

Gene: BRCA1 - Sequence: NG\_005905.2  
 Transcript: NM\_007294.3 - Protein: NP\_009225.1  
 Date : February 19, 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: 92500 | End: 92713 | Length: 213

```

. . . . .
tacttatatTTTaccgaaactggagacctccattagggcggaagagtgggggattggggac
. . . . .
ctcttcttacgactgctttggacaataggtagcgattctgaccttcgtacagcaattact
. . . . .
gtgatgcaataagccgcaactggaagagtagaggctagaggcaggcactttatggcaaa
. . . . .
ctcaggtagaattcttctcttccgtctctttccttttacgtcatccgggggcagactgg
. . . . .
gtggccaatccagagccccgagagacgcttggtctctttctgtccctcccatcctctgatt
      |-229      |-219      |-209      |-199      |-189      |-179
GTACCTTGATTTTCGTATTCTGAGAGGCTGCTGCTTAGCGGTAGCCCCTTGGTTTCCGTGG
      |-169      |-159      |-149      |-139      |-129      |-119
CAACGGAAGCGCGGAATTACAGATAAATTAAACTGCGACTGCGCGGCGTGAGCTCG
      |-109      |-99       |-89       |-79       |-69       |-59
CTGAGACTTCCTGGACGGGGACAGGCTGTGGGGTTTCTCAGATAACTGGGCCCTGCGC
      |-49       |-39       |-29       .        .        .        .
TCAGGAGGCCTTCACCTCTGCTCTGGGTAAAGgtagtagagtcccgggaaaggacagg
. . . . .
gggcccgaagtgatgctctggggtactggcgtgggagagtggatttccgaagctgacagat
. . . . .

```

gggtattctttgacgggggtagggcggaacctgagaggcgtaaggcgttgtgaaccct  
 . . . . .  
 ggggaggggggcagttttaggtcgcgagggaagcgctgaggatcaggaagggggcactg  
 . . . . .  
 agtgtccgtgggggaatcctcgtgataggaactggaatatgccttgagggggacactatg  
 . . . . .  
 tctttaaaaacgtcggctggtcatgaggtcagg

Exon 2 | Start: 93868 | End: 93967 | Length: 99

. . . . .  
 aggctaccaccacctacccggtcagtcactcctctgtagctttctctttcttgagaaag  
 . . . . .  
 gaaaagaccaaggggttggcagcaatatgtgaaaaattcagaatttatgttgtctaata  
 . . . . .  
 tacaaaaagcaacttctagaatctttaaaaataaaggacgttgtcattagttctttgggtt  
 . . . . .  
 tgtattattctaaaccttccaaatcttaaattactttattttaaaatgataaaatgaa  
 . . . . .  
 gttgtcattttataaacctttttaaaagatatatatatgtttttctaatgtgttaaag

-19	-9	1	11	21	31	41
TTCATTGGAACAGAAAGAAATGGATTATCTGCTCTTCGCGTTGAAGAAGTACAAAATGT						
		M	D	L	S	A
				L	R	V
				E	E	V
					Q	N
						V
		1			11	

51	61	71				
CATTAATGCTATGCAGAAATCTTAGAGTGTCCCATCTGgtaagtcagcacaagagtgtgta						
I	N	A	M	Q	K	I
				L	E	C
				P	I	C
		21				

. . . . .  
 ttaatttgggattcctatgattatctcctatgcaaatgaacagaattgaccttacataact  
 . . . . .

aggaagaaaagacatgtctagtaagattaggctattgtaattgctgattttcttaactg  
 . . . . .  
 aagaactttaaaaatatagaaaatgattccttggttctccatccactctgcctctccact  
 . . . . .  
 cctctccttttcaacacaaatcctgtggtccgggaaagacagggactctgtcttgattgg  
 . . . . .  
 ttctgcactggggcaggaatctagtttagattaactggc

Exon 3 | Start: 102204 | End: 102258 | Length: 54

. . . . .  
 aattcgtacgaactattatcaactaatcttttaaagtctgatagtagatagagtattga  
 . . . . .  
 agggatcaatataattctgttttgatatctgaaagctcactgaaggtaaggatcgtattc  
 . . . . .  
 tctgctgtattctcagttcctgacacagcagacatttaataaatattgaacgaacttgag  
 . . . . .  
 gccttatgttgactcagtcataacagctcaaagttgaacttattcactaagaatagcttt  
 . . . . .  
 atttttaataaattattgagcctcatttattttctttttctccccctaccctgctag  
 |81 |91 |101 |111 |121 |131 .  
 TCTGGAGTTGATCAAGGAACCTGTCTCCACAAAGTG TGACCACATATTTTGCAAgttaagt  
 L E L I K E P V S T K C D H I F C K  
 |31 |41  
 . . . . .  
 ttgaatgtgttatgtggctccattattagcttttggtttgtccttcataaaccaggaaa  
 . . . . .  
 cacctaactttatagaagctttactttcttcaattaagtgagaaacgaaaaatccaactcc  
 . . . . .  
 atttcattctttctcagagagtatatagttatcaaaagttggttgtaatcatagttcctg

. . . . .  
 gtaaagttttgacatatattatcttttttttttttttggagacaaagtctcgctctgtcg  
 . . . . .  
 cccaggctggagtgagtgaggcatgatcttggctcactgcaacctccgcccccg

Exon 4 | Start: 111450 | End: 111528 | Length: 78

. . . . .  
 tatgaatataatcattaaatagccatattaacttttattaagttttatgtgatcataaca  
 . . . . .  
 gtaagccatatgcatgtaagttcagttttcatagatcattgcttatgtagtttaggtttt  
 . . . . .  
 tgcttatgcagcatccaaaaacaattaggaaactattgcttgtaattcacctgccattac  
 . . . . .  
 tttttaaatggctcttaagggcagttgtgagattatcttttcatggctatttgccttttg

. . . . .  
 agtattctttctacaaaaggaagtaaattggtctttctttctttataatttatag  
 |141 |151 |161 |171 |181 |191  
 ATTTTGCATGCTGAAACTTCTCAACCAGAAGAAAGGGCCTTCACAGTGTCTTTATGTAA  
 F C M L K L L N Q K K G P S Q C P L C K  
 |51 |61

|201 |211 . . . . .  
 GAATGATATAACCAAAAGgtatataatttggtaatgatgctaggttgaagcaaccacag  
 N D I T K R  
 |71

. . . . .  
 taggaaaaagtagaaattatttaataacatagcggttcctataaaaccattcatcagaaaa  
 . . . . .  
 atttataaaaagagtttttagcacacagtaaattatttccaaagttattttcctgaaagtt  
 . . . . .  
 ttatgggacatctgccttatacaggtattagaaacttactgcctttctctaagtcttcta

. . . . .  
 gtgtaaaaacttgcagacttatgtaaagtagggctgtatcgccgtgccccattgtctgt  
 . . . . .  
 taatcttgtttttatatt

Exon 5 | Start: 113027 | End: 113116 | Length: 89

. . . . .  
 aaattaacaaaaaaaaaagtactctagttttctatgcaatgcattatatctgctgtgga  
 . . . . .  
 tttagggcagtattatatcagataattttaggcatttggtaggcttaaatgaatgacaaa  
 . . . . .  
 aagttactaaatcactgccatcacacggtttatacagatgtcaatgatgtattgattata  
 . . . . .  
 gaggttttctactgttgctgcatcttatttttattgtttacatgtcttttcttatttta  
 . . . . .  
 gtgtccttaaaagggtgataatcacttgctgagtggtgttttctcaaacaatttaatttcag

221	231	241	251	261	271
GAGCCTACAAGAAAGTACGAGATTTAGTCAACTTGTTGAAGAGCTATTGAAAAATCATTG					
S L Q E S T R F S Q L V E E L L K I I C					
		81			91

281	291	301	.	.	.	.	.
TGCTTTTCAGCTTGACACAGGTTTGGAGTgtaagtgttgaatatcccaagaatgacactc							
A F Q L D T G L E Y							
		101					

. . . . .  
 aagtgtgtccatgaaaactcaggaagtttgcaattactttctatgacgtggtgataa  
 . . . . .  
 gaccttttagtctaggttaatttttagttctgtatctgtaatctatttttaaaaaattact  
 . . . . .  
 ccactgggtctcacaccttattttatcaatcgtaagggtgcacatttttcacatcttaaca

. . . . .  
tctctgaaattgggaacattttactattgagggtgtgtcatttgtttaatttggtgtgctt

. . . . .  
tctttcttagtgatacagaaaataatagt

Exon 6 | Start: 113722 | End: 113862 | Length: 140

. . . . .  
tacattgttggtgtcttagcttttagtgaaatacagtattgataggcaaatttcttagtgt

. . . . .  
taaggtagaaaacaaggactctaaataactttgatggctgtgtatttgttttgtttcc

. . . . .  
taggagtaaaatttccagttgattttttaaaatttgatttttaaaaaaatcacaggtaa

. . . . .  
ccttaatgcattgtcttaacacaacaagagcatacatagggttctcttggtttctttg

. . . . .  
attataattcatacatttttctctaactgcaaacataatgttttcccttgatttttacag

          |311      |321      |331      |341      |351      |361  
ATGCAACAGCTATAATTTGCAAAAAAGGAAAAATAACTCTCCTGAACATCTAAAAGATG  
  A  N  S  Y  N  F  A  K  K  E  N  N  S  P  E  H  L  K  D  E  
                          |111                          |121

          |371      |381      |391      |401      |411      |421  
AAGTTTCTATCATCCAAAGTATGGGCTACAGAAACCGTGCCAAAAGACTTCTACAGAGTG  
  V  S  I  I  Q  S  M  G  Y  R  N  R  A  K  R  L  L  Q  S  E  
                          |131                          |141

          |431      |441      .      .      .      .      .      .  
AACCCGAAAATCCTTCCTTGgtaaaaccatttgttttcttcttcttcttcttcttctt  
  P  E  N  P  S  L

. . . . .  
cttttttttttctttttttttttgagatggagtccttgctctgtggcccaggctagaagc

. . . . .

agtcctcctgccttagcccccttagtagctgggattacaggcacgcgccaccatgccagg  
 . . . . .  
 ctaatTTTTgtatTTTTtagtagagacggggtttcatcatgttggccaggctgggtctcgaa  
 . . . . .  
 ctctaacctcaggtgatccaccacctcggctccccaattgctgggattacaggtgtg  
 . . . . .  
 agccactgtgcccggccggt

Exon 7 | Start: 118103 | End: 118209 | Length: 106

. . . . .  
 actactactattatTTTtagagactgggtctcactctgttgcttatgctggcttgaac  
 . . . . .  
 tcctggcctcaagcagtcctgctccagcctcccaaagtgctgggattataggcatgagct  
 . . . . .  
 accgctcccagccccagacattTTtagtgtgtaaattcctgggcattTTTTccaggcatca  
 . . . . .  
 tacatgttagctgactgatgatggtcaatttatTTTgtccatgggtgtcaagtttctcttc  
 . . . . .  
 aggaggaaaagcacagaactggccaacaattgcttgactgttctttaccatactgttttag

	451	461	471	481	491	501
CAGGAAAC	CAGTCT	CAGTGT	CCAAC	TCTCTA	ACCTTG	GAAC
Q	E	T	S	L	S	V
	151					
	511	521	531	541		
AAGCAG	CGGATA	CAACCT	CAAAAG	ACGTCT	GTCTAC	ATTGA
K	Q	R	I	Q	P	Q
	171					

. . . . .  
 ttttttaagtattttaataataattgctggattccttatcttatagttttgccccaaatct  
 . . . . .

tggtcataatTTGTATTTGTGGTAGGCAGCTTTGGGAAGTGAATTTTATGAGCCCTATGG  
 . . . . .  
 TGAGTTATAAAAAATGTAAAAGACGCAGTTCCACCTTGAAGAATCTTACTTTAAAAAGG  
 . . . . .  
 GAGCAAAAGAGGCCAGGCATGGTGGCTCACACCTGTAATCCAGCAGCTTTGGGAGGCCAA  
 . . . . .  
 AGTGGGTGGATCACCTGAGGTCGGGAGTTCGAGACCAGCCTAGCCA

Exon 8 | Start: 120694 | End: 120740 | Length: 46

. . . . .  
 TAGAAACGGGGTCTCAGCTTTGTTGGCCAGGCTGGTCTTGAACCTCTAACCTCAAATAAT  
 . . . . .  
 CCACCCATCTCGGCCTCCTCAAGTGCTGGGATTACAGGTGAGAGCCACTGTGCCTGGCGA  
 . . . . .  
 AGCCCATGCCTTTAACCAGCTTCTGTATTACATACTAGCTTAAGTACATTGTACCTGC  
 . . . . .  
 CACAGTAGATGCTCAGTAAATATTTCTAGTTGAATATCTGTTTTCAACAAGTACATTTT  
 . . . . .  
 TTAAACCCTTTTAATTAAGAAAAGTATTTGATTATTTTTTGGGGGAAATTTTTTAG  
 |551 |561 |571 |581 |591 . .  
 GATCTGATTCTTCTGAAGATACCGTTAATAAGGCAACTTATTGCAGGtgagtcaaagaga  
 S D S S E D T V N K A T Y C S  
 |191

. . . . .  
 ACCTTTGTCTATGAAGCTGGTATTTTCCTATTAGTTAATATTAAGGATTGATGTTTCTC  
 . . . . .  
 TCTTTTAAAAATATTTTAAGTATTTTAGGTTAGGATGTATGTGCAGTTTGTAT  
 . . . . .  
 ATAGGTAACACAGCAGTGGGATTTGGTGTATAGATTTTTTTCATCATCCGGGTACTAA



. . . . .  
gcataccccacagtttttgtttgctttctttctgaatttctccctcttcccaccttcct  
. . . . .  
ccctcaagtaggctgggtgtttctccagactagaatcatggatttg

Exon 9 | Start: 122061 | End: 122138 | Length: 77

. . . . .  
gtatttttagtagagatgggggtttcaccatgttggccaggctggtcttgaactcatgacc  
. . . . .  
tcaagtgggtccaccgcctcagcctcccaaagtgtggaattacaggcttgagccaccgt  
. . . . .  
gccagcaaccatttcatttcaactagaagtttctaaaggagagagcagctttcactaac  
. . . . .  
taaataagattggtcagctttctgtaatcgaaagagctaaaatgtttgatcttggtcatt  
. . . . .

tgacagttctgcatacatgtaactagtgtttcttattaggactctgtctttccctatag

601	611	621	631	641	651
TGTGGGAGATCAAGAATTGTTACAAATCACCCCTCAAGGAACCAGGGATGAAATCAGTTT					
V G D Q E L L Q I T P Q G T R D E I S L					
201			211		

661	. . . . .
GGATTCTGCAAAAAAGGgtaatggcaaagtttgccaacttaacaggcactgaaaagagag	
D S A K K A	
221	

. . . . .  
tgggtagatacagtactgtaattagattattctgaagaccatttgggacctttacaaccc  
. . . . .  
acaaaatctcttggcagagttagagtatcattctctgtcaaattgtcgtggtatggtctga  
. . . . .  
tagattttaatggtactagactaatgtacctataataagaccttctgtaactgattgttg

. . . . .  
ccctttcgtttttttttgtttgtttgtttgtttttttgagatggggtctcactctg  
. . . . .  
ttgccaggctggagtg

Exon 10 | Start: 123123 | End: 126549 | Length: 3426

. . . . .  
agttttctgatggccaatctgcttttaattcactcttagacgttagagaaataggtgtgg  
. . . . .  
tttctgcatagggaattctgaaattaaaaattaatggatcctaagtggaaataatct  
. . . . .  
aggtaaataggaattaaatgaaagagtatgagctacatcttcagtatacttggtagtta  
. . . . .  
tgagggttagtttcttaatatagccagttggttgatttccacctccaaggtgtatgaagt  
. . . . .  
atgtatttttttaatgacaattcagtttttgagtaccttggtatttttgtatattttcag

671	681	691	701	711	721
CTGCTTGTGAATTTTCTGAGACGGATGTAACAAATACTGAACATCATCAACCCAGTAATA					
A	C	E	F	S	E
	T	D	V	T	N
				T	E
				H	H
				Q	P
					S
					N
					N

731	741	751	761	771	781
ATGATTTGAACACCACTGAGAAGCGTGCAGCTGAGAGGCATCCAGAAAAGTATCAGGGTA					
D	L	N	T	T	E
		K	R	A	A
				E	R
				H	P
				E	K
				Y	Q
				G	S

791	801	811	821	831	841
GTTCTGTTTCAAACCTTGCATGTGGAGCCATGTGGCACAAATACTCATGCCAGCTCATTAC					
S	V	S	N	L	H
		V	E	P	C
			G	T	N
				T	H
				A	S
				S	S
				L	Q

851	861	871	881	891	901
AGCATGAGAACAGCAGTTTATTACTCACTAAAGACAGAATGAATGTAGAAAAGGCTGAAT					
H	E	N	S	S	L
		L	L	L	T
		K	D	R	M
			N	V	E
				K	A
				E	F

911	921	931	941	951	961
TCTGTAATAAAAGCAAACAGCCTGGCTTAGCAAGGAGCCAACATAACAGATGGGCTGGAA					
C	N	K	S	K	Q
	P	G	L	A	R
			S	Q	H
				N	R
				W	A
				G	S
		311			321
971	981	991	1001	1011	1021
GTAAGGAAACATGTAATGATAGGCGGACTCCCAGCACAGAAAAAAGGTAGATCTGAATG					
K	E	T	C	N	D
	R	R	T	P	S
			T	E	K
				K	V
				D	L
				N	A
		331			341
1031	1041	1051	1061	1071	1081
CTGATCCCCTGTGTGAGAGAAAAGAATGGAATAAGCAGAAACTGCCATGCTCAGAGAAATC					
D	P	L	C	E	R
	K	E	W	N	K
			Q	K	L
			P	C	S
				E	N
				P	
		351			361
1091	1101	1111	1121	1131	1141
CTAGAGATACTGAAGATGTTCTTGATAACACTAAATAGCAGCATTGAGAAAAGTTAATG					
R	D	T	E	D	V
	P	W	I	T	L
		N	S	S	I
			Q	K	V
				N	E
		371			381
1151	1161	1171	1181	1191	1201
AGTGGTTTTCCAGAAGTGATGAACTGTTAGGTTCTGATGACTCACATGATGGGGAGTCTG					
W	F	S	R	S	D
	E	L	L	G	S
		D	D	S	H
			D	G	E
			S	E	
		391			401
1211	1221	1231	1241	1251	1261
AATCAAATGCCAAAGTAGCTGATGTATTGGACGTTCTAAATGAGGTAGATGAATATTCTG					
S	N	A	K	V	A
	D	V	L	D	V
		L	N	E	V
			D	E	Y
			S	G	
		411			421
1271	1281	1291	1301	1311	1321
GTTCTTCAGAGAAAATAGACTTACTGGCCAGTGATCCTCATGAGGCTTTAATATGTAAAA					
S	S	E	K	I	D
	L	L	A	S	D
		P	H	E	A
			L	I	C
			K	S	
		431			441
1331	1341	1351	1361	1371	1381
GTGAAAGAGTTCACTCCAAATCAGTAGAGAGTAATATTGAAGACAAAATATTTGGGAAAA					
E	R	V	H	S	K
	S	V	E	S	N
		I	E	D	K
			I	F	G
			K	T	
		451			461
1391	1401	1411	1421	1431	1441
CCTATCGGAAGAAGGCAAGCCTCCCAACTTAAGCCATGTAAGTAAAAATCTAATTATAG					
Y	R	K	K	A	S
	L	P	N	L	S
		H	V	T	E
			N	L	I
			I	G	
		471			481

1451	1461	1471	1481	1491	1501
GAGCATTTGTTACTGAGCCACAGATAATACAAGAGCGTCCCCTCACAAATAAATTAAGC					
A	F	V	T	E	P
		Q	I	I	Q
			E	R	P
			L	T	N
			K	L	K
			R		
		491			501
1511	1521	1531	1541	1551	1561
GTAAAAGGAGACCTACATCAGGCCTTCATCCTGAGGATTTTATCAAGAAAGCAGATTTGG					
K	R	R	P	T	S
		G	L	H	P
		E	D	F	I
		K	K	A	D
		L	A		
		511			521
1571	1581	1591	1601	1611	1621
CAGTTCAAAAGACTCCTGAAATGATAAATCAGGGAACAAACGAGCAGATGGTC					
V	Q	K	T	P	E
		M	I	N	Q
		G	T	N	Q
		T	E	Q	N
		G			
		531			541
1631	1641	1651	1661	1671	1681
AAGTGATGAATATTACTAATAGTGGTCATGAGAATAAAACAAAAGGTGATTCTATTTCAGA					
V	M	N	I	T	N
		S	G	H	E
		N	K	T	K
		G	D	S	I
		Q			
		551			561
1691	1701	1711	1721	1731	1741
ATGAGAAAAATCCTAACCCAATAGAATCACTCGAAAAAGAATCTGCTTTCAAAACGAAAG					
E	K	N	P	N	P
		I	E	S	L
		E	K	E	S
		A	F	K	T
		K	A		
		571			581
1751	1761	1771	1781	1791	1801
CTGAACCTATAAGCAGCAGTATAAGCAATATGGAACCTGAATTAAATATCCACAATTCAA					
E	P	I	S	S	S
		I	S	N	M
		E	L	E	L
		N	I	H	N
		S	K		
		591			601
1811	1821	1831	1841	1851	1861
AAGCACCTAAAAAGAATAGGCTGAGGAGGAAGTCTTCTACCAGGCATATTCATGCGCTTG					
A	P	K	K	N	R
		L	R	R	K
		S	S	T	R
		H	I	H	A
		L	E		
		611			621
1871	1881	1891	1901	1911	1921
AACTAGTAGTCAGTAGAAATCTAAGCCACCTAATTGTACTGAATTGCAAATTGATAGTT					
L	V	V	S	R	N
		L	S	P	P
		N	C	T	E
		L	Q	I	D
		S	C		
		631			641
1931	1941	1951	1961	1971	1981
GTTCTAGCAGTGAAGAGATAAAGAAAAAAGTACAACCAAATGCCAGTCAGGCACAGCA					
S	S	S	E	E	I
		K	K	K	K
		Y	N	Q	M
		P	V	R	H
		S	R		
		651			661
1991	2001	2011	2021	2031	2041

GAAACCTACAACCTCATGGAAGGTAAAGAACCTGCAACTGGAGCCAAGAAGAGTAACAAGC  
 N L Q L M E G K E P A T G A K K S N K P  
 |671 |681

|2051 |2061 |2071 |2081 |2091 |2101  
 CAAATGAACAGACAAGTAAAAGACATGACAGCGATACTTTCCCAGAGCTGAAGTTAACAA  
 N E Q T S K R H D S D T F P E L K L T N  
 |691 |701

|2111 |2121 |2131 |2141 |2151 |2161  
 ATGCACCTGGTTCTTTTACTAAGTGTTCAAATACCAGTGAACCTAAAGAATTTGTCAATC  
 A P G S F T K C S N T S E L K E F V N P  
 |711 |721

|2171 |2181 |2191 |2201 |2211 |2221  
 CTAGCCTTCCAAGAGAAGAAAAAGAAGAGAACTAGAAACAGTTAAAGTGCTAATAATG  
 S L P R E E K E E K L E T V K V S N N A  
 |731 |741

|2231 |2241 |2251 |2261 |2271 |2281  
 CTGAAGACCCCAAAGATCTCATGTTAAGTGGAGAAAGGGTTTGCAAACTGAAAGATCTG  
 E D P K D L M L S G E R V L Q T E R S V  
 |751 |761

|2291 |2301 |2311 |2321 |2331 |2341  
 TAGAGAGTAGCAGTATTTTCATTGGTACCTGGTACTGATTATGGCACTCAGGAAAGTATCT  
 E S S S I S L V P G T D Y G T Q E S I S  
 |771 |781

|2351 |2361 |2371 |2381 |2391 |2401  
 CGTTACTGGAAGTTAGCACTCTAGGGAAGGCAAAAACAGAACCAAATAAATGTGTGAGTC  
 L L E V S T L G K A K T E P N K C V S Q  
 |791 |801

|2411 |2421 |2431 |2441 |2451 |2461  
 AGTGTGCAGCATTTGAAAACCCCAAGGGACTAATTCATGGTTGTTCCAAAGATAATAGAA  
 C A A F E N P K G L I H G C S K D N R N  
 |811 |821

|2471 |2481 |2491 |2501 |2511 |2521  
 ATGACACAGAAGGCTTTAAGTATCCATTGGGACATGAAGTTAACCACAGTCGGGAAACAA  
 D T E G F K Y P L G H E V N H S R E T S  
 |831 |841

|2531 |2541 |2551 |2561 |2571 |2581  
 GCATAGAAATGGAAGAAAGTGAACCTTGATGCTCAGTATTTGCAGAATACATTCAAGGTTT

I E M E E S E L D A Q Y L Q N T F K V S  
 |851 |861  
 |2591 |2601 |2611 |2621 |2631 |2641  
 CAAAGCGCCAGTCATTTGCTCCGTTTCAAATCCAGGAAATGCAGAAGAGGAATGTGCAA  
 K R Q S F A P F S N P G N A E E E C A T  
 |871 |881  
 |2651 |2661 |2671 |2681 |2691 |2701  
 CATTCTCTGCCACTCTGGGTCCTTAAAGAAACAAAGTCCAAAAGTCACTTTTGAATGTG  
 F S A H S G S L K K Q S P K V T F E C E  
 |891 |901  
 |2711 |2721 |2731 |2741 |2751 |2761  
 AACAAAAGGAAGAAAAATCAAGGAAAGAATGAGTCTAATATCAAGCCTGTACAGACAGTTA  
 Q K E E N Q G K N E S N I K P V Q T V N  
 |911 |921  
 |2771 |2781 |2791 |2801 |2811 |2821  
 ATATCACTGCAGGCTTTCCTGTGGTTGGTCAGAAAGATAAGCCAGTTGATAATGCCAAAT  
 I T A G F P V V G Q K D K P V D N A K C  
 |931 |941  
 |2831 |2841 |2851 |2861 |2871 |2881  
 GTAGTATCAAAGGAGGCTCTAGGTTTTGTCTATCATCTCAGTTCAGAGGCAACGAAACTG  
 S I K G G S R F C L S S Q F R G N E T G  
 |951 |961  
 |2891 |2901 |2911 |2921 |2931 |2941  
 GACTCATTACTCCAAATAAACATGGACTTTTACAAAACCCATATCGTATACCACCACTTT  
 L I T P N K H G L L Q N P Y R I P P L F  
 |971 |981  
 |2951 |2961 |2971 |2981 |2991 |3001  
 TTCCCATCAAGTCATTTGTTAAAACTAAATGTAAGAAAAATCTGCTAGAGGAAAACTTTG  
 P I K S F V K T K C K K N L L E E N F E  
 |991 |1001  
 |3011 |3021 |3031 |3041 |3051 |3061  
 AGGAACATTCAATGTCACCTGAAAGAGAAATGGGAAATGAGAACATTCCAAGTACAGTGA  
 E H S M S P E R E M G N E N I P S T V S  
 |1011 |1021  
 |3071 |3081 |3091 |3101 |3111 |3121  
 GCACAATTAGCCGTAATAACATTAGAGAAAATGTTTTTAAAGAAGCCAGCTCAAGCAATA  
 T I S R N N I R E N V F K E A S S S N I

		1031			1041
3131	3141	3151	3161	3171	3181
TTAATGAAGTAGGTTCCAGTACTAATGAAGTGGGCTCCAGTATTAATGAAATAGGTTCCA					
N E V G S S T N E V G S S I N E I G S S					
		1051			1061
3191	3201	3211	3221	3231	3241
GTGATGAAAACATTCAAGCAGAACTAGGTAGAAACAGAGGGCCAAAATTGAATGCTATGC					
D E N I Q A E L G R N R G P K L N A M L					
		1071			1081
3251	3261	3271	3281	3291	3301
TTAGATTAGGGGTTTTGCAACCTGAGGTCTATAAACAAAGTCTTCCTGGAAGTAATTGTA					
R L G V L Q P E V Y K Q S L P G S N C K					
		1091			1101
3311	3321	3331	3341	3351	3361
AGCATCCTGAAATAAAAAAGCAAGAATATGAAGAAGTAGTTCAGACTGTTAATACAGATT					
H P E I K K Q E Y E E V V Q T V N T D F					
		1111			1121
3371	3381	3391	3401	3411	3421
TCTCTCCATATCTGATTTTCAGATAACTTAGAACAGCCTATGGGAAGTAGTCATGCATCTC					
S P Y L I S D N L E Q P M G S S H A S Q					
		1131			1141
3431	3441	3451	3461	3471	3481
AGGTTTGTCTGAGACACCTGATGACCTGTTAGATGATGGTGAAATAAAGGAAGATACTA					
V C S E T P D D L L D D G E I K E D T S					
		1151			1161
3491	3501	3511	3521	3531	3541
GTTTTGCTGAAAATGACATTAAGGAAAGTTCTGCTGTTTTTAGCAAAAGCGTCCAGAAAAG					
F A E N D I K E S S A V F S K S V Q K G					
		1171			1181
3551	3561	3571	3581	3591	3601
GAGAGCTTAGCAGGAGTCCTAGCCCTTTCACCCATACACATTTGGCTCAGGGTTACCGAA					
E L S R S P S P F T H T H L A Q G Y R R					
		1191			1201
3611	3621	3631	3641	3651	3661
GAGGGGCCAAGAAATTAGAGTCCTCAGAAGAGAACTTATCTAGTGAGGATGAAGAGCTTC					
G A K K L E S S E E N L S S E D E E L P					
		1211			1221





. . . . .  
 gctaggacgtcatctttgactgaatgagctttaacatcctaattactggaggacttactt  
 . . . . .  
 ctggtttcattttataaaagcaaaccaggtgtcccaaagcaaggaatttaattcattttg  
 . . .  
 tgtgac

Exon 11 | Start: 126951 | End: 127040 | Length: 89

. . . . .  
 aaaggaactgcttcttaaacttgaacatgttcctcctaagggtgcttttcatagaaaaaa  
 . . . . .  
 gtccttcacacagctaggacgtcatctttgactgaatgagctttaacatcctaattactg  
 . . . . .  
 gtggacttacttctggtttcattttataaaagcaaaccaggtgtcccaaagcaaggaat  
 . . . . .  
 ttaatcattttgtgtgacatgaaagtaaatccagtcctgccaatgagaagaaaaagacac  
 . . . . .  
 agcaagttgcagcgtttatagtctgcttttacatctgaacctctgtttttgttatttaag

4101	4111	4121	4131	4141	4151
GTGAAGCAGCATCTGGGTGTGAGAGTGAAACAAGCGTCTCTGAAGACTGCTCAGGGCTAT					
E	A	A	S	G	C
	E	S	E	T	S
		V	S	E	D
			C	S	G
				L	S

4161	4171	4181	.	.	.	.	.
CCTCTCAGAGTGACATTTTAACCACTCAGgtaaaaagcgtgtgtgtgtgtgcacatgcgt							
S	Q	S	D	I	L	T	T
							Q

. . . . .  
 gtgtgtgggtgtcctttgcattcagtagtatgtatccacattcttaggtttgctgacatc  
 . . . . .  
 atctctttgaattaatggcacaattgtttgtggttcattgtctccttaaattagactgta

. . . . .  
 agcaccttgatggaactcatactacctttttatttcacacacacgcacacgcgcacacaca  
 . . . . .  
 gcctacacatacactgcctagctcattgtagcatactaaatactgattttaatgaataag  
 . . . . .  
 ctaaaccttcgaaaccatttgctaatacc

Exon 12 | Start: 135408 | End: 135580 | Length: 172

. . . . .  
 tattttttcactccctagcttttaaagaaaataaccaacttcaaaaggacatcacataa  
 . . . . .  
 catcaagtctatttgggggaatttgaggatttttccctcactaacatcatttggaata  
 . . . . .  
 atttcatgggcattaattgcatgaatgtggttagattaaaagggtgttcagctagaacttg  
 . . . . .  
 tagttccatactaggtgatttcaattcctgtgctaaaattaatttgatgatatttttc  
 . . . . .  
 atttaatggaaagcttctcaaagtatttcattttcttggtgccatttatcgtttttgaag

4191	4201	4211	4221	4231	4241
CAGAGGGATACCATGCAACATAACCTGATAAAGCTCCAGCAGGAAATGGCTGAAGTAGAA					
Q R D T M Q H N L I K L Q Q E M A E L E					
	1401			1411	

4251	4261	4271	4281	4291	4301
GCTGTGTTAGAACAGCATGGGAGCCAGCCTTCTAACAGCTACCCCTTCCATCATAAGTGAC					
A V L E Q H G S Q P S N S Y P S I I S D					
	1421			1431	

4311	4321	4331	4341	4351	.
TCTTCTGCCCTTGAGGACCTGCGAAATCCAGAACAAAGCACATCAGAAAAAGgtgtgtat					
S S A L E D L R N P E Q S T S E K A					
	1441			1451	

. . . . .  
 t g t t g g c c a a a c a c t g a t a t c t t a a g c a a a a t t c t t t c c t t c c c c t t t a t c t c t t c t g a  
 . . . . .  
 a g a g t a a g g a c c t a g c t c c a a c a t t t t a t g a t c c t t g c t c a g c a c a t g g g t a a t t a t g g a  
 . . . . .  
 g c c t t g g t t c t t g t c c c t g c t c a c a a c t a a t a t a c c a g t c a g a g g g a c c c a a g g c a g t c a  
 . . . . .  
 t t c a t g t t g t c a t c t g a g t a c t a c a a a g t a g a t g c t a t g g g g a g c c c a t g g a a g a t a  
 . . . . .  
 c a t g g t a t a c a a c a t a g c t c t t g c t c t a t t g g a a g c t a a g t g g a a t g g g a g a

Exon 13 | Start: 141369 | End: 141496 | Length: 127

. . . . .  
 t c a g g g c c a a g g a a t a t a g a t t t t t t t t t t c a g c c t t g t c t c a g c t g g g t g t c t t t a t t t  
 . . . . .  
 a c t c t g t c t t a a g t g t t c c t t t t a t t a t c a t t a t t a t t t t t a a t c a t t g a a t t c c a t t  
 . . . . .  
 t g g t g c t a g c a t c t g t c t g t t g c a t t g c t t g t g t t t a t a a a a t t c t g c c t g a t a t a c t t g  
 . . . . .  
 t t t a a a a c c a a t t t g t g t a t c a t a g a t t g a t g c t t t t g a a a a a a t c a g t a t t c t a a c c  
 . . . . .  
 t g a a t t a t c a c t a t c a g a a c a a g c a g t a a g t a g a t t t g t t t t c t c a t t c c a t t t a a a g

4361	4371	4381	4391	4401	4411
CAGTATTA	ACTTCACAGAAA	AGTAGTGAAT	ACCCTATAAG	CCAGAAATCC	CAGAAGGCCTTT
V	L	T	S	Q	K
		S	S	E	Y
		P	I	S	Q
		N	P	E	G
		L	S		
		1461			1471
4421	4431	4441	4451	4461	4471
CTGCTGACA	AAGTTTGAGGT	GTCTGCAGAT	AGTTCTACC	AGTAAAAATA	AAGAACCAGGAG
A	D	K	F	E	V
		S	A	D	S
		S	S	T	S
		K	N	K	E
		P	G	V	
		1481			1491

|4481 . . . . .  
 TGGAAAGgtaagaacatcaatgtaaagatgctgtggtatctgacatctttatttatatt  
 E R

. . . . .  
 gaactctgattgttaatTTTTTcaccatactttctccagtttttgcatacaggcatTT  
 . . . . .  
 atacacttttattgctctaggatacttcttttgTTtaacctatataggtttttgaacc  
 . . . . .  
 tataacataagctacaacatgagaaatgtgCGgtagatagatatgtcccttctgaaggt  
 . . . . .  
 cagaaaaaatataatggaggtaaaccctgaacaagcttggaactgatggtagacttct  
 . . .  
 tcaaggc

Exon 14 | Start: 143462 | End: 143653 | Length: 191

. . . . .  
 agcctcccgagtagctgagattacaggcgccagccaccacaccagctactgacctgctt  
 . . . . .  
 ttaaacagctgggagatatggtgcctcagaccaaccaaccccatgttatatgtcaaccc  
 . . . . .  
 tgacatattggcaggcaacatgaatccagacttctaggctgtcttgCGggctcttttttg  
 . . . . .  
 ccagtcatttctgatctctctgacatgagctgtttcatttatgctttggctgcccagcaa  
 . . . . .  
 gtatgatttgtcctttcacaattgggtggcgatggTTTTctccttcatttatctttctag

4491	4501	4511	4521	4531	4541
GTCATCCCTTCTAAATGCCCATCATTAGATGATAGGTGGTACATGCACAGTTGCTCTGG					
S S P S K C P S L D D R W Y M H S C S G					
	1501			1511	



tcatgtacccatTTTTCTCTTAACCTAactttattggtctTTTTaattcttaacagagac  
 . . . . .  
 cagaactttgtaattcaacattcatcgttgtgtaaattaaacttctcccattcctttcag  
 |4681 |4691 |4701 |4711 |4721 |4731  
 AGGGAACCCCTTACCTGGAATCTGGAATCAGCCTCTTCTCTGATGACCCTGAATCTGATC  
 G T P Y L E S G I S L F S D D P E S D P  
 |1561 |1571  
 |4741 |4751 |4761 |4771 |4781 |4791  
 CTTCTGAAGACAGAGCCCAGAGTCAGCTCGTGTGGCAACATACCATCTTCAACCTCTG  
 S E D R A P E S A R V G N I P S S T S A  
 |1581 |1591  
 |4801 |4811 |4821 |4831 |4841 |4851  
 CATTGAAAGTTCCCAATTGAAAGTTGCAGAATCTGCCAGAGTCCAGCTGCTGCTCATA  
 L K V P Q L K V A E S A Q S P A A A H T  
 |1601 |1611  
 |4861 |4871 |4881 |4891 |4901 |4911  
 CTACTGATACTGCTGGGTATAATGCAATGGAAGAAAGTGTGAGCAGGGAGAAGCCAGAAT  
 T D T A G Y N A M E E S V S R E K P E L  
 |1621 |1631  
 |4921 |4931 |4941 |4951 |4961 |4971  
 TGACAGCTTCAACAGAAAGGGTCAACAAAAGAATGTCCATGGTGGTGTCTGGCCTGACCC  
 T A S T E R V N K R M S M V V S G L T P  
 |1641 |1651  
 |4981 . . . . .  
 CAGAAGAATTTgtgagtgatccatatgtatctccctaagtactaagacttaacaacatt  
 E E F  
 |1661  
 . . . . .  
 ctggaaagagttttatgtaggtattgtcaattaataacctagaggaagaaatctagaaaa  
 . . . . .  
 caatcacagttctgtgtaatttaatttcgattactaatttctgaaaatttagatctagat  
 . . . . .  
 aaagctatagtggtgattatTTTatgtatatttacttgagaaaataattattaatatta  
 . . . . .  
 gtggaaaagctatactttgggtatgatataggactttcgaattggaattttcctttctat

. . .  
ctgtaaaagca

Exon 16 | Start: 150288 | End: 150376 | Length: 88

. . . . .  
gagacttcagggtgtcttagaattttttaaatgtaccctttctgagaaaaacagagactta

. . . . .  
aagctaggataactggtattctattttttttttttttttttttttacctccagcctgggt

. . . . .  
gacagagcaagactctgtctaaaaaaaaaaaaaaaaaattcactttaaatagttccagg

. . . . .  
acacgtgtagaacgtgcaggattgctacataggtaaacatatgccatgggtggaataacta

. . . . .  
gtattctgagctgtgtgctagaggttaactcatgataatggaatatttgatttaatttcag

          |4991      |5001      |5011      |5021      |5031      |5041  
ATGCTCGTGTACAAGTTTGCCAGAAAAACACCACATCACTTTAACTAATCTAATTACTGAA  
M L V Y K F A R K H H I T L T N L I T E  
                          |1671                                  |1681

          |5051      |5061      |5071      . . . . .  
GAGACTACTCATGTTGTTATGAAAACAGgtataccaagaacctttacagaataccttgca  
E T T H V V M K T D  
                          |1691

. . . . .  
tctgctgcataaaaccacatgaggcgaggcacggtggcgcatgcctgtaatcgagcact

. . . . .  
ttgggaggccgaggcgggcagatcacgagattaggagatcgagaccatcctggccagcat

. . . . .  
ggtgaaaccccgctcttactaaaaataaaaaattagctgggtgtggtcgcgtgcgcct

. . . . .  
gtagtcccagctactcgtgaggctgaggcaggagaatcacttgaaccggggagatggagg

. . . . .  
ttgcagtgagccgagatcatgccactgc

Exon 17 | Start: 154032 | End: 154110 | Length: 78

. . . . .  
tataatggagatctatagctagccttggcgtctagaagatgggtgttgagaagagggagt

. . . . .  
ggacagatatttcctctggtcttaacttcatatcagcctcccctagacttccaaatatcc

. . . . .  
atacctgctgggtataaattagtggtgttttcagcctctgattctgtcaccaggggtttta

. . . . .  
gaatcataaatccagattgatcttgggagtgtaaaaaactgaggctctttagcttcttag

. . . . .  
gacagcacttcctgattttgttttcaacttctaatacctttgagtgtttttcattctgcag

5081	5091	5101	5111	5121	5131
ATGCTGAGTTTGTGTGTGAACGGACACTGAAATATTTTCTAGGAATTGCGGGAGGAAAAT					
A E F V C E R T L K Y F L G I A G G K W					
				1701	1711

5141	5151	.	.	.	.	.	.	.	.
GGGTAGTTAGCTATTTCTgtaagtataataactatttctcccctcctccctttaacacctc									
V V S Y F W									

. . . . .  
agaattgcatttttacacctaacgtttaacacctaagggttttgctgatgctgagtctga

. . . . .  
gttacaaaaggctctttaattgtaataactaaactacttttatctttaatatcactttgtt

. . . . .  
cagataagctgggtgatgctgggaaaatgggtctctttttataactaataggacctaactctg

. . . . .  
ctcctagcaatggttagcatatgagctagggatttatttaaatagtcggcaggaatccatgt



. . . .  
gcagcaggcaaacttata

Exon 18 | Start: 154610 | End: 154651 | Length: 41

. . . . .  
ataactaataggacctaactctgctcctagcaatgtagcatatgagctagggatttat  
. . . . .  
aatagtcggcaggaatccatgtgcagcaggcaaacttataatgtttaaatcaacatcaa  
. . . . .  
ctctgtctccagaaggaaactgctgctacaagccttattaagggtgtggccttagagg  
. . . . .  
gaaggacctctcctctgtcattcttctgtgctctttgtgaatcgctgacctctctatc  
. . . . .  
tccgtgaaaagagcacgttcttctgctgtatgtaacctgtcttttctatgatctcttag  
|5161 |5171 |5181 |5191 . . .  
GGGTGACCCAGTCTATTAAAGAAAGAAAAATGCTGAATGAGgtaagtacttgatgttaca  
V T Q S I K E R K M L N E  
|1721 |1731  
. . . . .  
aactaaccagagatattcattcagtcatatagttaaaaatgtatttgcttccttccatca  
. . . . .  
atgcaccactttccttaacaatgcacaaatcttccatgataatgaggatcatcaagaatt  
. . . . .  
atgcaggcctgcactgtgggtcatacctataatcccagcgctttgggaggctgaggcgct  
. . . . .  
tggatcacctgatgtcgggagttcaagaccagcctgaccaacatggagaaacccgtttc  
. . . . .  
tactaaaaatacaaaattagccgggcttggtggcacttgcc

Exon 19 | Start: 160848 | End: 160932 | Length: 84

```
. . . . .
gcattcaaaagattctcctgcctcagcctcccaagtagctgggattacaggtgcctgcc
. . . . .
ccacgccaactaatttttgtatttttagtagagatgaggtttcaccatgttggtcaga
. . . . .
ctggtgtcgaactcctgacctcaagtgatctgcctgcctcagtcctccaaagtgctagga
. . . . .
ttacaggggtgagccactgcgctggcctgaatgccttaaatacgactgtctgctccac
. . . . .
ttcattgaaggaagcttctctttctcttatcctgatgggtgtgtttggtttctttcag
      |5201      |5211      |5221      |5231      |5241      |5251
CATGATTTTGAAGTCAGAGGAGATGTGGTCAATGGAAGAAACCAAGGTCCAAAGCGA
H D F E V R G D V V N G R N H Q G P K R
                        |1741                        |1751

      |5261      |5271      . . . . .
GCAAGAGAATCCCAGGACAGAAAGgtaaagctccctccctcaagttgacaaaaatctcac
A R E S Q D R K

. . . . .
cccaccactctgtattccactcccctttgcagagatgggccgcttcattttgtaagactt
. . . . .
attacatacacacagtgtctagatactttcacacaggttcttttttactcttccatcc
. . . . .
caaccacataaataagtattgtctctactttatgaatgataaaactaagagatttagaga
. . . . .
ggctgtgtaatattggattcccgtctcgggttcagatcttagctgataagtggaagagctg
. . . . .
ggactttaagcagatgagaatcta
```

Exon 20 | Start: 166866 | End: 166921 | Length: 55

```
. . . . .
cttggcctgattggtgacaaaagtgagatgctcagtccttgaatgacaaagaatgcctgt
. . . . .
agagtgcagggtcaactacatatgcacttcaagaagatcttctgaaatccagtagtgttct
. . . . .
ggacattggactgcttgctccctgggaagtagcagcagaaatcatcaggtggtgaacagaa
. . . . .
gaaaaagaaaagctcttcctttttgaaagtctgttttttgaataaaagccaatattcttt
. . . . .
tataactagattttccttctctccattcccctgtccctctctcttcctctcttcttcag
|5281 |5291 |5301 |5311 |5321 |5331 .
ATCTTCAGGGGGCTAGAAATCTGTTGCTATGGGCCCTTCACCAACATGCCACAGgtaag
I F R G L E I C C Y G P F T N M P T D
|1761 |1771
. . . . .
agcctgggagaaccccagagttccagcaccagcctttgtcttacatagtgaggattattata
. . . . .
agcaagatcccacgatgggggttcctcagattgctgaaatgttctagaggctattctatt
. . . . .
tctctaccactctccaaacaaaacagcacctaaatgttatcctatggcaaaaaaaacta
. . . . .
taccttgcccccttctcaagagcatgaagggtggttaatagttaggattcagtatgttat
. . . . .
gtgttcagatggcggttgagctgctgttagtgccaacatgttagtgagaaaaatc
```

Exon 21 | Start: 168789 | End: 168863 | Length: 74

. . . . .  
 attaatggaaattagatctttgattttttttctttcaagcattttatttgagagactat  
 . . . . .  
 caaaccttataccaagtggccttatggagactgataaccagagtacatggcatatcagtg  
 . . . . .  
 gcaaattgacttaaaatccatacccctactattttaagaccattgtcctttggagcagag  
 . . . . .  
 agacagactctccattgagaggtcttgctataagccttcacccggagagtgtagggtag  
 . . . . .  
 agggcctgggttaagtatgcagattactgcagtgattttacatctaaatgtccatttttag  
 . . . . .  
 |5341 |5351 |5361 |5371 |5381 |5391  
 ATCAACTGGAATGGATGGTACAGCTGTGTGGTGCTTCTGTGGTGAAGGAGCTTTCATCAT  
 Q L E W M V Q L C G A S V V K E L S S F  
 |1781 |1791  
 . . . . .  
 |5401  
 TCACCCTTGGCACAgtaagtattgggtgccctgtcagagagggaggacacaatattctct  
 T L G T  
 |1801  
 . . . . .  
 cctgtgagcaagactggcacctgtcagtccttatggatgcccctactgtagcctcagaag  
 . . . . .  
 tcttctctgcccacatacctgtgccaaaagactccatctgtaagggatgggtaaggattt  
 . . . . .  
 gagaactgcacatattaaatatactgagggaagactttttccctctaactctttttccca  
 . . . . .  
 tatgtccctccccctcctctctgtgactgccccagcatactgtgtttcaacaaatcatca  
 . . . . .  
 agaaatgatgggct

Exon 22 | Start: 170280 | End: 170341 | Length: 61

. . . . .  
 ccaggagtttgagaccagcctgggcaacatggcaaaaccctgtctctacaaaaatacaa  
 . . . . .  
 aaaaattagccaggggtggtggtacgtgtctgtagttccagctacttaggaggctgagat  
 . . . . .  
 ggaaggattgcttgagcccaggaggcagaggtggcagtgagctgagatcacaccactgca  
 . . . . .  
 ctccagcctgggtgacagagcaagaccctgtctcaaaaacaaacaaaaaatgatgaag  
 . . . . .  
 tgacagttccagtagtcctactttgacactttgaatgctcttccttcctggggatccag  
 |5411 |5421 |5431 |5441 |5451 |5461  
 GGTGTCCACCCAATTGTGGTTGTGCAGCCAGATGCCTGGACAGAGGACAATGGCTTCCAT  
 G V H P I V V V Q P D A W T E D N G F H  
 |1811 |1821  
 . . . . .  
 Ggtaaggctgcctgcatgtacctgtgctatatggggtccttttgcattgggtttggtttatc  
 A  
 . . . . .  
 actcattacctggtgcttgagtagcacagttcttggcacattttaaatatttggtgaatg  
 . . . . .  
 aatggctaaaatgtctttttgatgtttttattgttatttgttttatattgtaaaagtaat  
 . . . . .  
 acatgaactgtttccatgggggtgggagtaagatatgaatgttcacacaaaaacataaat  
 . . . . .  
 caaggccgggcatggtggctcatgcctataattccagcactttgggaggtcaagatggag  
 .  
 g

Exon 23 | Start: 172181 | End: 173689 | Length: 1508

. . . . .  
 taaaaatacaaaaattagctgggtgtgatggcatgtgcctgtaattccagctactcagga  
 . . . . .  
 ggcagagacaggagaattgcttgaacccaggaggcggaggtgaatgagccgagattgcg  
 . . . . .  
 ccatcacactctagcctcggcgacagagcaagactccgtctcaaaaaaaaaaaaaaaaaa  
 . . . . .  
 ttagcttctacctcattaatcctaagaactcatacaaccaggaccctggagtcgattgat  
 . . . . .  
 tagagcctagtccaggagaatgaattgacactaatctctgcttgtgttctctgtctccag  
 |5471 |5481 |5491 |5501 |5511 |5521  
 CAATTGGGCAGATGTGTGAGGCACCTGTGGTGACCCGAGAGTGGGTGTTGGACAGTGTAG  
 I G Q M C E A P V V T R E W V L D S V A  
 |1831 |1841  
 |5531 |5541 |5551 |5561 |5571 |5581  
 CACTCTACCAGTGCCAGGAGCTGGACACCTACCTGATACCCAGATCCCCACAGCCACT  
 L Y Q C Q E L D T Y L I P Q I P H S H Y  
 |1851 |1861  
 |5591 |+11 |+21 |+31 |+41 |+51  
 ACTGACTGCAGCCAGCCACAGGTACAGAGCCACAGGACCCCAAGAATGAGCTTACAAAGT  
 \*  
 |+61 |+71 |+81 |+91 |+101 |+111  
 GGCCTTTCCAGGCCCTGGGAGCTCCTCTACTCTTCAGTCCTTCTACTGTCCTGGCTACT  
 |+121 |+131 |+141 |+151 |+161 |+171  
 AAATATTTTATGTACATCAGCCTGAAAAGGACTTCTGGCTATGCAAGGGTCCCTTAAAGA  
 |+181 |+191 |+201 |+211 |+221 |+231  
 TTTTCTGCTTGAAGTCTCCCTTGGAATCTGCCATGAGCACAAAATTATGGTAATTTTTC  
 |+241 |+251 |+261 |+271 |+281 |+291  
 ACCTGAGAAGATTTTAAAACCATTTAAACGCCACCAATTGAGCAAGATGCTGATTCATTA  
 |+301 |+311 |+321 |+331 |+341 |+351  
 TTTATCAGCCCTATTCTTTCTATTGAGGCTGTTGTTGGCTTAGGGCTGGAAGCACAGAGT  
 |+361 |+371 |+381 |+391 |+401 |+411

GGCTTGGCCTCAAGAGAATAGCTGGTTTCCCTAAGTTTACTTCTCTAAAACCTGTGTTC

      |+421      |+431      |+441      |+451      |+461      |+471  
ACAAAGGCAGAGAGTCAGACCCTTCAATGGAAGGAGAGTGCTTGGGATCGATTATGTGAC

      |+481      |+491      |+501      |+511      |+521      |+531  
TTAAAGTCAGAATAGTCCTTGGGCAGTTCTCAAATGTTGGAGTGGAACATTGGGGAGGAA

      |+541      |+551      |+561      |+571      |+581      |+591  
ATTCTGAGGCAGGTATTAGAAATGAAAAGGAACTTGAAACCTGGGCATGGTGGCTCACG

      |+601      |+611      |+621      |+631      |+641      |+651  
CCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGGGCAGATCACTGGAGGTCAGGAGTTCTG

      |+661      |+671      |+681      |+691      |+701      |+711  
AAACCAGCCTGGCCAACATGGTGAAACCCCATCTCTACTAAAAATACAGAAATTAGCCGG

      |+721      |+731      |+741      |+751      |+761      |+771  
TCATGGTGGTGGACACCTGTAATCCCAGCTACTCAGGTGGCTAAGGCAGGAGAATCACTT

      |+781      |+791      |+801      |+811      |+821      |+831  
CAGCCCGGAGGTGGAGGTTGCAGTGAGCCAAGATCATACCACGGCACTCCAGCCTGGGT

      |+841      |+851      |+861      |+871      |+881      |+891  
GACAGTGAGACTGTGGCTCAAAAAAAAAAAAAAAAAAGGAAAATGAACTAGAAGAGAT

      |+901      |+911      |+921      |+931      |+941      |+951  
TTCTAAAAGTCTGAGATATATTTGCTAGATTTCTAAAGAATGTGTTCTAAAACAGCAGAA

      |+961      |+971      |+981      |+991      |+1001     |+1011  
GATTTTCAAGAACCGTTTCCAAAGACAGTCTTCTAATTCCTCATTAGTAATAAGTAAAA

      |+1021     |+1031     |+1041     |+1051     |+1061     |+1071  
TGTTTATTGTTGTAGCTCTGGTATATAATCCATTCTCTTAAAAATATAAGACCTCTGGCA

      |+1081     |+1091     |+1101     |+1111     |+1121     |+1131  
TGAATATTTTCATATCTATAAAATGACAGATCCCACCAGGAAGGAAGCTGTTGCTTTCTTT

      |+1141     |+1151     |+1161     |+1171     |+1181     |+1191  
GAGGTGATTTTTTTCCTTTGCTCCCTGTTGCTGAAACCATACAGCTTCATAAATAATTTT

      |+1201     |+1211     |+1221     |+1231     |+1241     |+1251  
GCTTGCTGAAGGAAGAAAAAGTGTTTTTCATAAACCCATTATCCAGGACTGTTTATAGCT

      |+1261     |+1271     |+1281     |+1291     |+1301     |+1311  
GTTGGAAGGACTAGGTCTTCCCTAGCCCCCAGTGTCAGGGCAGTGAAGACTTGATT

```

      |+1321      |+1331      |+1341      |+1351      |+1361      |+1371
GTACAAAATACGTTTTGTAAATGTTGTGCTGTTAACTGCAAATAAACTGGTAGCAAA

      |+1381 . . . . .
CACTTCCAccatgaatgactgttcttgagacttaggccagccgactttctcagagccttt

. . . . .
tcactgtgcttcagtctccactctgtaaaatgggggtaatgatagtatctacctcctag

. . . . .
gatttattgaggcagcttaaataccttttgatattcctgttgctgcaaaacaaattgtt

. . . . .
gcaaggtcagaagtctgaggtggctcaactgtttctttgtttcaggtttcatgaggccaa

. . . . .
aataaagtggttcgcagggcgtgttccttctagaggctctgggtccttgagttctagg

. . .
actaagat

```

```

GBK 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

```