+121        |+131        |+141        |+151  
 GCCAAAAATGGGAGACAATTTACATGGACTTTGGAAAATATTTTTTCCTTTGCATTC  
  
              |+161        |+171        |+181        |+191        |+201        |+211  
 ATCTCTCAAACCTTAGTTTTTATCTTTGACCAACCGAACATGACCAAAAACCAAAGTGCA  
  
              |+221        |+231        |+241        |+251        |+261        |+271  
 TTCAACCTTACCAAAAAAAAAAAAAAAAAAGAATAAATAAAATAACTTTTAAAAAAGGA  
  
              |+281        |+291        |+301        |+311        |+321        |+331  
 AGCTTGGTCCACTTGCTTGAAGACCCATGCGGGGGTAAGTCCCTTTCTGCCCGTTGGGCT  
  
              |+341        |+351        |+361        |+371        |+381        |+391  
 TATGAAACCCCAATGTGCCCCTTTCTGCTCCTTTCTCCACCCCCCTTGGGGCCTCCCC  
  
              |+401        |+411        |+421        |+431        |+441        |+451  
 TCCACTCCTTCCCAAATCTGTCTCCCAGAAGACACAGGAAACAATGTATTGTCTGCCCA  
  
              |+461        |+471        |+481        |+491        |+501        |+511  
 GCAATCAAAGGCAATGCTCAAACACCCAAGTGGCCCCCACCCTCAGCCCGCTCCTGCCCG  
  
              |+521        |+531        |+541        |+551        |+561        |+571  
 CCCAGCACCCCCAGGCCCTGGGGGACCTGGGGTTCTCAGACTGCCAAAGAAGCCTTGCCA  
  
              |+581        |+591        |+601        |+611        |+621        |+631  
 TCTGGCGCTCCCATGGCTCTTGCAACATCTCCCCTTCGTTTTTGAGGGGGTCATGCCGGG  
  
              |+641        |+651        |+661        |+671        |+681        |+691  
 GGAGCCACCAGCCCCTCACTGGGTTCTGGAGGAGAGTCAGGAAGGGCCACGACAAAGCAGA  
  
              |+701        |+711        |+721        |+731        |+741        |+751  
 AACATCGGATTTGGGGAACGCGTGTCAATCCCTTGTGCCGAGGGCTGGGCGGGAGAGAC  
  
              |+761        |+771        |+781        |+791        |+801        |+811  
 TGTTCTGTTCTTGTGTAAGTGTGTTGCTGAAAGACTACCTCGTTCTTGTCTTGATGTGT  
  
              |+821        |+831        |+841        |+851        |+861        |+871

CACCGGGGCAACTGCCTGGGGGCGGGGATGGGGGCAGGGTGGAAGCGGCTCCCCATTTTA

          |+881      |+891      |+901      |+911      |+921      |+931  
TACCAAAGGTGCTACATCTATGTGATGGGTGGGGTGGGGAGGGAATCACTGGTGCTATAG

          |+941      |+951      |+961      |+971      |+981      |+991  
AAATTGAGATGCCCCCAGGCCAGCAAATGTCCTTTTTTGTCAAAGTCTATTTTTATT

          |+1001     |+1011     |+1021     |+1031     |+1041     |+1051  
CCTTGATATTTTTCTTTTTTTTTTTTTTTTTTGTGGATGGGGACTTGTGAATTTTTCTA

          |+1061     |+1071     |+1081     |+1091     |+1101     |+1111  
AAGGTGCTATTTAACATGGGAGGAGAGCGTGTGCGGCTCCAGCCCAGCCCGCTGCTCACT

          |+1121     |+1131     |+1141     |+1151     |+1161     |+1171  
TTCCACCCTCTCTCCACCTGCCTCTGGCTTCTCAGGCCTCTGCTCTCCGACCTCTCTCCT

          |+1181     |+1191     |+1201     |+1211     |+1221     |+1231  
CTGAAACCCTCCTCCACAGCTGCAGCCCATCCTCCCGGCTCCCTCCTAGTCTGTCTGCG

          |+1241     |+1251     |+1261     |+1271     |+1281     |+1291  
TCCTCTGTCCCGGGTTTCAGAGACAACCTCCCAAAGCACAAAGCAGTTTTTCCCCTAG

          |+1301     |+1311     |+1321     |+1331     |+1341     |+1351  
GGGTGGGAGGAAGCAAAAGACTCTGTACCTATTTTGTATGTGTATAATAATTTGAGATGT

          |+1361     |+1371     |+1381     |+1391     |+1401     .  
TTTTAATTATTTTGATTGCTGGAATAAAGCATGTGGAATGACCCAAACATAAtccgcag

. . . . .  
tggcctcctaatttccttctttggagttgggggaggggtagacatggggaaggggctttg

. . . . .  
gggtgatgggcttgccttccattcctgccctttccctccccactattctcttctagatcc

. . . . .  
ctccataacccccactcccctttctctcacccttcttataaccgcaaactttctacttct

. . . . .  
ctttcattttctattcttgcaatttccttgcaccttttccaaatcctcttctcccctgca

. . . . .  
ataccatacaggcaatccacgtgcacaacacacacacactcttcacatctg

LRG Parser: Version: 0.3, Version Date: 11/02/2015

Reader: Version: 0.3, Version Date: 11/02/2015  
Writer: Version: 0.3, Version Date: 11/02/2015  
Control: Version: 0.3, Version Date: 11/02/2015