

Gene: BRCA1 - Sequence: NG_005905.2
Transcript: NM_007294.3 - Protein: NP_009225.1
Date : February 26, 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: 92501 | End: 92713 | Length: 212

```
. . . . .
ttactttatatttaccgaaactggagacctccattagggcggaagagtgggggattggga

. . . . .
cctcttcttacgactgctttggacaataggtagcgattctgaccttcgtacagcaattac

. . . . .
tgtgatgcaataagccgcaactggaagagtagaggctagagggcaggcactttatggcaa

. . . . .
actcaggtagaattcttcctcttccgtctctttccttttacgtcatccgggggcagactg

. . . . .
ggtggccaatccagagccccgagagacgcttggtctctttctgtccctcccacctctgat

|-229    |-219    |-209    |-199    |-189    |-179
GTACCTTGATTTCGTATTCTGAGAGGCTGCTGCTTAGCGGTAGCCCCTTGGTTTCCGTGG

|-169    |-159    |-149    |-139    |-129    |-119
CAACGGAAGCGCGGAATTACAGATAAATTAAGTAACTGCGACTGCGCGCGTGAGCTCG

|-109    |-99     |-89     |-79     |-69     |-59
CTGAGACTTCCTGGACGGGGACAGGCTGTGGGGTTTCTCAGATAACTGGGCCCTGCGC

|-49     |-39     |-29     . . . . .
TCAGGAGGCCTTACCCCTCTGCTCTGGGTAAAGgtagtagagtcccgggaaaggacagg

. . . . .
gggccaagtgatgctctgggggtactggcgtgggagagtggatttccgaagctgacagat

. . . . .
gggtattctttgacgggggtagggcggaacctgagaggcgtaaggcgttgtgaaccct

. . . . .
```

ggggaggggggcagttttaggtcgcgagggaagcgctgaggatcaggaagggggcactg

.
agtgtccgtgggggaatcctcgtgataggaactggaatatgccttgagggggacactatg

.
tctttaaaaacgtcggctggtcatgaggtcagg

Exon 2 | Start: 93869 | End: 93967 | Length: 98

```
. . . . .
aaggctaccaccacctacccggtcagtcactcctctgtagctttctctttcttgagaaa

. . . . .
ggaaaagacccaagggttggcagcaatatgtgaaaaaattcagaatttatgttgtctaa

. . . . .
ttacaaaaagcaacttctagaatctttaaaaataaaggacgttggtcattagttctttggt

. . . . .
ttgtattattctaaaaccttccaaatcttaaatttactttattttaaaatgataaaatga

. . . . .
agttgtcattttataaaccttttaaaaagatatatatatgtttttctaattgtgttaaa

|-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      . . . . .
CATTAATGCTATGCAGAAAAATCTTAGAGTGTCCCATCTGgtaagtcagcacaagagtgtg
I N A M Q K I L E C P I C
      |21

. . . . .
ttaatttgggattcctatgattatctcctatgcaaatgaacagaattgaccttacatact

. . . . .
agggagaagaaagacatgtctagtaagattaggctattgtaattgctgattttcttaactg

. . . . .
aagaactttaaaaatatagaaaatgattccttgttctccatccactctgcctctccact

. . . . .
cctctccttttcaacacaaatcctgtggtccgggaaagacagggactctgtcttgattgg

. . . . .
ttctgcactggggcaggaatctagtttagattaactggc
```

Exon 3 | Start: 102205 | End: 102258 | Length: 53

```
. . . . .
gaattcgtacgaactattatcaactaatcttttaaagtctgatgatagtatagagtattg
. . . . .
aagggatcaatataattctgttttgatatctgaaagctcactgaaggtaaggatcgtatt
. . . . .
ctctgctgtattctcagttcctgacacagcagacatttaataaatattgaacgaactga
. . . . .
ggccttatgttgactcagtcataacagctcaaagttgaacttattcactaagaatagctt
. . . . .
tatttttaataaattattgagcctcatttattttctttttctccccctaccctgcta
|81      |91      |101     |111     |121     |131     .
TCTGGAGTTGATCAAGGAACCTGTCTCCACAAAGTGTGACCACATATTTTGCAAgtaagt
L E L I K E P V S T K C D H I F C K
      |31              |41
. . . . .
ttgaatgtgttatgtggctccattattagcttttgttttgtccttcataaaccaggaaa
. . . . .
cacctaactttatagaagctttactttcttcaattaagtgagaacgaaaaatccaactcc
. . . . .
atttcattctttctcagagagtatatagttatcaaaagttggttgtaatcatagttcctg
. . . . .
gtaaagttttgacatatattatcttttttttttttttgagacaaagtctcgctctgtcg
. . . . .
cccaggctggagtgagtgcatgatcttggctcactgcaacctccgcccccg
```

Exon 4 | Start: 111451 | End: 111528 | Length: 77

```
. . . . .
ttatgaatatatcattaaatagccatattaacttttattaagttttatgtgatcataac

. . . . .
agtaagccatatgcatgtaagttcagttttcatagatcattgcttatgtagtttaggttt

. . . . .
ttgcttatgcagcatccaaaacaattaggaactattgcttgtaattcacctgccatta

. . . . .
ctttttaaatggctcttaagggcagttgtgagattatctttcatggctatttgcctttt

. . . . .
gagtattctttctacaaaaggaagtaaattggttctttctttctttataatttata

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

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

. . . . .
taggaaaaagtagaaattatttaataacatagcgttcctataaaaccattcatcagaaaa

. . . . .
atttataaaagagtttttagcacacagtaaattatttccaaagttattttcctgaaagtt

. . . . .
ttatgggacatctgccttatacaggtattagaaacttactgcctttctctaagtcttcta

. . . . .
gtgtaaaaacttgcgacttatgtaaagtagggctgtatcgccgtgccccattgtctgt

. . . . .
taatcttgtttttatatt
```

Exon 5 | Start: 113028 | End: 113116 | Length: 88

.
taaattaaacaaaaaaaaaagtactctagttttctatgcaatgcattatatctgctgtgg

.
atttagggcagtattatatcagataattttaggcatttggtaggcctaaatgaatgacaa

.
aaagttactaaatcactgccatcacacggttttatacagatgtcaatgatgtattgattat

.
agaggttttctactgttgctgcatcttatttttattgtttacatgtcttttcttatttt

.
agtgtccttaaaaggttgataatcacttgctgagtggtgtttctcaaacaatttaatttca

 |221 |231 |241 |251 |261 |271
GAGCCTACAAGAAAGTACGAGATTTAGTCAACTTGGTTGAAGAGCTATTGAAAAATCATTG
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
TGCTTTTCAGCTTGACACAGGTTTGGAGTgtaagtgttgaatatccaagaatgacactc
A F Q L D T G L E Y
 |101

.
aagtgtgtccatgaaaactcaggaagtttgcacaattactttctatgacgtggtgataa

.
gaccttttagtctaggttaatttttagttctgtatctgtaattctatttttaaaaaattact

.
cccactgggtctcacaccttattttatcaatcgtaagggtgcacatttttcacatcttaaca

.
tctctgaaattgggaacattttactattgagggtgtgtcatttgtttaatttggtgtgctt

.
tctttcttagtgatacagaaaataatagt

Exon 6 | Start: 113723 | End: 113862 | Length: 139

.
ttacattgttggtgtccttagctttagtgaaatacagtattgataggcaaatttcttagtg

.
ttaaggtagaaaacaaggactctaaataactttgatggctgtgtatttgttttgtttc

.
ctaggagtaaaatttccagttgattttttaaaatttgatttttaaaaaaatcacaggta

.
accttaatgcattgtccttaacacaacaagagcatacatagggtttctcttggtttcttt

.
gattataattcatacatttttctctaactgcaaacataatgttttccttgattttaca

311	321	331	341	351	361
ATGCAAACAGCTATAATTTTGCAAAAAGGAAAAATAACTCTCCTGAACATCTAAAAGATG					
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

.
ctaatttttgtatttttagtagagacggggtttcatcatgttggccaggctggtctcgaa

.
ctcctaacctcaggtgatccaccacctcggctccccaattgctgggattacaggtgtg

.
agccactgtgcccggccggt

Exon 7 | Start: 118104 | End: 118209 | Length: 105

.
tactactactattatTTTgtagagactgggtctcactctgttgcttatgctgggtcttgaa

.
ctcctggcctcaagcagtcctgctccagcctcccaaagtgctgggattataggcatgagc

.
taccgctcccagccccagacatttttagtgtgtaaattcctgggcattttttccaggcatc

.
atacatgttagctgactgatgatggccaattttttgtccatgggtgtcaagtttctctt

.
caggaggaaaagcacagaactggccaacaattgcttgactgttctttaccatactgttta

451	461	471	481	491	501
CAGGAAACCAGTCTCAGTGTCCA	ACTCTCTAACCTTGGA	ACTGTGAGAACTCTGAGGACA			
Q E T S L S V Q L S N L G T V R T L R T					
151			161		

511	521	531	541	.	.
AAGCAGCGGATACAACCTCAAAAGACGTCTGTCTACATTGAATTGG	gtaagggtctcagg				
K Q R I Q P Q K T S V Y I E L G					
171			181		

.
ttttttaagtattttaataataattgctggattccttatcttatagttttgccaanaatct

.
tggtcataattttgtattttgtggtaggcagctttgggaagtgaattttatgagccctatgg

.
tgagttataaaaaatgtaaaagacgcagttccaccttgaagaatcttacttttaaaaagg

.
gagcaaaagaggccaggcatggtggctcacacctgtaatccagcactttgggaggccaa

.
agtgggtggatcacctgaggtcgggagttcgagaccagcctagcca

Exon 8 | Start: 120695 | End: 120740 | Length: 45

.
gtagaaacgggggtctcactttgttggccaggctggtcttgaactcctaacctcaaataa

.
tccacccatctcggcctcctcaagtgtctgggattacaggtgagagccactgtgcctggcg

.
aagcccatgcctttaaccacttctctgtattacatactagcttaactagcattgtacctg

.
ccacagtagatgctcagtaaataatttctagttgaatatctgtttttcaacaagtacattt

.
ttttaacccttttaattaagaaaacttttattgatttatttttgggggaaatttttta

 |551 |561 |571 |581 |591 . .
GATCTGATTCTTCTGAAGATACCGTTAATAAGGCAACTTATTGCAGgtgagtcaaagaga
S D S S E D T V N K A T Y C S
 |191

.
acctttgtctatgaagctggtattttcctatttagttaatattaaggattgatgtttctc

.
tctttttaaaaatattttaacttttatttttaggttcagggatgtatgtgcagtttgttat

.
ataggtaaacacacgacttgggatttggtgtatagattttttcatcatccgggtactaa

.
gcataccccacagttttttgtttgctttctttctgaatttctccctcttcccaccttcct

.
ccctcaagtaggctggtgttttctccagactagaatcatggtattgg

Exon 9 | Start: 122062 | End: 122138 | Length: 76

.
tgtatTTTTtagtagagatggggTTTcaccatgttggccaggctggtcttgaactcatgac

.
ctcaagtggTccaccgcctcagcctcccaaagtgctggaattacaggcttgagccaccg

.
tgcccagcaaccatttcatttcaactagaagtttctaaaggagagagcagctttcactaa

.
ctaaataagattggTcagctttctgtaatcgaaagagctaaaatgtttgatcttggtcat

.
ttgacagttctgcatacatgtaactagtgtttcttattaggactctgtcttttcctata

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	

.
tgggtagatacagtactgtaattagattattctgaagaccatttgggacctttacaacc

.
acaaaatctcttggcagagtttagagtatcattctctgtcaaatgtcgtgggtatggtctga

.
tagatttaaattggtactagactaatgtacctataataagaccttctgtaactgattgttg

.
ccctttcgtTTTTTTTTTgtttgtttgtttgtttTTTTTgagatggggTctcactctg

.
ttgcccaggctggagtg

Exon 10 | Start: 123124 | End: 126549 | Length: 3425

.
cagttttctgatggccaatctgcttttaattcactccttagacgttagagaaataggtgtg

.
gtttctgcatagggaaaattctgaaattaaaaatttaatggatcctaagtggaaataatc

.
taggtaaataggaattaaatgaaagagtatgagctacatcttcagtataacttggtagttt

.
atgaggttagtttctctaatatagccagttggttgatttccacctccaaggtgtatgaag

.
tatgtatttttttaatgacaattcagtttttgagtaccttgttatttttgatatattttca

|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
|231 |241

|731 |741 |751 |761 |771 |781
ATGATTGTAACACCACTGAGAAGCGTGCAGCTGAGAGGCATCCAGAAAAGTATCAGGGTA
D L N T T E K R A A E R H P E K Y Q G S
|251 |261

|791 |801 |811 |821 |831 |841
GTTCTGTTTCAAACCTGCATGTGGAGCCATGTGGCACAATACTCATGCCAGCTCATTAC
S V S N L H V E P C G T N T H A S S L Q
|271 |281

|851 |861 |871 |881 |891 |901
AGCATGAGAACAGCAGTTTATTACTCACTAAAGACAGAATGAATGTAGAAAAGGCTGAAT
H E N S S L L L T K D R M N V E K A E F
|291 |301

|911 |921 |931 |941 |951 |961
TCTGTAATAAAAAGCAAACAGCCTGGCTTAGCAAGGAGCCAACATAACAGATGGGCTGGAA
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
CTGATCCCCTGTGTGAGAGAAAAGAATGGAATAAGCAGAACTGCCATGCTCAGAGAATC					
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
CTAGAGATACTGAAGATGTTCTTGATAACACTAAATAGCAGCATTGAGAAAGTTAATG					
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
GAGCATTTGTTACTGAGCCACAGATAATACAAGAGCGTCCCTCACAAAATAAATTAAGC					
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

ATGCACCTGGTTCTTTTACTAAGTGTTCAAATACCAGTGAACCTTAAAGAATTTGTCAATC
 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
 CTAGCCTTCCAAGAGAAGAAAAAGAAGAGAACTAGAAACAGTTAAAGTGTCTAATAATG
 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
 CAAAGCGCCAGTCATTTGCTCCGTTTTCAAATCCAGGAAATGCAGAAGAGGAATGTGCAA
 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
 CATTCTCTGCCCACTCTGGGTCCTTAAAGAAACAAAGTCCAAAAGTCACTTTTGAATGTG

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
 AACAAAAGGAAGAAAATCAAGGAAAGAATGAGTCTAATATCAAGCCTGTACAGACAGTTA
 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
 GACTCATTACTCCAAATAAACATGGACTTTTACAAAACCCATATCGTATACCACCCTTT
 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
 TTCCCATCAAGTCATTGTGTTAAACTAAATGTAAGAAAAATCTGCTAGAGGAAAACTTTG
 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
 AGGAACATTCAATGTACCTGAAAGAGAAAATGGGAAATGAGAACATTCCAAGTACAGTGA
 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
 GTGATGAAAAATTCAAGCAGAACTAGGTAGAAAACAGAGGGCCAAAATTGAATGCTATGC
 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	E	V	V
		Q	T	V	N
				T	D
				F	
		1111			1121
3371	3381	3391	3401	3411	3421
TCTCTCCATATCTGATTTCAGATAACTTAGAACAGCCTATGGGAAGTAGTCATGCATCTC					
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
GTTTTGCTGAAATGACATTAAGGAAAGTTCTGCTGTTTTTAGCAAAAGCGTCCAGAAAG					
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
3671	3681	3691	3701	3711	3721
CCTGCTTCCAACACTTGTTATTTGGTAAAGTAAACAATATACCTTCTCAGTCTACTAGGC					
C	F	Q	H	L	L
		F	G	K	V
		N	N	I	P
		S	Q	S	T
				R	H
		1231			1241
3731	3741	3751	3761	3771	3781
ATAGCACCGTTGCTACCGAGTGTCTGTCTAAGAACACAGAGGAGAATTTATTATCATTGA					
S	T	V	A	T	E
		C	L	S	K
		N	T	E	E
		N	L	L	S
				L	K
		1251			1261

Exon 11 | Start: 126952 | End: 127040 | Length: 88

```
. . . . .
taaaggaactgcttcttaaacttgaaacatgttcctcctaagggtgcttttcatagaaaaa
. . . . .
agtccttcacacagctaggacgtcatctttgactgaatgagctttaacatcctaattact
. . . . .
gggtggacttacttctggtttcattttataaaaagcaaataccaggtgtcccaaagcaaggaa
. . . . .
tttaatcattttgtgtgacatgaaagtaaataccagtcctgccaatgagaagaaaaagaca
. . . . .
cagcaaggtgcagcgtttatagtctgcttttacatctgaacctctgtttttgttatattaa

      |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
              |1371                                |1381

      |4161      |4171      |4181      . . . . .
CCTCTCAGAGTGACATTTTAACCACTCAGgtaaaaagcgtgtgtgtgtgtgcacatgcgt
      S  Q  S  D  I  L  T  T  Q
              |1391

. . . . .
gtgtgtgggtgtcctttgcattcagtagtatgtatcccacattcttaggtttgctgacatc
. . . . .
atctctttgaattaatggcacaattgtttgtggttcattgtctccttaaattagactgta
. . . . .
agcaccttgatggaactcatactaccttttatttcacacacacgcacacgcgcacacaca
. . . . .
gcctacacatacactgccttagctcattgtagcatactaaatactgattttaatgaataag
. . . . .
ctaaaccttcgaaacccatttgctaatacc
```

Exon 12 | Start: 135409 | End: 135580 | Length: 171

.
ttatTTTTtTcactccctagctTTTaaagaaaataaccaacttcaaaaggacatcacaata

.
acatcaagtctatTTgggggaatttgaggattTTTTccctcactaacatcatttggaat

.
aatttcatgggcattaattgcatgaatgtggttagattaaaagggtgttcagctagaactt

.
gtagttccatactaggtgatttcaattcctgtgctaaaattaattgtatgatatatTTT

.
catttaatggaaagcttctcaaagtatttcatttTcttggtgccatttatcgTTTTtgaa

|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
GCTGTGTTAGAACAGCATGGGAGCCAGCCTTCTAACAGCTACCCCTTCATCATAAGTGAC
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

.
tgttggccaaacactgatatcttaagcaaaattctttccttcccctttatctccttctga

.
agagtaaggacctagctccaacattttatgatccttgctcagcacatgggtaattatgga

.
gccttggttcttgtccctgctcacaactaatataaccagtcagagggaccaaggcagtca

.
ttcatgttgctcatctgagtacctacaacaagtagatgctatggggagcccatggaagata

.
catggtataacaacatagctcttgctctattggaagctaagtggaatggggaga

Exon 13 | Start: 141370 | End: 141496 | Length: 126

```
. . . . .
ttcaggccaaggaatatagatttttttttcagccttgctctcagctgggtgtctttatt

. . . . .
tactctgtcttaaagtgttccttttattatcattattttttaatcattgaattccat

. . . . .
ttggtgctagcatctgtctgttgcatgcttggtgtttataaaattctgcctgatatactt

. . . . .
gtttaaaaaccaatttgtgtatcatagattgatgcttttgaaaaaatcagtattctaac

. . . . .
ctgaattatcactatcagaacaaagcagtaaagtagatttgtttctcattccatttaa

    |4361      |4371      |4381      |4391      |4401      |4411
CAGTATTAAC TTCACAGAAAAGTAGTGAATACCCTATAAGCCAGAATCCAGAAGGCCTTT
  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
CTGCTGACAAGTTTGAGGTGTCTGCAGATAGTTCTACCAGTAAAAATAAAGAACCAGGAG
  A  D  K  F  E  V  S  A  D  S  S  T  S  K  N  K  E  P  G  V
                        |1481                                |1491

    |4481
TGGAAAGgtaagaacatcaatgtaaagatgctgtggtatctgacatctttatttatatt
  E   R

. . . . .
gaactctgattgttaatttttttcaccatactttctccagttttttgcatacaggcattt

. . . . .
atacactttttattgctctaggatacttcttttgtttaatcctatataggttttttgaacc

. . . . .
tataacataagctacaacatgagaaatgtgcggttagatagatatgtcccttctgaaggt

. . . . .
cagaaaaaaatataatggaggtaaaacctgaacaagcttggaactgatggtagacttct

. . .
tcaaggc
```

Exon 14 | Start: 143463 | End: 143653 | Length: 190

.
cagcctcccgagtagctgagattacaggcgccagccaccacaccagctactgacctgct

.
tttaaacagctgggagatatggtgcctcagaccaaccaaccccatgttatatgtcaacc

.
ctgacatatattggcaggcaacatgaatccagacttctaggctgtcttgctgggctctttttt

.
gccagtcatttctgatctctctgacatgagctgtttcatttatgctttggctgccagca

.
agtatgatttgcctttcacaattggtggcgatggttttctccttcatttatctttcta

4491	4501	4511	4521	4531	4541
GT	CAT	CCCC	TTCT	AAATG	CCCATC
AT	AG	ATG	ATAG	GTGG	TACATG
CA	CAG	TTG	CTCT	GG	
S	S	P	S	K	C
P	S	K	C	P	S
L	D	D	R	W	Y
M	H	S	C	S	G
1501			1511		

4551	4561	4571	4581	4591	4601
GAG	TCT	TCA	GA	ATA	GA
AA	CT	CA	AG	AG	AG
CT	CA	AG	AG	CT	CA
TA	AG	GT	TT	GT	GA
TG	GA				
S	L	Q	N	R	N
Y	P	S	Q	E	E
L	I	K	V	V	D
V	E				
1521			1531		

4611	4621	4631	4641	4651	4661
GG	AG	CA	AC	AG	CT
GG	AA	AG	CT	GG	CC
AC	AC	GC	AT	TT	GA
CG	AA	AC	AT	CT	TA
CT	TA	CT	TA	CT	TA
E	Q	Q	L	E	E
S	G	P	H	D	L
T	E	T	S	Y	L
P	R				
1541			1551		

|4671
GCAAGATCTAGgtaatatattcatctgctgtattggaacaaacactttgattttactctga
Q D L E

.
atcctacataaagatatcttggttaaccaacttttagatgtactagtctatcatggacac

.
ttttgttatacttaattaagcccacttttagaaaaatagctcaagtgttaatcaaggttta

.
cttgaaaattattgaaactgttaatccatctatatattttaattaatggtttaactaatgat

.

tttgaggatgagggagtcttggtgtactctaaatgtattatttcaggccaggcatagtgg

. . .
ctcacgcctgt

Exon 15 | Start: 146746 | End: 147056 | Length: 310

.
ttctaaaattatactattcctatgactaaacctttgcatatatcttttatctccctagga

.
tatatttctaaaactagcattgttgactgaaagtgtaaatacgtgttaagggtgtttgcta

.
cataatgccatatttccttttttaggaaactaagctactttggatttccaccaacactgta

.
ttcatgtaccattttttctttaacctaactttattggctttttaattcttaacagaga

.
ccagaactttgtaattcaacattcatcgttggtgttaaattaaacttctcccattcctttca

 |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
CTTCTGAAGACAGAGCCCCAGAGTCAGCTCGTGTGGCAACATACCATCTTCAACCTCTG
 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
CATTGAAAGTTCCCCAATTGAAAGTTGCAGAACTGCCCAGAGTCCAGCTGCTGCTCATA
 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
CAGAAGAATTTgtgagtgatatccatatgtatctccctaatagacttaacaacatt
 E E F
 |1661

.
ctggaaagagttttatgtaggtattgtcaattaataacctagaggaagaatctagaaa

.
caatcacagttctgtgtaatttaatttcgattactaatttctgaaaatttagatctagat

.
aaagctatagtgtggattattttatgtatatatttacttgagaaaataattattaaatatta

.
gtggaaaagctatactttgggtatgatataggactttcgaattggaattttcctttctat

. . .
ctgtaaaagca

Exon 16 | Start: 150289 | End: 150376 | Length: 87

```
. . . . .
cgagacttcaggtgtcttagaattttttaaatgtacccttctgagaaaaacagagactt

. . . . .
aaagctaggataactgggtattctatTTTTTTTTTTTTTTTTTTTTTtacctccagcctggg

. . . . .
tgacagagcaagactctgtctaaaaaaaaaaaaaaaaaattcactttaaatagttccag

. . . . .
gacacgtgtagaacgtgcaggattgctacataggtaaacatatgccatggtggaataact

. . . . .
agtattctgagctgtgtgctagaggtaactcatgataatggaatatttgatttaatttca

      |4991      |5001      |5011      |5021      |5031      |5041
ATGCTCGTGTACAAGTTTGCCAGAAAACACCACATCACTTTAACTAATCTAATTACTGAA
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

. . . . .
tctgctgcataaaaccacatgaggcgaggcacggtggcgcatgcctgtaatcgcagcact

. . . . .
ttgggaggccgaggcgggcagatcacgagattaggagatcgagaccatcctggccagcat

. . . . .
ggtgaaaccccgctcttactaaaaaataaaaaaattagctgggtgtggtcgcggtgcgcct

. . . . .
gtagtcccagctactcgtgaggctgaggcaggagaatcacttgaaccggggagatggagg

. . . . .
ttgcagtgagccgagatcatgccactgc
```

Exon 17 | Start: 154033 | End: 154110 | Length: 77

```
. . . . .
ctataatggagatctatagctagccttggcgtctagaagatgggtgttgagaagaggag
. . . . .
tggacagatatttcctctgtgtcttaacttcatatcagcctcccctagacttccaaatatc
. . . . .
catacctgctgggtataaattagtggtgttttcagcctctgattctgtcaccaggggtttt
. . . . .
agaatcataaatccagattgatcttgggagtgtaaaaactgaggctcttttagcttctta
. . . . .
ggacagcacttcctgattttgttttcaacttctaatacctttgagtgtttttcattctgca
      |5081      |5091      |5101      |5111      |5121      |5131
ATGCTGAGTTTGTGTGTAACGGACACTGAAATATTTTCTAGGAATTGCGGGAGGAAAAT
  A  E  F  V  C  E  R  T  L  K  Y  F  L  G  I  A  G  G  K  W
                        |1701                                |1711

      |5141      |5151 . . . . .
GGGTAGTTAGCTATTTCTgtaagtataataactatcttctcccctcctccctttaacacctc
  V  V  S  Y  F  W

. . . . .
agaattgcatttttacacctaacgtttaacacctaagggtttttgctgatgctgagtctga
. . . . .
gttacaaaagggtctttaattgtaataactaaactacttttatctttaatatcactttgtt
. . . . .
cagataagctgggtgatgctgggaaaatgggtctcttttataactaataggacctaatactg
. . . . .
ctcctagcaatgttagcatatgagctagggtatttatttaatatgctggcaggaatccatgt
. . . . .
gcagcaggcaaacttata
```

Exon 18 | Start: 154611 | End: 154651 | Length: 40

```
. . . . .
tataactaataggacctaactctgctcctagcaatgtagcatatgagctagggatttatt
. . . . .
taatagtcggcaggaatccatgtgcagcaggcaaacttataatgtttaaattaaacatca
. . . . .
actctgtctccagaaggaaactgctgctacaagccttattaaagggtgtggctttagag
. . . . .
ggaaggacctctcctctgtcattcttcctgtgctcttttgtgaatcgctgacctctctat
. . . . .
ctccgtgaaaagagcacgttcttctgctgtatgtaacctgtcttttctatgatctcttta

      |5161      |5171      |5181      |5191      . . .
GGGTGACCCAGTCTATTAAAGAAAGAAAAATGCTGAATGAGgtaagtacttgatgttaca
  V  T  Q  S  I  K  E  R  K  M  L  N  E
      |1721                        |1731

. . . . .
aactaaccagagatatcattcagtcatatagttaaaaatgtatttgcttccttccatca
. . . . .
atgcaccactttccttaacaatgcacaaattttccatgataatgaggatcatcaagaatt
. . . . .
atgcaggcctgcactgtgggtcatacctataatcccagcgctttgggaggctgaggcgct
. . . . .
tggatcacctgatgtcgggagttcaagaccagcctgaccaacatggagaaaccccgtttc
. . . . .
tactaaaaatacaaaattagccgggcttggtggcacttgcc
```

Exon 19 | Start: 160849 | End: 160932 | Length: 83

```
. . . . .
tgcattcaaaagattctcctgcctcagcctccaagtagctgggattacaggtgcctgcc
. . . . .
accacgcccactaattttttgtattttttagtagagatgaggtttcaccatgttggtcag
. . . . .
actggtgtcgaactcctgacctcaagtgatctgcctgcctcagtctcccaaagtgctagg
. . . . .
attacaggggtgagccactgcgctggcctgaatgccttaaataatgacgtgtctgtcca
. . . . .
cttcattgaaggaagcttctctttctcttatcctgatgggttggtttggtttctttca

      |5201      |5211      |5221      |5231      |5241      |5251
CATGATTTTGAAGTCAGAGGAGATGTGGTCAATGGAAGAAACCACCAAGGTCCAAAGCGA
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

. . . . .
cccaccactctgtattccactcccctttgcagagatgggcccgttcattttgtaagactt
. . . . .
attacatacatcacagtgctagatactttcacacaggttcttttttactcttccatcc
. . . . .
caaccacataaataagtattgtctctactttatgaatgataaaactaagagatttagaga
. . . . .
ggctgtgtaatttggattcccgtctcgggttcagatcttagctgataagtggaagagctg
. . . . .
ggactttaagcagatgagaatcta
```

Exon 20 | Start: 166867 | End: 166921 | Length: 54

```
. . . . .
ccttggcctgattggtgacaaaagtgagatgctcagtccttgaatgacaaagaatgcctg
. . . . .
tagagtgcaggtcaactacatatgcacttcaagaagatcttctgaaatccagtagtggtc
. . . . .
tggacattggactgcttgtccctgggaagtagcagcagaaatcatcaggtggtgaacaga
. . . . .
agaaaaagaaaagctcttcctttttgaaagtctgttttttgaataaaagccaatatctt
. . . . .
ttataactagattttccttctctccattcccctgtccctctctcttcctctcttcttcca

      |5281      |5291      |5301      |5311      |5321      |5331 .
ATCTTCAGGGGGCTAGAAATCTGTGCTATGGGCCCTTCACCAACATGCCACAGgtaag
I  F  R  G  L  E  I  C  C  Y  G  P  F  T  N  M  P  T  D
      |1761                                |1771

. . . . .
agcctgggagaaccccagagttccagcaccagcctttgtcttacatagtgaggattata
. . . . .
agcaagatcccacgatgggggttcctcagattgctgaaatgttctagaggctattctatt
. . . . .
tctctaccactctccaaacaaaacagcacctaaatgttatcctatggcaaaaaaaacta
. . . . .
taccttggtcccccttctcaagagcatgaagggtggttaatagttaggattcagtatgttat
. . . . .
gtgttcagatggcggttgagctgctgttagtgccaacatgttagtgagaaaaatc
```

Exon 21 | Start: 168790 | End: 168863 | Length: 73

.
aattaatggaaattagatctttgattttttttctttcaagcattttatttgagagacta

.
tcaaaccttataccaagtggccttatggagactgataaccagagtacatggcatatcagt

.
ggcaaattgacttaaaatccatacccctactattttaagaccattgtcctttggagcaga

.
gagacagactctccattgagaggtcttgctataagccttcacccggagagtgtagggtta

.
gagggcctgggttaagtatgcagattactgcagtgattttacatctaaatgtccatttta

 |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

.
gagaactgcacatattaaatatactgaggggaagactttttccctctaactctttttcca

.
tatgtccctccccctcctctctgtgactgccccagcatactgtgtttcaacaaatcatca

.
agaaatgatgggct

cccaggagtttgagaccagcctgggcaacatggcaaaacctgtctctaccaaaaataca
.
.aaaaaatttagccaggggtggtggtactgttctgtagtccagctacttaggaggctgaga
.
tggaaggattgcttgagcccaggaggcagaggtggcagtgcgatcacaccactgc
.
actccagcctgggtgacagagcaagaccctgtctcaaaaacaacaaaaaaaaaatgatgaa
.
gtgacagttccagtagtcctactttgacactttgaatgctctttccttctctggggatcca

|5411 |5421 |5431 |5441 |5451 |5461
GGTGTCCACCAATTGTTGTTGTGCAGCCAGATGCCCTGGACAGAGGACAATGGCTTCAT
G V H P I V V V Q P D A W T E D N G F H
|1811 |1821

၁၀၄

Exon 23 | Start: 172182 | End: 173689 | Length: 1507

```
. . . . .
ctaaaaatacaaaaattagctgggtgtgatggcatgtgcctgtaattccagctactcagg

. . . . .
aggcagagacaggagaattgcttgaacccaggaggcggaggttgaatgagccgagattgc

. . . . .
gccatcacactctagcctcggcgacagagcaagactccgtctcaaaaaaaaaaaaaaaaaa

. . . . .
attagcttctacctcattaatcctaagaactcatacaaccaggaccctggagtcgattga

. . . . .
ttagagcctagtccaggagaatgaattgacactaatctctgcttggttctctgtctcca

|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
CACTCTACCAGTGCCAGGAGCTGGACACCTACCTGATACCCGAGATCCCCACAGCCACT
 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
GGCCTTTCCAGGCCCTGGGAGCTCCTCTCACTCTTCAGTCCTTCTACTGTCTGGCTACT

|*121     |*131     |*141     |*151     |*161     |*171
AAATATTTTATGTACATCAGCCTGAAAAGGACTTCTGGCTATGCAAGGGTCCCTTAAAGA

|*181     |*191     |*201     |*211     |*221     |*231
TTTTCTGCTTGAAGTCTCCCTTGGAATCTGCCATGAGCACAAAATTATGGTAATTTTTC

|*241     |*251     |*261     |*271     |*281     |*291
ACCTGAGAAGATTTTAAAAACCATTTAAACGCCACCAATTGAGCAAGATGCTGATTCATTA

|*301     |*311     |*321     |*331     |*341     |*351
TTTATCAGCCCTATTCTTTCTATTCAAGGCTGTTGTTGGCTTAGGGCTGGAAGCACAGAGT
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*361	*371	*381	*391	*401	*411
GGCTTGGCCTCAAGAGAATAGCTGGTTTCCCTAAGTTTACTTCTCTAAAAACCTGTGTTC					
*421	*431	*441	*451	*461	*471
ACAAAGGCAGAGAGTCAGACCCTTCAATGGAAGGAGAGTGCTTGGGATCGATTATGTGAC					
*481	*491	*501	*511	*521	*531
TTAAAGTCAGAATAGTCCTTGGGCAGTTCTCAAATGTTGGAGTGGAACATTGGGGAGGAA					
*541	*551	*561	*571	*581	*591
ATTCTGAGGCAGGTATTAGAAATGAAAAGGAACTTGAAACCTGGGCATGGTGGCTCAGC					
*601	*611	*621	*631	*641	*651
CCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGGGCAGATCACTGGAGGTCAGGAGTTCG					
*661	*671	*681	*691	*701	*711
AAACCAGCCTGGCCAACATGGTGAACCCCATCTCTACTAAAAATACAGAAATTAGCCGG					
*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
GATTTTCAAGAACCGTTTCCAAAGACAGTCTTCTAATTCTCATTAGTAATAAGTAAAA					
*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

GTTGAAGGACTAGGTCTTCCTAGCCCCCAGTGTGCAAGGGCAGTGAAGACTTGATT

 |*1321 |*1331 |*1341 |*1351 |*1361 |*1371
 GTACAAAATACGTTTTGTAAATGTTGTGCTGTAACTGCAAATAAACTTGGTAGCAAA

 |*1381
 CACTTCCAccatgaatgactgttcttgagacttaggccagccgactttctcagagccttt

 tcactgtgcttcagtctcccactctgtaaaatgggggtaatgatagtatctacctcctag

 gatttattgaggcagcttaaataccttttgatattcctgttgctgcaaaacaaattgtt

 gcaaggtcagaagtctgaggtggctcaactgtttctttgtttcaggtttcatgaggccaa

 aataaaggtgttcgcagggcgtgttccttctagaggctctgggtccttgagttctagg

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GBK Parser: Version: 1.1, Version Date: 11/02/2015
Reader: Version: 1, Version Date: 11/02/2015
Writer: Version: 1, Version Date: 11/02/2015
Control: Version: 1, Version Date: 11/02/2015