

Gene: BRCA2 - Sequence: NG_012772.3
 Transcript: NM_000059.3 - Protein: NP_000050.2
 Date : February 19, 2015

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

Exon 1 | Start: 5001 | End: 5188 | Length: 187

```

. . . . .
tgtcccgagggcgccacccaaacatgagctggagcaaaaagaaagggatgggggacttg

. . . . .
gagtaggcataagggcgcccccctccaagcagggtggcctgggactcttaagggtcagcga

. . . . .
gaagagaacacacactccagctcccgtttattcggtcagatactgacggttgggatgcc

. . . . .
tgacaagggaatttcctttcgccacactgagaaataccgcgagcgccccaccaggcctga

. . . . .
cttccgggtggtgctgtgctgctgctgcgtcacggcgtcacgtggccagcgcggtt

      |-219      |-209      |-199      |-189      |-179      |-169
GTGGCGCGAGCTTCTGAAACTAGGCGGCAGAGCGGAGCCGCTGTGGCACTGCTGCGCCT

      |-159      |-149      |-139      |-129      |-119      |-109
CTGCTGCGCCTCGGGTGTCTTTTGCGGCGGTGGGTGCGCGCCGGGAGAAGCGTGAGGGGA

      |-99       |-89       |-79       |-69       |-59       |-49
CAGATTTGTGACCGGCGCGGTTTTTGTGAGCTTACTCCGGCCAAAAAAGAACTGCACCTC

TGGAGCGGgtagtggtggtggttagtgggttgggacgagcgcgctttccgcagtcccagt

. . . . .
ccagcgtggcgggggagcgcctcacgccccgggtcgctgccgaggcttcttgcccttttg

. . . . .

```


agtacgtcacagtgttgcttagaaccataaaactgttccttatgtgtgtataaatccagtt

 aacaacataatcatcgtttgaggttaaccacatgataaatatagaacgtctagtggata

 aagaggaaactggccccttgactagcagtaggaacaattactaacaacatcagaagcatta

 atgttactttatggcagaagttgtccaactttttggtttcagtact

Exon 3 | Start: 8598 | End: 8846 | Length: 248

.
 ttgattccgaacctgaataataagaaaataaaatataaatgaaaaataaaagctaa

 tatatacagcttttaataatatagttaaataatgccatcttgtaacttttgtaactcttggt

 acacctttctatagattcgcaagagaatggattaatgatcttgtttaataatatgcctt

 aacaaaagtaatccatagtcagatcttaagcatttttttccttatgatctttaactgtt

 ctgggtcacaaatttgctgtcactgggttaaaactaagggtgggatttttttttaaatag

71	81	91	101	111	121
ATTTAGGACCAATAAGTCTTAATTGGTTTGAAGAACTTTCTTCAGAAGCTCCACCCTATA					
L	G	P	I	S	L
			N	W	F
			E	E	L
			S	S	E
			A	P	P
			Y	N	
		31			41

131	141	151	161	171	181
ATTCTGAACCTGCAGAAGAATCTGAACATAAAAAACAACAATTACGAACCAAACCTATTTA					
S	E	P	A	E	E
		S	E	H	K
		N	N	N	Y
		E	P	N	L
		F	K		
		51			61

191	201	211	221	231	241
AAACTCCACAAAGGAAACCATCTTATAATCAGCTGGCTTCAACTCCAATAATATTCAAAG					
T	P	Q	R	K	P
		S	Y	N	Q
		L	A	S	T
		P	I	I	F
		K	E		
		71			81

|251 |261 |271 |281 |291 |301
AGCAAGGGCTGACTCTGCCGCTGTACCAATCTCCTGTAAAAGAATTAGATAAATTCAAAT
Q G L T L P L Y Q S P V K E L D K F K L
|91 |101

|311
 TAGACTTAGgtaagtaatgcaatatggtagactgggggagaactacaaactaggaatttag
 D L G

gcaaacctgtgttaaaatcttagctcattcattaattgtgtcatgctgggcaaatcagtc
tctctggcctctttttcctcactcgaaaaatggagacgatgaaaataatgtctcataggt
ttggattaaattaataatgttaggtacttagtaaatgttctctttcatccttcctttgat
aaatttgccaactgagatttgctgaattacgtcttttctatgccaaaaaacctaggact
tgttttgat

Exon 4 | Start: 14597 | End: 14705 | Length: 108

caccacaccagcgttttctttgtagaggcagagtctcactctgttgctcaggcaggtgtt
gaactcctgcctcaagcaatcctcccacctcagcctcccagagccctcaaattataagcc
actgtgctcggggcatcctttttggggggtaatcagcaaactgaaaaacctcttcttaca
actccctatacattctcattcccagtatagaggagactttttgtttttaaacatttcaa
agaatgcaaatttataatccagagtatatacattctcactgaattattgtactgtttcag

AAGCAGATGATGTTTCCTGTCCACTTCTAAATTCTTGCTTAGTGAAA Ggtatgatgaag
A D D V S C P L L N S C L S E S

ctatttatattaaaatattttaaatgaaacattttcctacatatatttggttctataaagatg
aatctgatttttatgctaataattttggctaagagcctggtagaatcttacatttttaa
ataatcttttaggttgagtcctttaatagaatagttttacattagaacatgtaagttg
ttgttcttgatggtgaattggctggttttctgtatattctgtgatttttaagtaaca
aaaataacagtgggtgaaaagcagtaagtcagtccttgaattatcaattt

Exon 5 | Start: 15622 | End: 15671 | Length: 49
BE AWARE: This section overlaps with the following exon

gactgctgggattacaggcgtgaaccactgtgtcccggcctactttacaaaatttttgagt
ttaaatacacggtttccagcagctgaaatttgtgagtacatatgtgttggcattttaaa
catcacttgatgattatttaatgcttcatgagagatttactttttaaaatgtaataaaa
atatctaaaagtagtattccaacaatttatatgaatgagaatcttcttttaaaaataaga

taaactagtttttgccagttttttaaaataacctaagggatttgctttgttttatttttag

 |431 |441 |451 |461 |471 .< .
TCCTGTTGTTCTACAATGTACACATGTAACACCACAAAGAGATAAGTCAGgtatgattaa
P V V L Q C T H V T P Q R D K S V
 |151

. .
aaacaatgctttttatttcttagaataactagaaatgtaataaaaaataaaaacttaacaatt

. .
ttcccctttttttacccccagtggtatgtgggagtttgtttcatacaccaaagtttgta

. .
aggtaaatattctacctgggtttatttttatgacttagtaattgagaatttgacaatagcg

. .
ttatacctttgccctgagatttacaaatctgtacctagcattctgcctcatcacaggcaat

. .
tcagtaaacggttaagtgaataaagagtgaatgaaaaataaatatcctta

Exon 6 | Start: 15763 | End: 15803 | Length: 40
BE AWARE: This section overlaps with the following exon

. .
tgcttcatgagagatttactttttaaaatgtaataataaaatatctaaaagtagtattcca

. .
acaatttatatgaatgagaatcttcttttaaaaataagataaaactagtttttgccagttt

. .
tttaaaataacctaagggatttgctttgttttatttttagtcctgttggttctacaatgtac

. .
acatgtaacaccacaaagagataagtcaggtatgattaaaaacaatgctttttatttctta

. .
gaataactagaaatgtaataaaaaataaaaacttaacaattttcccctttttttacccccag

 |481 |491 |501 |511 .< .
TGGTATGTGGGAGTTTGTTCATACACCAAAGTTTGTGAAGgtaaattattctacctggtt

V C G S L F H T P K F V K
 |161 |171

.
 tatttttatgacttagtaattgagaatttgacaatagcgttatacctttgccctgagatt

 taaaaatctgtacctagcattctgcctcatcacaggcaattcagtaaacgttaagtgaat

 aaagagtgaatgaaaaataatatccttaatgatcagggcatttctataaaaaataaact

 attttcttctcctcccagggctcgtcagacaccaaataatatttctgaaagtctaggagctg

 aggtgatcctgatatgtcttgggtcaagttctttagctaca

Exon 7 | Start: 16020 | End: 16134 | Length: 114

.
 ataaaaataaaacttaacaattttcccctttttttacccccagtggtatgtgggagtttg

 tttcatcaccaaagtttgtgaaggtaaataattctacctggtttatttttatgacttagt

 aattgagaatttgacaatagcgttatacctttgccctgagatttacaatctgtacctag

 cattctgcctcatcacaggcaattcagtaaacgttaagtgaataaagagtgaatgaaaa

 ataatatccttaatgatcagggcatttctataaaaaataaactattttcttctcctccag

|521 |531 |541 |551 |561 |571
 GGTTCGTCAGACACCAAAACATATTTCTGAAAGTCTAGGAGCTGAGGTGGATCCTGATATG
 G R Q T P K H I S E S L G A E V D P D M
 |181 |191

|581 |591 |601 |611 |621 |631 .
 TCTTGGTCAAGTTCTTTAGCTACACCACCCACCTTAGTTCTACTGTGCTCATAGgtaaat

S W S S S L A T P P T L S S T V L I V
 |201 |211

.
 aatagcaaatgtgtattttacaagaaagagcagatgaggttgataattgtcatctctaata

 cttctgttaaaaggaaatatgaaaagaaaatattagataatgtctttgataagtgtgtta

 gtaactgacaataatTTTtattctattaagtgtagattggaataaatacaatacatttag

 tggtagtccagtgggtgtcaagcattatgttttagtacgatgtgattaacgtagaatagct

 tacaaatattcctttactggcctatataagcgTTtaagaggcagtatttgggtgtg

Exon 8 | Start: 18964 | End: 19013 | Length: 49

.
 ttaagtcataacctcacagcatcatctgactttccaactcattgtggacagtattaccat

 aaagtaatgatcaccaagccatatcttaccaccttgtagtagtactaaggaagtaagta

 tagtttattcactgtgttgattgacctttctaattactatacttaagtacttgaatcaat

 tcattttgtttcaaagtgtgtcatgtaatcaaatagtatgtgctttttgatgtctgaca

 aaaaataagtttttgattctagtataataacacataaatttttatcttacag

|641 |651 |661 |671 |681 . .
 TCAGAAATGAAGAAGCATCTGAAACTGTATTTCTCATGATACTACTGCTgtaagtaa
 R N E E A S E T V F P H D T T A
 |221

.
 atgacattgattagactgttgaaattgctaacaattttggaatgccttgtaaattattt

.
 atcttacatnttttaatttcctaactctgtaatttatctaagcctttgagaaagtctctaaa

 cctggtcctatatgtgattttaacttcctgtgaaactctgctgtctctctgttaaagttg

 catatatacaatatataaccgtagtagccctattcatgggggtatacattccaatatcccca

 gtgaatgcttgaaaccttagatagtagaccgaaccctatatatatatatataa

Exon 9 | Start: 20440 | End: 20551 | Length: 111

.
 attcatgtcccaagtgggatggagcaagatgggtgcaagtttttttctccatttcatt

 ttcctttcctaagatttccacatcctagtggtgcaagatttcatcacactactcaggatg

 acacacaatttaaaacttactaattgcttacttctggaattttccattaaaaatttttgg

 acctaggttgattgcagataactgaaatcaccaaagtgaaccatggataaggggggac

 tactactatatgtgcattgagagtttttatactagtgattttaactataatttttgcag

691	701	711	721	731	741
AATGTGAAAAGCTATTTTCCAATCATGATGAAAGTCTGAAGAAAAATGATAGATTTATC					
N	V	K	S	Y	F
	S	N	H	D	E
	S	L	K	K	N
	D	R	F	I	
231		241			

751	761	771	781	791	.
GCTTCTGTGACAGACAGTGAAAACACAAATCAAAGAGAAGCTGCAAGTCATGgtaagtcc					
A	S	V	T	D	S
E	N	T	N	Q	R
E	A	A	S	H	G
251		261			

.
 tctgttttagttgaactacagggttttttgttggtgtgtgttttgatttttttttttggagg

.
 tggagtcttgctctgtcaccctgatctcggtttaccgcaacctctgcctcccgtgctca

 agcgatcctgcctcagcttgccaagtagctgagattacaagcatgcaccaccatgccc

 ctattgtatTTTTtagtagagatggcatttcacatgttggccaggctggtctcaa

 cgtgagccaccatgccagcctgaactactctTTTtaattggcaccattgaa

Exon 10 | Start: 21793 | End: 22908 | Length: 1115

.
 agtaagattaggtgagTTTTaattgtgtagaactgctaaagaaaggTTTTtagggattgt

 tgtatgaataaaaggctTTtaggttcattggaatcaggggaatcaggcttactagaagaa

 caggagaaggggtgactgaccgaaaaataaaatgccagtgactcagaataaccctTTtaa

 tactgatatgtaatatTTtagcacattctacataaactgtttctatgagaaaggTTgtgag

 aataatataaattatatggcTTataaaatattaatgtgcttctgttttatactTTtaacag

801	811	821	831	841	851
GATTTGAAAAACATCAGGGAATTCATTTAAAGTAAATAGCTGCAAAGACCACATTGGAA					
F	G	K	T	S	G
	N	S	F	K	V
			N	S	C
			K	D	H
				I	G
				K	
		271		281	

861	871	881	891	901	911
AGTCAATGCCAAATGTCCTAGAAGATGAAGTATATGAAACAGTTGTAGATACCTCTGAAG					
S	M	P	N	V	L
	E	D	E	V	Y
			E	T	V
			V	D	T
				S	E
				E	
		291		301	

921	931	941	951	961	971
AAGATAGTTTTTCATTATGTTTTTCTAAATGTAGAACAAAAATCTACAAAAAGTAAGAA					

D S F S L C F S K C R T K N L Q K V R T
|311 |321
|981 |991 |1001 |1011 |1021 |1031
CTAGCAAGACTAGGAAAAAATTTCCATGAAGCAAACGCTGATGAATGTGAAAAATCTA
S K T R K K I F H E A N A D E C E K S K
|331 |341
|1041 |1051 |1061 |1071 |1081 |1091
AAAACCAAGTGAAAGAAAAATACTCATTGTATCTGAAGTGAACCAAATGATACTGATC
N Q V K E K Y S F V S E V E P N D T D P
|351 |361
|1101 |1111 |1121 |1131 |1141 |1151
CATTAGATTCAAATGTAGCAAATCAGAAGCCCTTTGAGAGTGAAGTGACAAAATCTCCA
L D S N V A N Q K P F E S G S D K I S K
|371 |381
|1161 |1171 |1181 |1191 |1201 |1211
AGGAAGTTGTACCGTCTTTGGCCTGTGAATGGTCTCAACTAACCCTTTCAGGTCTAAATG
E V V P S L A C E W S Q L T L S G L N G
|391 |401
|1221 |1231 |1241 |1251 |1261 |1271
GAGCCCAGATGGAGAAAAATACCCCTATTGCATATTTCTTCATGTGACCAAAATATTTAG
A Q M E K I P L L H I S S C D Q N I S E
|411 |421
|1281 |1291 |1301 |1311 |1321 |1331
AAAAAGACCTATTAGACACAGAGAACAAAAGAAAGATTTTCTTACTTCAGAGAATT
K D L L D T E N K R K K D F L T S E N S
|431 |441
|1341 |1351 |1361 |1371 |1381 |1391
CTTTGCCACGTATTTCTAGCCTACCAAAATCAGAGAAGCCATTAAATGAGGAAACAGTGG
L P R I S S L P K S E K P L N E E T V V
|451 |461
|1401 |1411 |1421 |1431 |1441 |1451
TAAATAAGAGAGATGAAGAGCAGCATCTTGAATCTCATACAGACTGCATTCTTGAGTAA
N K R D E E Q H L E S H T D C I L A V K
|471 |481
|1461 |1471 |1481 |1491 |1501 |1511
AGCAGGCAATATCTGGAACCTTCTCCAGTGGCTTCTTCATTTAGGGTATCAAAAAGTCTA
Q A I S G T S P V A S S F Q G I K K S I

	491		501
1521	1531	1541	1551
1561	1571		
TATTCAGAATAAGAGAATCACCTAAAGAGACTTCAATGCAAGTTTTTCAGGTCATATGA			
F R I R E S P K E T F N A S F S G H M T			
	511		521
1581	1591	1601	1611
1621	1631		
CTGATCCAAACTTTAAAAAAGAACTGAAGCCTCTGAAAGTGGACTGGAAATACATACTG			
D P N F K K E T E A S E S G L E I H T V			
	531		541
1641	1651	1661	1671
1681	1691		
TTTGCTCACAGAAGGAGGACTCCTTATGTCCAAATTTAATTGATAATGGAAGCTGGCCAG			
C S Q K E D S L C P N L I D N G S W P A			
	551		561
1701	1711	1721	1731
1741	1751		
CCACCACCACACAGAATTCTGTAGCTTTGAAGAATGCAGGTTTAATATCCACTTTGAAAA			
T T T Q N S V A L K N A G L I S T L K K			
	571		581
1761	1771	1781	1791
1801	1811		
AGAAAACAAATAAGTTTATTTATGTATACATGATGAAACATCTTATAAAGGAAAAAAA			
K T N K F I Y A I H D E T S Y K G K K I			
	591		601
1821	1831	1841	1851
1861	1871		
TACCGAAAGACCAAAATCAGAACTAATTAAGTTCAGCCAGTTTGAAGCAAATGCTT			
P K D Q K S E L I N C S A Q F E A N A F			
	611		621
1881	1891	1901	.
			.
TTGAAGCACCCTTACATTTGCAAATGCTGATTGAGgtacctctgtcctttttttttgt			
E A P L T F A N A D S G			
	631		
.			
aaatagtacatatagttttatagatgacgattccttctgtgtttttttctgcttttttaa			
.			
atcttcatatcttatatttaaccttaggcacatctgtatacatgattgttttaggtcttt			
.			
aattaccagtgtttagaatcaggtcactcaaacatggtagataagtttgcatagtttgtg			

.
tatatccatcactcttgagacagttttattttaagttccgggtacatgtgcaggatgtg
.
caggtttgttacataagtaaactgtatgccatgttgg

Exon 11 | Start: 25786 | End: 30717 | Length: 4931

.
tcaccctcctgggctcaagcagtccttgacctcacctcctgagtaactggcaccacag
.
gcatacaccaccacaccagctaatttttatttttcatagagtcatggctcactatgtt
.
gccaggctagtctcgaactcctgggctcaagcagtcctcctgcctcagcctccaaaag
.
tgctgagattacaggcatgagccactgtgcccaaactacctttttaacttagtgaaa
.
atatttagtgaaatgtgattgatgtactttaattttgtcactttgtgttttatgtttag

1911	1921	1931	1941	1951	1961
GTTTATTGCATTCTTCTGTGAAAAGAAGCTGTTACAGAATGATTCTGAAGAACCAACTT					
L	L	H	S	S	V
		K	R	S	C
		S	Q	N	D
				S	E
				E	P
				T	L
	641			651	

1971	1981	1991	2001	2011	2021
TGTCCTTAAGTAGCTCTTTTGGGACAATTCTGAGGAAATGTTCTAGAAATGAAACATGTT					
S	L	T	S	S	F
		G	T	I	L
		R	K	C	S
			R	N	E
				T	C
				S	
	661			671	

2031	2041	2051	2061	2071	2081
CTAATAATACAGTAATCTCTCAGGATCTTGATTATAAAGAAGCAAAATGTAATAAGGAAA					
N	N	T	V	I	S
		Q	D	L	D
		Y	K	E	A
			K	C	N
				K	E
				K	
	681			691	

2091	2101	2111	2121	2131	2141
AACTACAGTTATTTATTACCCAGAAAGCTGATTCTCTGTCATGCCTGCAGGAAGGACAGT					
L	Q	L	F	I	T
		P	E	A	D
		S	L	S	C
			L	Q	E
				G	Q
				C	
	701			711	

2151	2161	2171	2181	2191	2201
GTGAAAATGATCCAAAAAGCAAAAAAGTTTCAGATATAAAAGAAGAGGTCTTGGCTGCAG					
E	N	D	P	K	S
			K	V	S
			D	I	K
			E	E	V
			L	A	A
					A
		721			731
2211	2221	2231	2241	2251	2261
CATGTCACCCAGTACAACATTCAAAAGTGGAATACAGTGATACTGACTTTCAATCCAGAG					
C	H	P	V	Q	H
			S	K	V
			E	Y	S
			D	T	D
			F	Q	S
					Q
					K
		741			751
2271	2281	2291	2301	2311	2321
AAAGTCTTTTATATGATCATGAAAATGCCAGCACTCTTATTTTAACTCCTACTTCCAAGG					
S	L	L	Y	D	H
			E	N	A
			S	T	L
			I	L	T
			P	T	S
					K
					D
		761			771
2331	2341	2351	2361	2371	2381
ATGTTCTGTCAAACCTAGTCATGATTTCTAGAGGCAAAGAATCATACAAAATGTCAGACA					
V	L	S	N	L	V
			M	I	S
			R	G	K
			E	S	Y
			K	M	S
					D
					K
		781			791
2391	2401	2411	2421	2431	2441
AGCTCAAAGGTAACAATTATGAATCTGATGTTGAATTAACCAAAAATATTCCCATGGAAA					
L	K	G	N	N	Y
			E	S	D
			V	E	L
			T	K	N
			I	P	M
					E
					K
		801			811
2451	2461	2471	2481	2491	2501
AGAATCAAGATGTATGTGCTTTAAATGAAAATTATAAAACGTTGAGCTGTTGCCACCTG					
N	Q	D	V	C	A
			L	N	E
			N	Y	K
			N	V	E
			L	L	P
					P
					E
		821			831
2511	2521	2531	2541	2551	2561
AAAAATACATGAGAGTAGCATCACCTTCAAGAAAGGTACAATTCAACCAAAACACAAATC					
K	Y	M	R	V	A
			S	P	S
			R	K	V
			Q	F	N
			Q	N	T
					N
					L
		841			851
2571	2581	2591	2601	2611	2621
TAAGAGTAATCCAAAAAATCAAGAAGAACTACTTCAATTTCAAAAATAACTGTCAATC					
R	V	I	Q	K	N
			Q	E	E
			T	T	S
			I	S	K
			I	T	V
					N
					P
		861			871
2631	2641	2651	2661	2671	2681
CAGACTCTGAAGAAGCTTTTCTCAGACAATGAGAATAATTTGTCTTCCAAGTAGCTAATG					
D	S	E	E	L	F
			S	D	N
			E	N	N
			F	V	F
			Q	V	A
					N
					E
		881			891

2691	2701	2711	2721	2731	2741
AAAGGAATAATCTTGCTTTAGGAAATACTAAGGAACTTCATGAAACAGACTTGACTTGTG					
R	N	N	L	A	L
			G	N	T
			K	E	L
			H	E	T
			D	L	T
			C	V	
	901			911	
2751	2761	2771	2781	2791	2801
TAAACGAACCCATTTTCAAGAACTCTACCATGGTTTTATATGGAGACACAGGTGATAAAC					
N	E	P	I	F	K
			N	S	T
			M	V	L
			Y	G	D
			T	G	D
			K	Q	
	921			931	
2811	2821	2831	2841	2851	2861
AAGCAACCCAAGTGTCAATTAAGAAAGATTTGGTTTATGTTCTTGCAGAGGAGACAAAA					
A	T	Q	V	S	I
			K	K	D
			L	V	Y
			V	L	A
			E	E	N
			K	N	
	941			951	
2871	2881	2891	2901	2911	2921
ATAGTGTAAGCAGCATATAAAATGACTCTAGGTCAAGATTTAAATCGGACATCTCCT					
S	V	K	Q	H	I
			K	M	T
			L	G	Q
			D	L	K
			S	D	I
			S	L	
	961			971	
2931	2941	2951	2961	2971	2981
TGAATATAGATAAAATACCAGAAAAAATAATGATTACATGAACAAATGGGCAGGACTCT					
N	I	D	K	I	P
			E	K	N
			N	D	Y
			M	N	K
			W	A	G
			L	L	
	981			991	
2991	3001	3011	3021	3031	3041
TAGGTCCAATTTCAAATCACAGTTTGGAGGTAGCTTCAGAACAGCTTCAAATAAGGAAA					
G	P	I	S	N	H
			S	F	G
			G	S	F
			R	T	A
			S	N	K
			E	I	
	1001			1011	
3051	3061	3071	3081	3091	3101
TCAAGCTCTCTGAACATAACATTAAGAAGAGCAAAATGTTCTTCAAAGATATTGAAGAAC					
K	L	S	E	H	N
			I	K	K
			S	K	M
			F	F	K
			D	I	E
			E	E	Q
	1021			1031	
3111	3121	3131	3141	3151	3161
AATATCCTACTAGTTTAGCTTGTGTTGAAATTGTAAATACCTTGGCATTAGATAATCAAAA					
Y	P	T	S	L	A
			C	V	E
			I	V	N
			T	L	A
			L	D	N
			Q	K	
	1041			1051	
3171	3181	3191	3201	3211	3221
AGAAACTGAGCAAGCCTCAGTCAATTAATACTGTATCTGCACATTTACAGAGTAGTGTAG					
K	L	S	K	P	Q
			S	I	N
			T	V	S
			A	H	L
			Q	S	S
			V	V	
	1061			1071	
3231	3241	3251	3261	3271	3281

TTGTTTCTGATTGTAAAAATAGTCATATAACCCCTCAGATGTTATTTTCCAAGCAGGATT
 V S D C K N S H I T P Q M L F S K Q D F
 |1081 |1091

|3291 |3301 |3311 |3321 |3331 |3341
 TTAATTCAAACCATAATTTAACACCTAGCCAAAAGGCAGAAATTACAGAACTTTCTACTA
 N S N H N L T P S Q K A E I T E L S T I
 |1101 |1111

|3351 |3361 |3371 |3381 |3391 |3401
 TATTAGAAGAATCAGGAAGTCAGTTTGAATTTACTCAGTTTAGAAAACCAAGCTACATAT
 L E E S G S Q F E F T Q F R K P S Y I L
 |1121 |1131

|3411 |3421 |3431 |3441 |3451 |3461
 TGCAGAAGAGTACATTTGAAGTGCCTGAAAACCAGATGACTATCTTAAAGACCACTTCTG
 Q K S T F E V P E N Q M T I L K T T S E
 |1141 |1151

|3471 |3481 |3491 |3501 |3511 |3521
 AGGAATGCAGAGATGCTGATCTTCATGTCATAATGAATGCCCCATCGATTGGTCAGGTAG
 E C R D A D L H V I M N A P S I G Q V D
 |1161 |1171

|3531 |3541 |3551 |3561 |3571 |3581
 ACAGCAGCAAGCAATTTGAAGGTACAGTTGAAATTAACGGAAGTTTGCTGGCCTGTTGA
 S S K Q F E G T V E I K R K F A G L L K
 |1181 |1191

|3591 |3601 |3611 |3621 |3631 |3641
 AAAATGACTGTAACAAAAGTGCTTCTGTTTATTTAACAGATGAAAATGAAGTGGGGTTTA
 N D C N K S A S G Y L T D E N E V G F R
 |1201 |1211

|3651 |3661 |3671 |3681 |3691 |3701
 GGGGCTTTTATTCTGCTCATGGCACAAAACCTGAATGTTTCTACTGAAGCTCTGCAAAAAG
 G F Y S A H G T K L N V S T E A L Q K A
 |1221 |1231

|3711 |3721 |3731 |3741 |3751 |3761
 CTGTGAAACTGTTTAGTGATATTGAGAATATTAGTGAGGAACTTCTGCAGAGGTACATC
 V K L F S D I E N I S E E T S A E V H P
 |1241 |1251

|3771 |3781 |3791 |3801 |3811 |3821
 CAATAAGTTTATCTTCAAGTAAATGTCATGATTCTGTTGTTTCAATGTTTAAGATAGAAA

I S L S S S K C H D S V V S M F K I E N
 |1261 |1271

 |3831 |3841 |3851 |3861 |3871 |3881
 ATCATAATGATAAACTGTAAGTGAAAAAATAATAAATGCCAACTGATATTACAAAATA
 H N D K T V S E K N N K C Q L I L Q N N
 |1281 |1291

 |3891 |3901 |3911 |3921 |3931 |3941
 ATATTGAAATGACTACTGGCACTTTTGTGAAGAAATTACTGAAAATTACAAGAGAAAATA
 I E M T T G T F V E E I T E N Y K R N T
 |1301 |1311

 |3951 |3961 |3971 |3981 |3991 |4001
 CTGAAAATGAAGATAACAAATATACTGCTGCCAGTAGAAATTCTCATAACTTAGAATTG
 E N E D N K Y T A A S R N S H N L E F D
 |1321 |1331

 |4011 |4021 |4031 |4041 |4051 |4061
 ATGGCAGTGATTCAAGTAAAAATGATACTGTTGTATTTCATAAAGATGAAACGGACTTGC
 G S D S S K N D T V C I H K D E T D L L
 |1341 |1351

 |4071 |4081 |4091 |4101 |4111 |4121
 TATTTACTGATCAGCACAAACATATGTCTTAAATTATCTGGCCAGTTTATGAAGGAGGGAA
 F T D Q H N I C L K L S G Q F M K E G N
 |1361 |1371

 |4131 |4141 |4151 |4161 |4171 |4181
 ACACTCAGATTAAAGAAGATTTGTGAGATTTAACTTTTTTGAAGTTGCGAAAGCTCAAG
 T Q I K E D L S D L T F L E V A K A Q E
 |1381 |1391

 |4191 |4201 |4211 |4221 |4231 |4241
 AAGCATGTCATGGTAATACTTCAAATAAAGAACAGTTAACTGCTACTAAAACGGAGCAAA
 A C H G N T S N K E Q L T A T K T E Q N
 |1401 |1411

 |4251 |4261 |4271 |4281 |4291 |4301
 ATATAAAGATTTTGAGACTTCTGATACATTTTTTCAGACTGCAAGTGGGAAAAATATTA
 I K D F E T S D T F F Q T A S G K N I S
 |1421 |1431

 |4311 |4321 |4331 |4341 |4351 |4361
 GTGTCGCCAAAGAGTCATTTAATAAAAATTGTAAATTTCTTTGATCAGAAACCAGAAGAAT
 V A K E S F N K I V N F F D Q K P E E L

	1441		1451
4371	4381	4391	4401
4411	4421		
TGCATAACTTTTCCTTAAATTCTGAATTACATTCTGACATAAGAAAGAACAAAATGGACA			
H N F S L N S E L H S D I R K N K M D I			
	1461		1471
4431	4441	4451	4461
4471	4481		
TTCTAAGTTATGAGGAAACAGACATAGTTAAACACAAAATACTGAAAGAAAAGTGTCCCAG			
L S Y E E T D I V K H K I L K E S V P V			
	1481		1491
4491	4501	4511	4521
4531	4541		
TTGGTACTGGAAATCAACTAGTGACCTTCCAGGGACAACCCGAACGTGATGAAAAGATCA			
G T G N Q L V T F Q G Q P E R D E K I K			
	1501		1511
4551	4561	4571	4581
4591	4601		
AAGAACCTACTCTATTGGGTTTTTCATACAGCTAGCGGAAAAAAGTTAAAATTGCAAAGG			
E P T L L G F H T A S G K K V K I A K E			
	1521		1531
4611	4621	4631	4641
4651	4661		
AATCTTTGGACAAAGTGAAAAACCTTTTTGATGAAAAAGAGCAAGGTACTAGTGAAATCA			
S L D K V K N L F D E K E Q G T S E I T			
	1541		1551
4671	4681	4691	4701
4711	4721		
CCAGTTTTAGCCATCAATGGGCAAAGACCCTAAAGTACAGAGAGGCCTGTAAAGACCTTG			
S F S H Q W A K T L K Y R E A C K D L E			
	1561		1571
4731	4741	4751	4761
4771	4781		
AATTAGCATGTGAGACCATTGAGATCACAGCTGCCCCAAAGTGTAAGAAATGCAGAATT			
L A C E T I E I T A A P K C K E M Q N S			
	1581		1591
4791	4801	4811	4821
4831	4841		
CTCTCAATAATGATAAAAAACCTTGTCTATTGAGACTGTGGTGCCACCTAAGCTCTTAA			
L N N D K N L V S I E T V V P P K L L S			
	1601		1611
4851	4861	4871	4881
4891	4901		
GTGATAATTTATGTAGACAAACTGAAAATCTCAAAACATCAAAAAGTATCTTTTTGAAAG			
D N L C R Q T E N L K T S K S I F L K V			
	1621		1631

|4911 |4921 |4931 |4941 |4951 |4961
 TTAAAGTACATGAAAATGTAGAAAAAGAAACAGCAAAAAGTCCTGCAACTTGTTACACAA
 K V H E N V E K E T A K S P A T C Y T N
 |1641 |1651

|4971 |4981 |4991 |5001 |5011 |5021
 ATCAGTCCCCTTATTTCAGTCATTGAAAATTCAGCCTTAGCTTTTACACAAGTTGTAGTA
 Q S P Y S V I E N S A L A F Y T S C S R
 |1661 |1671

|5031 |5041 |5051 |5061 |5071 |5081
 GAAAACTTCTGTGAGTCAGACTTCATTACTTGAAGCAAAAAAATGGCTTAGAGAAGGAA
 K T S V S Q T S L L E A K K W L R E G I
 |1681 |1691

|5091 |5101 |5111 |5121 |5131 |5141
 TATTTGATGGTCAACCAGAAAGAATAAATACTGCAGATTATGTAGGAAATTATTTGTATG
 F D G Q P E R I N T A D Y V G N Y L Y E
 |1701 |1711

|5151 |5161 |5171 |5181 |5191 |5201
 AAAATAATTCAAACAGTACTATAGCTGAAAATGACAAAAATCATCTCTCCGAAAAACAAG
 N N S N S T I A E N D K N H L S E K Q D
 |1721 |1731

|5211 |5221 |5231 |5241 |5251 |5261
 ATACTTATTTAAGTAACAGTAGCATGTCTAACAGCTATTCTCTACCATTCTGATGAGGTAT
 T Y L S N S S M S N S Y S Y H S D E V Y
 |1741 |1751

|5271 |5281 |5291 |5301 |5311 |5321
 ATAATGATTCAGGATATCTCTCAAAAAATAAACTTGATTCTGGTATTGAGCCAGTATTGA
 N D S G Y L S K N K L D S G I E P V L K
 |1761 |1771

|5331 |5341 |5351 |5361 |5371 |5381
 AGAATGTTGAAGATCAAAAAAACACTAGTTTTTCCAAAGTAATATCCAATGTAAAAGATG
 N V E D Q K N T S F S K V I S N V K D A
 |1781 |1791

|5391 |5401 |5411 |5421 |5431 |5441
 CAAATGCATACCCACAAACTGTAAATGAAGATATTTGCGTTGAGGAACTTGTGACTAGCT
 N A Y P Q T V N E D I C V E E L V T S S
 |1801 |1811

|5451 |5461 |5471 |5481 |5491 |5501
 CTTCACCCTGCAAAAATAAAAATGCAGCCATTAAATTGTCCATATCTAATAGTAATAATT
 S P C K N K N A A I K L S I S N S N N F
 |1821 |1831

|5511 |5521 |5531 |5541 |5551 |5561
 TTGAGGTAGGGCCACCTGCATTTAGGATAGCCAGTGGTAAAAATCGTTTGTGTTTCACATG
 E V G P P A F R I A S G K I V C V S H E
 |1841 |1851

|5571 |5581 |5591 |5601 |5611 |5621
 AAACAATTAAAAAAGTGAAAGACATATTTACAGACAGTTTCAGTAAAGTAATTAAGGAAA
 T I K K V K D I F T D S F S K V I K E N
 |1861 |1871

|5631 |5641 |5651 |5661 |5671 |5681
 ACAACGAGAATAAATCAAAAATTTGCCAAACGAAAATTATGGCAGGTTGTACGAGGCAT
 N E N K S K I C Q T K I M A G C Y E A L
 |1881 |1891

|5691 |5701 |5711 |5721 |5731 |5741
 TGGATGATTCAGAGGATATTCTTCATAACTCTCTAGATAATGATGAATGTAGCACGCATT
 D D S E D I L H N S L D N D E C S T H S
 |1901 |1911

|5751 |5761 |5771 |5781 |5791 |5801
 CACATAAGGTTTTTGCTGACATTGAGAGTGAAGAAATTTTACAACATAACCAAAATATGT
 H K V F A D I Q S E E I L Q H N Q N M S
 |1921 |1931

|5811 |5821 |5831 |5841 |5851 |5861
 CTGGATTGGAGAAAGTTTCTAAAATATCACCTTGTGATGTTAGTTTGGAAACTTCAGATA
 G L E K V S K I S P C D V S L E T S D I
 |1941 |1951

|5871 |5881 |5891 |5901 |5911 |5921
 TATGTAAATGTAGTATAGGGAAGCTTCATAAGTCAGTCTCATCTGCAAATACTTGTGGGA
 C K C S I G K L H K S V S S A N T C G I
 |1961 |1971

|5931 |5941 |5951 |5961 |5971 |5981
 TTTTTCAGCACAGCAAGTGGAATCTGTCCAGGTATCAGATGCTTCATTACAAAACGCAA
 F S T A S G K S V Q V S D A S L Q N A R
 |1981 |1991

|5991 |6001 |6011 |6021 |6031 |6041

GACAAGTGTTTTCTGAAATAGAAGATAGTACCAAGCAAGTCTTTTCCAAAGTATTGTTTA
Q V F S E I E D S T K Q V F S K V L F K
|2001 |2011
|6051 |6061 |6071 |6081 |6091 |6101
AAAGTAACGAACATTAGACCAGCTCACAAGAGAAGAAAATACTGCTATACGTACTCCAG
S N E H S D Q L T R E E N T A I R T P E
|2021 |2031
|6111 |6121 |6131 |6141 |6151 |6161
AACATTTAATATCCCAAAAGGCTTTTCATATAATGTGGTAAATTCATCTGCTTTCTCTG
H L I S Q K G F S Y N V V N S S A F S G
|2041 |2051
|6171 |6181 |6191 |6201 |6211 |6221
GATTTAGTACAGCAAGTGGAAAGCAAGTTTCCATTTTAGAAAGTTCCTTACACAAAGTTA
F S T A S G K Q V S I L E S S L H K V K
|2061 |2071
|6231 |6241 |6251 |6261 |6271 |6281
AGGGAGTGTTAGAGGAATTTGATTTAATCAGAACTGAGCATAGTCTTCACTATTACCTA
G V L E E F D L I R T E H S L H Y S P T
|2081 |2091
|6291 |6301 |6311 |6321 |6331 |6341
CGTCTAGACAAAATGTATCAAAAATACTTCCTCGTGTGATAAGAGAAACCCAGAGCACT
S R Q N V S K I L P R V D K R N P E H C
|2101 |2111
|6351 |6361 |6371 |6381 |6391 |6401
GTGTAAACTCAGAAATGGAAAAACCTGCAGTAAAGAATTTAAATTATCAAAATAACTTAA
V N S E M E K T C S K E F K L S N N L N
|2121 |2131
|6411 |6421 |6431 |6441 |6451 |6461
ATGTTGAAGGTGGTTCTTCAGAAAATAATCACTCTATTAAAGTTTCTCCATATCTCTCTC
V E G G S S E N N H S I K V S P Y L S Q
|2141 |2151
|6471 |6481 |6491 |6501 |6511 |6521
AATTTCAACAAGACAAACAACAGTTGGTATTAGGAACCAAAGTGTCACCTGTTGAGAACA
F Q Q D K Q Q L V L G T K V S L V E N I
|2161 |2171
|6531 |6541 |6551 |6561 |6571 |6581
TTCATGTTTTGGGAAAAGAACAGGCTTCACCTAAAAACGTAAAAATGGAAATTGGTAAAA

H V L G K E Q A S P K N V K M E I G K T
|2181 |2191
|6591 |6601 |6611 |6621 |6631 |6641
CTGAAACTTTTTCTGATGTTCTGTGAAAACAAATATAGAAGTTTGTCTACTTACTCCA
E T F S D V P V K T N I E V C S T Y S K
|2201 |2211
|6651 |6661 |6671 |6681 |6691 |6701
AAGATTGAGAAAACACTTTTGAACAGAGCAGTAGAAATTGCTAAAGCTTTTATGGAAG
D S E N Y F E T E A V E I A K A F M E D
|2221 |2231
|6711 |6721 |6731 |6741 |6751 |6761
ATGATGAACTGACAGATTCTAAACTGCCAAGTCATGCCACACATTCTCTTTTACATGTC
D E L T D S K L P S H A T H S L F T C P
|2241 |2251
|6771 |6781 |6791 |6801 |6811 |6821
CCGAAAATGAGGAAATGGTTTTGTCAAATTCAAGAATTGGAAAAAGAAGAGGAGAGCCCC
E N E E M V L S N S R I G K R R G E P L
|2261 |2271
|6831 |6841.
TTATCTTAGTGgtaagtggttcatttttacctttcgtgttgccaatcactattttaaag
I L V G
|2281
.
tgtttattcagtagacttggtatgctaacaattaagagtggtataaactatgtcttttca
.
gccatttttgtgtagtcagtttgggggagtgatggtttgatatacagatacacagattcag
.
tattcgtatacagatttgatatcttggtatacagattcgatatctctgaatctgtatacc
.
aagaaatcatgttttaagggctcaatatattttcaaaaagattattagtagtataataattg
. . . .
agaaattactgt

Exon 12 | Start: 34079 | End: 34174 | Length: 95

.
 tttttcaaaaagtagcttatctgtggtatctggtagcatctgtttatcctatttaggatt

 tatcctgttttagaccctgttaaatagtgggttttaaagtggtcaaacagacaaaaaat

 gtaattgacattgaagactgactttactctttcaaacattaggtcactatttggtgtaag

 tatttttgtttaacatttaaagagtcaatacttttagctttaaaaaaatggcttatagact

 tttgagaaataaaactgatattatttgccttaaaaacatatatgaaatatttcttttag
 |6851 |6861 |6871 |6881 |6891 |6901
 GAGAACCCTCAATCAAAAAGAACTTATTAAATGAATTTGACAGGATAATAGAAAATCAAG
 E P S I K R N L L N E F D R I I E N Q E
 |2291 |2301
 |6911 |6921 |6931
 AAAAATCCTTAAAGGCTTCAAAAAGCACTCCAGATGgtaaaattagctttttatttatat
 K S L K A S K S T P D G
 |2311

 ctgttctccctctataggtatggtatataatattctgacctcaggtgatccacctgcctc

 tcaaagtgtgggattacagacatgagccactgtgcctaataaggacctctttatactc

 ttaaaaattactgaggacctaaaagagcatttgtttatgtggaatatatctattgatatt

 taccatattagaaatgtaaattgattaatgttaaaattagtaatatattatgcgttggtcat

 ttggaagatatgagttcactgagttatgcggatctt

Exon 13 | Start: 36348 | End: 36417 | Length: 69

.
 aattttaaatttggtatttgcatcagaaattagctaacacctttgagttatgatggttaa

 catcaactgactaaatttatgctgatttctgttgatgcttgactgtgagttatttggt

 gcatagtcattatcaatttgtgaatcaatttattttcatagttaacatttattgagcatc

 tgttacattcactgaaaattgtaaagcctataattgtctcaaatttttgtgtatttaca

 gtaacatggatattctcttagattttaactaatatgtaatatataaataattgtttcctag
 |6941 |6951 |6961 |6971 |6981 |6991
 GCACAATAAAAGATCGAAGATTGTTTATGCATCATGTTTCTTTAGAGCCGATTACCTGTG
 T I K D R R L F M H H V S L E P I T C V
 |2321 |2331
 |7001
 TACCCTTTTCGgtaagacatgttttaaatttttctaaattctaatacagtatgagaaaagtc
 P F R

 tcgtttttataaatgaacatttctaaaaataatgacactaacgttaagaagttaacactt

 cccgttttataaaatttataaaatactttggtagtattttatagtgtgttcatatcatt

 attttattttttaattttatgacagctttgtaaagtagacagattttatttctaattttat

 ggatgaagtactaagggttgagaggaattaaggaaattgctccgaatcagttaacaaaaag
 . . .
 attgcagata

Exon 14 | Start: 44382 | End: 44809 | Length: 427

.
gggtagacatttccctgggtgaaggaggtaaggagtactatgatggaattagaggggacac

.
actgagaggggtccacacttgacagactctcttctattatgtgttatgtgaggttagattgt

.
aaagtcaaaggctagccttgaaaaatgtgatattgttttggaatggcaaccatggtgaat

.
acaaaacagttaccagaatagtatcaccatgtagcaaatgagggtctgcaacaaaggcat

.
attcctaaatatttatatgtgtactagtcaataaacttatatattttctccccattgcag

 |7011 |7021 |7031 |7041 |7051 |7061
CACAACTAAGGAACGTCAAGAGATACAGAATCCAAATTTTACCGCACCTGGTCAAGAATT
T T K E R Q E I Q N P N F T A P G Q E F
 |2341 |2351

 |7071 |7081 |7091 |7101 |7111 |7121
TCTGTCTAAATCTCATTGTATGAACATCTGACTTTGGAAAAATCTTCAAGCAATTTAGC
L S K S H L Y E H L T L E K S S S N L A
 |2361 |2371

 |7131 |7141 |7151 |7161 |7171 |7181
AGTTTCAGGACATCCATTTTATCAAGTTTCTGCTACAAGAAATGAAAAAATGAGACACTT
V S G H P F Y Q V S A T R N E K M R H L
 |2381 |2391

 |7191 |7201 |7211 |7221 |7231 |7241
GATTACTACAGGCAGACCAACCAAAGTCTTTGTTCCACCTTTTAAACTAAATCACATTT
I T T G R P T K V F V P P F K T K S H F
 |2401 |2411

 |7251 |7261 |7271 |7281 |7291 |7301
TCACAGAGTTGAACAGTGTGTTAGGAATATTAACCTTGAGGAAAAACAGACAAAAGCAAAA
H R V E Q C V R N I N L E E N R Q K Q N
 |2421 |2431

 |7311 |7321 |7331 |7341 |7351 |7361
CATTGATGGACATGGCTCTGATGATAGTAAAAATAAGATTAATGACAATGAGATTTCATCA
I D G H G S D D S K N K I N D N E I H Q

	2441		2451
7371	7381	7391	7401
7411	7421		

GTTTAACAAAAACAACCTCCAATCAAGCAGCAGCTGTAACCTTCACAAAAGTGTGAAGAAGA
 F N K N N S N Q A A A V T F T K C E E E
 |2461 |2471

|7431
 ACCTTTAGgtattgtatgacaatttgtgtgatgaatttttgcctttcagttagatatttc
 P L D

.
 cgttgtaaataatgtcctgatggttttcccccttggtggtggttaattttaagccctt

 tttaatgttttagattttctaaatccaaagattaggtttaaattattctaagtgttcttt

 caagataacttcttgggacttgttaaaaaaattagacacacaatctaggactgctgt

 tactggaatatattttctatcatgctactaattttctttttaaatgtgataaaaatagg
 . . .
 gccgggcg

Exon 15 | Start: 45949 | End: 46130 | Length: 181

.
 ctcctcccagattcaagtgattctcctgcctcagcctgctgaatagctgggattacaggc

 atgtgccaccacacctggctacttttgtgttttttttacttttatatattttttttgt

 ttagtagagacagggtttctccattttggtcaggctggtcttgaactcccgacctcagat

 gatctgcccgcctcagcctcccaaagtgtgggattacaggcgtgagccactgtgcctgg

ccaggggttgctgttttttaaatttcaattttatgttctgtaagtatttattccttgatag

 |7441 |7451 |7461 |7471 |7481 |7491
ATTTAATTACAAGTCTTCAGAATGCCAGAGATATACAGGATATGCGAATTAAGAAGAAAC
 L I T S L Q N A R D I Q D M R I K K K Q
 |2481 |2491

 |7501 |7511 |7521 |7531 |7541 |7551
AAAGGCAACGCGTCTTCCACAGCCAGGCAGTCTGTATCTTGCAAAAACATCCACTCTGC
 R Q R V F P Q P G S L Y L A K T S T L P
 |2501 |2511

 |7561 |7571 |7581 |7591 |7601 |7611
CTCGAATCTCTCTGAAAGCAGCAGTAGGAGGCCAAGTTCCTCTGCGTGTCTCATAAAC
 R I S L K A A V G G Q V P S A C S H K Q
 |2521 |2531

AGgtatgtgtttgtctacaatactgatggcttttatgacagagtgttaattttatttcatt

aactagtatctacaaatggcctttgtttaagaatgaacacattagtcaggaatggatga

atgaaatcatcatattttctaattagcctgcagtggcagcctctggccccttgctaggcc

tgcctcatcctgctaaagtgatctgtgcttcaaattactacttcttttcccccttcaaa

tctttcttattttgtcattgtaaatgctctcagctaggtgttaaagtagtcttactgata

.
tt

Exon 16 | Start: 47263 | End: 47450 | Length: 187

agagattttgtaaaacatcacattttttatcctcacagtaccttcctatggcagattta

.
 gcaggaggcgtataaacggggtggaaaaggtacagcagactgtggaatgtatggatcatt

 tatattacattaaaatttttagtttctagtaaataacttaaatgtttttgtagtgaagat

 tctagtagttaatgaaaatttttggtaaattcagttttggtttgttataaattgtttttat

 tgtgtgatacatgtttacttttaaatgtttttcttttttgtgtgtgtttattttgtgtag
 |7621 |7631 |7641 |7651 |7661 |7671
 CTGTATACGTATGGCGTTTCTAAACATTGCATAAAAATTAACAGCAAAAATGCAGAGTCT
 L Y T Y G V S K H C I K I N S K N A E S
 |2541 |2551
 |7681 |7691 |7701 |7711 |7721 |7731
 TTTCAGTTTCACACTGAAGATTATTTGGTAAGGAAAGTTTATGGACTGGAAAAGGAATA
 F Q F H T E D Y F G K E S L W T G K G I
 |2561 |2571
 |7741 |7751 |7761 |7771 |7781 |7791
 CAGTTGGCTGATGGTGGATGGCTCATACCCTCCAATGATGGAAGGCTGGAAAAGAAGAA
 Q L A D G G W L I P S N D G K A G K E E
 |2581 |2591
 |7801
 TTTTATAGgtactctatgcaaaaagattgtgtgttaacttttatgtattccctcatccct
 F Y R
 |2601

 ctttcttctcttaactgtctctcgaactaaaaagttggctagaaatcaaatttttatgca

 ttttaattgttttaagtgacattatgggttaagcattctgtagaagtcttttgaaaagtgctg

 tttgtcctgggggttaataactggattttcttgatttgggacatttttcttaggcattt

 ataaatatagcccaattttataaagttaaatttggccgggtacagtggctcatgcctgtaa
 . . .
 tcccagca

Exon 17 | Start: 52044 | End: 52214 | Length: 170

```
. . . . .
tctcactatgttgccctaggctggtctcaaacttctggcctcaagcaatcctcctgcctca

. . . . .
gcttcccaaatgctgggagtataggcatgagccaccatgctcagcaatgaagtttttat

. . . . .
cagtatgatactttgatacatgtcaaataattttctgaaattatattgtagatcatatga

. . . . .
actcataaaaacttaatgatcttgaacaatgtagttttgtacagagaatagttgtagtt

. . . . .
gttgaattcagtatcatcctatgtggtttttatgataatatctacttttatttgttcag

      |7811      |7821      |7831      |7841      |7851      |7861
GGCTCTGTGTGACACTCCAGGTGTGGATCCAAAGCTTATTTCTAGAATTTGGGTTTATAA
A L C D T P G V D P K L I S R I W V Y N
                        |2611                                |2621

      |7871      |7881      |7891      |7901      |7911      |7921
TCACTATAGATGGATCATATGGAAACTGGCAGCTATGGAATGTGCCTTTCCTAAGGAATT
H Y R W I I W K L A A M E C A F P K E F
                        |2631                                |2641

      |7931      |7941      |7951      |7961      |7971      .
TGCTAATAGATGCCTAAGCCCAGAAAGGGTGCTTCTTCAACTAAAATACAGgcaagttta
A N R C L S P E R V L L Q L K Y R
                        |2651

. . . . .
aagcattacattacgtaatcatatacggcagtatggttaaggtttctgtgtagtctgtga

. . . . .
cttccatgtcaaatgttgcacaagccagttgtcagtgacagttgccatcccacactgct

. . . . .
gttctcctgtcatccctagcccccatttaagagagatcacacattcatgcattgcttgct
```

.
tccctctttccccacccctccttaacctcttgatgtatgagaagaatatgagttactaa
.
tttgatccactatgtgggattgctaataaagcatttttgcattttatttt

Exon 18 | Start: 52700 | End: 53054 | Length: 354

.
tgcttccctctttccccacccctccttaacctcttgatgtatgagaagaatatgagtta
.
ctaatttgatccactatgtgggattgctaataaagcatttttgcattttattttttgc
.
ttttaaaaataattgatattttaacaatatgaacaatatattcctagctacaaaatttt
.
taattctcagtatttcttagataaattcagtttttattctcagttattcagtgacttggt
.
taaacagtgggaattctagagtcacacttcctaaaatatgcatttttgttttcacttttag

7981	7991	8001	8011	8021	8031
ATATGATACGGA	AATTGATAGA	AGCAGAAGAT	CGGCTATA	AAAAAGATA	AATGGAAAGGGA
Y D T E I	D R S R R	S A I K K	I M E R D		
2661			2671		

8041	8051	8061	8071	8081	8091
TGACACAGCTG	CAAAACACTT	GTCTCTGT	TTCTGACATA	ATTTTCATTG	AGCGCAAA
D T A A K	T L V L C	V S D I I	S L S A N		
2681			2691		

8101	8111	8121	8131	8141	8151
TATATCTGAA	ACTTCTAGCA	ATAAACTAG	TAGTGCAGAT	ACCCAAAAAG	TGGCCATTAT
I S E T S	S N K T S	S A D T Q	K V A I I		
2701			2711		

8161	8171	8181	8191	8201	8211
TGAACTTACAG	ATGGGTGATG	TGTTAAGGCC	AGTTAGATCCT	CCCCCTCTTA	GCTGT
E L T D G	W Y A V K	A Q L D P	P L L A V		
2721			2731		

8221	8231	8241	8251	8261	8271
CTTAAAGAATGGCAGACTGACAGTTGGTCAGAAGATTATTCTTCATGGAGCAGAACTGGT					
L	K	N	G	R	L
T	V	G	Q	K	I
I	L	H	G	A	E
L	V				
2741			2751		

8281	8291	8301	8311	8321	8331.
GGGCTCTCCTGATGCCTGTACACCTCTTGAAGCCCAGAATCTCTTATGTTAAAGgtaaa					
G	S	P	D	A	C
T	P	L	E	A	P
E	S	L	M	L	K
2761			2771		

```

. . . . .
ttaatttgcaactcttggtaaaaatcagtcattgattcagttaaattctagaagttttaca
. . . . .
tttaaattttaaatgcttactaaggatgctcaatttcttagatgtactgataattttagt
. . . . .
ataaaaagcatattcttcagacagttaaagttttgtgagttttgggaggtccagaga
. . . . .
tctttcttgagcttaataatgcatttccaattaaaaagcaaaataaattgaccattt
. . . . .
gattttgggtatctgtagcttgctgccctcttggtctcatagctttgctttgatca

```

Exon 19 | Start: 59923 | End: 60078 | Length: 155

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. . . . .
tgtattttaaaactattatgtttaaatcgaagttccttttatctgttttctaatagaaca
. . . . .
tttaaatagcattaagaacttgtagcagtataaacaatatgtttgagaagtactatattg
. . . . .
tgaaaatattttcactttttatacagtttttacttatttactgtcttactaatcttccta
. . . . .
agactttttaaagtgaatatttttaaggcagttctagaagaatgaaaactcttatgatat
. . . . .
ctgtaatagaattgaatacatatttaactactaaatcaatatatttattaatttgtccag

```


.
 actaatctcagcctcccaaagttctgggattacagatgtgagccactgtgcctggcctga

 tacaattaacttgaatgttatatatgtgacttttttggtgtgtgtaacacattattacag

 |8491 |8501 |8511 |8521 |8531 |8541
 TGGATGGAGAAGACATCATCTGGATTATACATATTTGCAATGAAAGAGAGGAAGAAAAG
 W M E K T S S G L Y I F R N E R E E E K
 |2831 |2841

 |8551 |8561 |8571 |8581 |8591 |8601
 GAAGCAGCAAAATATGTGGAGGCCCAACAAAAGAGACTAGAAGCCTTATTCATAAAATT
 E A A K Y V E A Q Q K R L E A L F T K I
 |2851 |2861

 |8611 |8621 |8631
 CAGGAGGAATTTGAAGAACATGAAGgtaaaattagttatatggtacacattgttatttct
 Q E E F E E H E E
 |2871

 aatatgagaacaaagtcttagagactttgaatttaacatttttaagttaaattgtttt

 tattttgagtagtaaattgactttatttttagtatctagggtattcttttttggtgtta

 gacaaagaatagcaacaaggacagaaatatcaggtctaagccatttgtaatatattttcc

 tgaattcttacctatatgatgtggcttttgcattttgatggtagttattagctttca

 tgtgttattatgcctggaactagga

Exon 21 | Start: 66191 | End: 66312 | Length: 121

.
 aaattaaccgggcatgttgatgtgcctctaattcccagctactcgggaggctgaggcagg

agaaccacttgaaccaggagacagaggttgcaatgagccgagatcacaccactgcactc

 ccagattgggtgacagagtgagaccctgtctcaaaaaaaaaaaaaagaaaaacttttag

 cagttatatagtttcttatctttaaatctcccttctttgggtgttttatgcttggttctt

 tagtttttagttgcttttgaatttacagtttagtgaattaataatccttttgttttcttag

 |8641 |8651 |8661 |8671 |8681 |8691
 AAAACACAACAAAACCATATTTACCATCACGTGCACTAACAAGACAGCAAGTTCGTGCTT
 N T T K P Y L P S R A L T R Q Q V R A L
 |2881 |2891

 |8701 |8711 |8721 |8731 |8741 |8751
 TGCAAGATGGTGCAGAGCTTTATGAAGCAGTGAAGAATGCAGCAGACCCAGCTTACCTTG
 Q D G A E L Y E A V K N A A D P A Y L E
 |2901 |2911

 AGgtgagagagtaagaggacatataatgaggcttgatgattattcaaggtgagaagctgt

 tttagactctctggccatcacaggaaggagtgattgaaatgctgcatttctcaaaaggg

 atgtgtacatttctgggattttcagtgatgtgccagacgagtggtggtatgttttcaa

 ctatataccgagtagaggatgggaggggttctagaattttatatattaattaaatttggtt

 taaaatgcaggcaaaacttgttttattttgtccctcctgtactctgaagcaaaaaact
 .
 tt

Exon 22 | Start: 68838 | End: 69036 | Length: 198
 BE AWARE: This section overlaps with the following exon

.
catagatctatttctcaaaataatgagcattcagatattagccatctgtaatgtagttgg

.
tgatgattatgattattagagtacattttataattggaggatcatttttgccgtagggaaa

.
tagaattattaatagtttgaggcacctgagaatattatgtgagaaactgattacattaac

.
cacacccttaagatgagctctaattttgttgattttgtcctgtttaagccatctagtta

.
caatagatggaacttttttgttctgattgctttttattccaatatcttaaagggtcacag

 |8761 |8771 |8781 |8791 |8801 |8811
GGTTATTTTCAGTGAAGAGCAGTTAAGAGCCTTGAATAATCACAGGCAAATGTTGAATGAT
G Y F S E E Q L R A L N N H R Q M L N D
 |2921 |2931

 |8821 |8831 |8841 |8851 |8861 |8871
AAGAAACAAGCTCAGATCCAGTTGGAAATTAGGAAGGCCATGGAATCTGCTGAACAAAAG
K K Q A Q I Q L E I R K A M E S A E Q K
 |2941 |2951

 |8881 |8891 |8901 |8911 |8921 |8931
GAACAAGGTTTATCAAGGGATGTCACAACCGTGTGGAAGTTGCGTATTGTAAGCTATTCA
E Q G L S R D V T T V W K L R I V S Y S
 |2961 |2971

 |8941 |8951
AAAAAAGAAAAAGATTGAGgtaagtatgtaaatgctttgtttttatcagttttattaact
K K E K D S V
 |2981

.
taaaaaatgaccttactaacaanaatgattataaatccagataaagtataaagttagtta

.
tatcagagaagcaaaatccactactaatgccacaaagagataatataaaagaggatctg

.
tatttattttgaaacaaacatttaaatgataatcacttcttccattgcatctttctcatc

.
 tttctccaaacagttatactgagtatttggcgtccatcatcagatttatattctctgtta

 acagaaggaaagagataca

Exon 23 | Start: 69271 | End: 69434 | Length: 163
 BE AWARE: This section overlaps with the following exon

.
 caagggatgtcacaaccgtgtggaagttgcgtattgtaagctattcaaaaaagaaaaag

 attcaggtgaagtatgtaaagctttgtttttatcagttttattaacttaaaaaatgacct

 tactaacaaaatgattataaatccagataaagtataaagttagtttatatcagagaagca

 aaatccactactaatgccacaaagagataatataaaagaggatctgtattttattttgaa

 acaaacatttaaatgataatcacttcttcattgcatctttctcatctttctccaaacag

8961	8971	8981	8991	9001	9011
TTATACTGAGTATTTGGCGTCCATCATCAGATTATATTCTCTGTTAACAGAAGGAAAAGA					
I L S I W R P S S D L Y S L L T E G K R					
2991				3001	

9021	9031	9041	9051	9061	9071
GATACAGAATTTATCATCTTGCAACTTCAAAATCTAAAAGTAAATCTGAAAGAGCTAACA					
Y R I Y H L A T S K S K S K S E R A N I					
3011				3021	

9081	9091	9101	9111
TACAGTTAGCAGCGACAAAAAACTCAGTATCAACAACACTACCGgtacaaacctttcatt				
Q L A A T K K T Q Y Q Q L P				
3031				

.
 gtaatttttcagttttgataagtgcttgtagtttatggaatctccatatgttgaatttt

.
 tgttttgttttctgtaggtttcagatgaaatttatttcagatttaccagccacgggagc

 cccttcacttcagcaaatttttagatccagactttcagccatcttggttctgaggtggacc

 taataggatttgctggttctggttgtaaaaaaacaggtaatgcacaatatagttaatttt

 ttttattgattcttttaaaaaacattgtcttttaaaatctctta

Exon 24 | Start: 69528 | End: 69666 | Length: 138

.
 aatcacttcttccattgcatctttctcatctttctcaaacagttatactgagtatttgg

 cgtccatcatcagatttatattctctgttaacagaaggaaagagatacagaatttatcat

 cttgcaacttcaaaatctaaaagtaaatctgaaagagctaatacagtagcagcgaca

 aaaaaaactcagtatcaacaactaccggtacaaacctttcattgtaatttttcagttttg

 ataagtgccttgtagtttatggaatctccatatgttgaatttttgtttctgttag

9121	9131	9141	9151	9161	9171
GTTTCAGATGAAATTTTATTTTCAGATTTACCAGCCACGGGAGCCCCTTCACTTCAGCAAA					
V	S	D	E	I	L
F	Q	I	Y	Q	P
R	E	P	L	H	F
S	K				
3041		3051			

9181	9191	9201	9211	9221	9231
TTTTTAGATCCAGACTTTCAGCCATCTTGTTCTGAGGTGGACCTAATAGGATTTGTCGTT					
F	L	D	P	D	F
Q	P	S	C	S	E
V	D	L	I	G	F
V					
3061		3071			

9241	9251
TCTGTTGTGAAAAAACAGgtaatgcacaatatagttaatttttttattgattctttta									
S	V	V	K	K	T	G			

|3081

```
. . . . .
aaaaacattgtcttttaaaatctcttatgattagttggagctaccagttggcaaatttgc
. . . . .
tagctaactagtgatctgaaagtaagcctctttgaacctctgatttttcatgaaaagcaa
. . . . .
ttctctcaattctatattatttcaagggtacaagttacatcctagtctgtgtacttaat
. . . . .
tttatagaaattgtccttaattttattttctgcaatttatgttttcttactatttctggt
. . . . .
gtatgtgtttatcccattg
```

Exon 25 | Start: 84210 | End: 84454 | Length: 244

```
. . . . .
gccaaatttatctccagaaatcttgcaaaatctgtactcctgttagcaatgtgtgcgta
. . . . .
tacctgcttccacatgacctcagtaaaagaatgtgttgcatattggtattgaaatttta
. . . . .
gcactgtaagcaacaggtcattttggaaaacctgagctttcgccaaattcagctattttg
. . . . .
atttgcttttattattagcatataccaaaataaataggcatattagagtttcctttcttg
. . . . .
catcttaaaattcatctaacacatctataataacattcttttctttttttccattctag
```

```
      |9261      |9271      |9281      |9291      |9301      |9311
GACTTGCCCTTTTCGTCTATTTGTCAGACGAATGTTACAATTTACTGGCAATAAAGTTT
  L  A  P  F  V  Y  L  S  D  E  C  Y  N  L  L  A  I  K  F  W
              |3091                      |3101
```

```
      |9321      |9331      |9341      |9351      |9361      |9371
GGATAGACCTTAATGAGGACATTATTAAGCCTCATATGTTAATTGCTGCAAGCAACCTCC
  I  D  L  N  E  D  I  I  K  P  H  M  L  I  A  A  S  N  L  Q
```

	3111		3121
9381	9391	9401	9411
9421	9431		
AGTGGCGACCAGAATCCAAATCAGGCCTTCTTACTTTATTTGCTGGAGATTTTCTGTGT			
W	R	P	E
S	K	S	G
L	L	T	L
F	A	G	D
F	S	V	F
	3131		3141
9441	9451	9461	9471
9481	9491		
TTTCTGCTAGTCCAAAAGAGGGCCACTTTCAAGAGACATTCAACAAAATGAAAAATACTG			
S	A	S	P
K	E	G	H
F	Q	E	T
F	N	K	M
K	N	T	V
	3151		3161
9501.			
TTGAGGtaagggttacttttcagcatcaccacacattttggtatttttctattttgacagt			
E			
.			
ccagttatcaaggaaatagctttttatacaaattggatagttgaggtagtgtgaggtaaa			
.			
gtttaatcatatattaattgcccatgaacctcaggagatgggggaatggggaaatgacag			
.			
caactagaaagagaagaatgacttgaagggaatgagttaggagaaattgtgagaaggat			
.			
gttcagaaatgcagactttgtaagcaaactggaaattggttacaagaataatatgagtta			
.			
tctgt			

Exon 26 | Start: 86419 | End: 86565 | Length: 146

.
agcttcaatccagatcatatgacagcttgctgattaaactagatgacagagaagatctct
.
ttccttcagatacacatactttttctctgttcccctctccctatcagctagattccccta
.
aatcactgatactgggttttgtaattttgcatcggcatgtttgacaattggtatcacattt

.
 agggtttttcattcttttttggtccaaacttttcatttctgcttttaaggaaatacttt

 tggaaacataaatatgtgggtttgcaatttataaagcagcttttccacttattttcttag

 |9511 |9521 |9531 |9541 |9551 |9561
 AATATTGACATACTTTGCAATGAAGCAGAAAACAAGCTTATGCATATACTGCATGCAAAT
 N I D I L C N E A E N K L M H I L H A N
 |3171 |3181

 |9571 |9581 |9591 |9601 |9611 |9621
 GATCCCAAGTGGTCCACCCCACTAAAGACTGTACTTCAGGGCCGTACACTGCTCAAATC
 D P K W S T P T K D C T S G P Y T A Q I
 |3191 |3201

 |9631 |9641
 ATTCCTGGTACAGGAAACAAGCTTCTGgtaagttaatgtaaactcaaggaatattataag
 I P G T G N K L L
 |3211

 aagtatatatggaggccatcgtatattctgttgatatacctagtaaacatggtaaaatgta

 attaaacttaattagaaaatgtggttggttatgtggctcctgtaagtatagttatttagaa

 attttatttattgaagcaagatatgaaactctgggtgcacactttccaaacaggtgcttt

 catttacatgtgattgaaaagtgtttttgtcatttatttcactgttccatacaattagg

 gttgtttctaagctgtttgtaagctgt

Exon 27 | Start: 87683 | End: 89193 | Length: 1510

.
 aataataaaaaacaaaagattaaagcataagtgacgtcccctacctccttttttatcttt

.
tactgtgattattcttcatcttccttccttttcatgtcattttatatgttcttatgtaaa

.
attactttcatctagaataggaataatgtgaactgaaatcacctaacctattaggagtta

.
ggggaggaggactgtgtgtaatatttgctgcttaaataattttcaatgaaaagttacttt

.
gatttagttttttatgttactacataattatgataggctacgttttcatttttttatcag

|9651 |9661 |9671 |9681 |9691 |9701
ATGTCTTCTCCTAATTGTGAGATATATTATCAAAGTCCTTTATCACTTTGTATGGCCAAA
M S S P N C E I Y Y Q S P L S L C M A K
|3221 |3231

|9711 |9721 |9731 |9741 |9751 |9761
AGGAAGTCTGTTTCCACACCTGTCTCAGCCCAGATGACTTCAAAGTCTTGTAAGGGGAG
R K S V S T P V S A Q M T S K S C K G E
|3241 |3251

|9771 |9781 |9791 |9801 |9811 |9821
AAAGAGATTGATGACCAAAAGAACTGCAAAAAGAGAAGAGCCTTGATTCTTGAGTAGA
K E I D D Q K N C K K R R A L D F L S R
|3261 |3271

|9831 |9841 |9851 |9861 |9871 |9881
CTGCCTTTACCTCCACCTGTTAGTCCCATTGTACATTTGTTTCTCCGGCTGCACAGAAG
L P L P P P V S P I C T F V S P A A Q K
|3281 |3291

|9891 |9901 |9911 |9921 |9931 |9941
GCATTTTCAGCCACCAAGGAGTTGTGGCACCAAAATACGAAACACCCATAAAGAAAAAAGAA
A F Q P P R S C G T K Y E T P I K K K E
|3301 |3311

|9951 |9961 |9971 |9981 |9991 |10001
CTGAATTCTCCTCAGATGACTCCATTTAAAAAATTCAATGAAATTTCTCTTTTGAAAAGT
L N S P Q M T P F K K F N E I S L L E S
|3321 |3331

|10011 |10021 |10031 |10041 |10051 |10061
AATTCAATAGCTGACGAAGAACTTGCATTGATAAATACCCAAGCTCTTTTGTCTGGTTCA
N S I A D E E L A L I N T Q A L L S G S
|3341 |3351

|10071 |10081 |10091 |10101 |10111 |10121
 ACAGGAGAAAAACAATTTATATCTGTCAGTGAATCCACTAGGACTGCTCCCACCAGTTCA
 T G E K Q F I S V S E S T R T A P T S S
 |3361 |3371

|10131 |10141 |10151 |10161 |10171 |10181
 GAAGATTATCTCAGACTGAAACGACGTTGTACTACATCTCTGATCAAAGAACAGGAGAGT
 E D Y L R L K R R C T T S L I K E Q E S
 |3381 |3391

|10191 |10201 |10211 |10221 |10231 |10241
 TCCCAGGCCAGTACGGAAGAATGTGAGAAAAATAAGCAGGACACAATTACAATAAAAAA
 S Q A S T E E C E K N K Q D T I T T K K
 |3401 |3411

|10251 | +1 | +11 | +21 | +31 | +41 | +51
 TATATCTAAGCATTGCAAAGGCGACAATAAATTATTGACGCTTAACCTTCCAGTTTAT
 Y I *

 | +61 | +71 | +81 | +91 | +101 | +111
 AAGACTGGAATATAATTTCAAACCACACATTAGTACTTATGTTGCACAATGAGAAAAGAA

 | +121 | +131 | +141 | +151 | +161 | +171
 ATTAGTTTCAAATTTACCTCAGCGTTTGTGTATCGGGCAAAAATCGTTTTGCCCGATTCC

 | +181 | +191 | +201 | +211 | +221 | +231
 GTATTGGTATACTTTTGCTTCAGTTGCATATCTTAAACTAAATGTAATTTATTAATAA

 | +241 | +251 | +261 | +271 | +281 | +291
 TCAAGAAAAACATCTTTGGCTGAGCTCGGTGGCTCATGCCTGTAATCCCAACACTTTGAG

 | +301 | +311 | +321 | +331 | +341 | +351
 AAGCTGAGGTGGGAGGAGTGCTTGAGGCCAGGAGTTCAAGACCAGCCTGGGCAACATAGG

 | +361 | +371 | +381 | +391 | +401 | +411
 GAGACCCCATCTTTACAAAGAAAAAAAAAAGGGGAAAAGAAAATCTTTAAATCTTTGG

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

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

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

AGTTTAGCTACTATTTTAGGGGATTTTTTTTAGAGGTAACCTACTATGAAATAGTTCTCC

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

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

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

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

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

 |+901
TGGCACTGATTctgcctgctttattttagcgctatcacaggaccagagcctatgcctt

.
tttaaacttaccacaaaagcagaagattaattcaatttaagatgatactctcatttgta

.
cgtccttttttttttttttggagatggagtccttgctttgtcgcccatgctggagtgag

.
tggcatgatcctggctcactgcagcctccacttcccgggttcacgtaattctccacctc

.
aagcctccctagtagctgggattacagggacgcaccaccatgccagctaatttttgc

. . .
ttttagtagag

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