Gene: BRCA1 - Sequence:  $NG_005905.2$ Transcript:  $NM_007294.3$  - Protein:  $NP_009225.1$ Date: February 19, 2015

 $1^{st}$  line: Base numbering. Full stops for intronic +/-5, 10, 15...  $2^{nd}$  line: Base sequence. lower case Introns, upper case Exons  $3^{rd}$  line: Amino acid sequence. Printed on FIRST base of codon  $4^{th}$  line: Amino acid numbering. Numbered on  $1^{st}$  and increments of 10 Exon 1 | Start: 92501 | End: 92713 | Length: 212  $\verb|tacttatatttaccgaaactggagacctccattagggcggaaagagtgggggattgggac||$  $\verb|ctcttcttacgactgctttggacaataggtagcgattctgaccttcgtacagcaattact|\\$  $\tt gtgatgcaataagccgcaactggaagagtagaggctagaggcaaggcactttatggcaaa$  $\verb|ctcaggtagaattcttcctcttccgtctcttttccttttacgtcatccgggggcagactgg|$ GTACCTTGATTTCGTATTCTGAGAGGCTGCTGCTTAGCGGTAGCCCCTTGGTTTCCGTGG |-159 l-169 l-149 |-139 |-129 |-119 CAACGGAAAAGCGCGGGAATTACAGATAAATTAAAACTGCGACTGCGCGCGTGAGCTCG |-99 |-89 |-79 l-69  $\tt CTGAGACTTCCTGGACGGGGGACAGGCTGTGGGGTTTCTCAGATAACTGGGCCCCTGCGC$ |-39 |-29  ${\tt TCAGGAGGCCTTCACCCTCTGCTCTGGGTAAAGgtagtagagtcccgggaaagggacagg}$  $\tt gggcccaagtgatgctctggggtactggcgtgggagagtggatttccgaagctgacagat$ 

gggtattctttgacggggggtaggggcggaacctgagaggcgtaaggcgttgtgaaccct
tctttaaaaacgtcggctggtcatgaggtcagg
Exon 2   Start: 93869   End: 93967   Length: 98
-19  -9  1  11  21  31  41 TTCATTGGAACAGAAAGAAATGGATTTATCTGCTCTTCGCGTTGAAGAAGTACAAAATGT  M D L S A L R V E E V Q N V  1  11
51  61  71
ttaatttgggattcctatgattatctcctatgcaaatgaacagaattgaccttacatact

${\tt agggaagaaaagacatgtctagtaagattaggctattgtaattgctgattttcttaactg}$
aagaactttaaaaatatagaaaatgattccttgttctccatcca
cctctccttttcaacacaaatcctgtggtccgggaaagacagggactctgtcttgattgg
ttctgcactggggcaggaatctagtttagattaactggc
Exon 3   Start: 102205   End: 102258   Length: 53
agggatcaatataattctgttttgatatctgaaagctcactgaaggtaaggatcgtattc
tctgctgtattctcagttcctgacacagcagacatttaataaata
gccttatgttgactcagtcataacagctcaaagttgaacttattcactaagaatagcttt
atttttaaataaattattgagcctcatttattttctttttctccccccctaccctgctag
81  91  101  111  121  131 .
${\tt TCTGGAGTTGATCAAGGAACCTGTCTCCACAAAGTGTGACCACATATTTTGCAAgtaagt}$
L E L I K E P V S T K C D H I F C K
ttgaatgtgttatgtggctccattattagcttttgtttttgtccttcataacccaggaaa
cacctaactttatagaagctttactttcttcaattaagtgagaacgaaaaatccaactcc
atttcattctttctcagagagtatatagttatcaaaagttggttg

Exon 4   Start: 111451   End: 111528   Length: 77
141   151   161   171   181   191   ATTTTGCATGCTGAAACTTCTCAACCAGAAGAAAGGGCCTTCACAGTGTCCTTTATGTAA F C M L K L L N Q K K G P S Q C P L C K   51   61

gtgtaaaaacttgcagacttatgtaaagtagggctgtatcgccgtgcccccattgtctgt
taatcttgtttttatatt
Exon 5   Start: 113028   End: 113116   Length: 88
gtgtccttaaaaggttgataatcacttgctgagtgtgtttctcaaacaatttaatttcag
221   231   241   251   261   271   366
281   291   301
A F Q L D T G L E Y

tctttcttagtgatacagaaaataatagt
Exon 6   Start: 113723   End: 113862   Length: 139
311   321   331   341   351   361 ATGCAAACAGCTATAATTTTGCAAAAAAAGGAAAATAACTCTCCTGAACATCTAAAAGATG ANSYNFAKKENNSPEHLKDE   1111   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
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$

${\tt agtcctcctgccttagcccccttagtagctgggattacaggcacgcgccaccatgccagg}$
agccactgtgcccggccggt
Exon 7   Start: 118104   End: 118209   Length: 105
actactactattattttgtagagactgggtctcactctgttgcttatgctggtcttgaac
tcctggcctcaagcagtcctgctccagcctcccaaagtgctgggattataggcatgagct
accgctcccagccccagacattttagtgtgtaaattcctgggcattttttccaggcatca
tacatgttagctgactgatgatggtcaatttattttgtccatggtgtcaagtttctcttc
aggaggaaaagcacagaactggccaacaattgcttgactgttctttaccatactgtttag
451   461   471   481   491   501 CAGGAAACCAGTCTCAGTGTCCAACTCTCTAACCTTGGAACTGTGAGAACTCTGAGGACA 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  AAGCAGCGGATACAACCTCAAAAGACGTCTGTCTACATTGAATTGGgtaagggtctcagg K Q R I Q P Q K T S V Y I E L G
K Q R I Q P Q K T S V Y I E L G  171    181
$\tt ttttttaagtatttaataattagttggattccttatcttatagttttgccaaaaatct$

tggtcataatttgtatttgtggtaggcagctttgggaagtgaattttatgagccctatgg
tgagttataaaaaatgtaaaagacgcagttcccaccttgaagaatcttactttaaaaagg
gagcaaaagaggccaggcatggtggctcacacctgtaatcccagcactttgggaggccaa
agtgggtggatcacctgaggtcgggagttcgagaccagcctagcca
Exon 8   Start: 120695   End: 120740   Length: 45
tagaaacgggggtctcactttgttggccaggctggtcttgaactcctaacctcaaataat
ccacccatctcggcctcctcaagtgctgggattacaggtgagagccactgtgcctggcga
agcccatgcctttaaccacttctctgtattacatactagcttaactagcattgtacctgc
551  561  571  581  591 GATCTGATTCTTCTGAAGATACCGTTAATAAGGCAACTTATTGCAGgtgagtcaaagaga S D S S E D T V N K A T Y C S  191
tctttttaaaaatattttaacttttattttaggttcagggatgtatgt

gcataccccacagtttttgtttgctttctttctgaatttctccctcttcccaccttcct
Exon 9   Start: 122062   End: 122138   Length: 76
tgacagttctgcatacatgtaactagtgtttcttattaggactctgtcttttccctatag
601   611   621   631   641   651   1651
661
tgggtagatacagtactgtaattagattattctgaagaccatttgggacctttacaaccc

ttgcccaggctggagtg
Exon 10   Start: 123124   End: 126549   Length: 3425
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
ATGATTTGAACACCACTGAGAAGCGTGCAGCTGAGAGGCATCCAGAAAAGTATCAGGGTA D L N T T E K R A A E R H P E K Y Q G S   251   261
791
851

TCTGTAATAAAAGCAAACAGCCTGGCTTAGCAAGGAGCCAACATAACAGATGGGCTGGAA C N K S K Q P G L A R S Q H N R W A G S |311 GTAAGGAAACATGTAATGATAGGCGGACTCCCAGCACAGAAAAAAAGGTAGATCTGAATG K E T C N D R R T P S T E K K V D L N A |1061 CTGATCCCCTGTGTGAGAGAAAGAATGGAATAAGCAGAAACTGCCATGCTCAGAGAATC D P L C E R K E W N K Q K L P C S E N P |351 |361 |1111 |1131 CTAGAGATACTGAAGATGTTCCTTGGATAACACTAAATAGCAGCATTCAGAAAGTTAATG R D T E D V P W I T L N S S I Q K V N E AGTGGTTTTCCAGAAGTGATGAACTGTTAGGTTCTGATGACTCACATGATGGGGAGTCTG W F S R S D E L L G S D D S H D G E S E AATCAAATGCCAAAGTAGCTGATGTATTGGACGTTCTAAATGAGGTAGATGAATATTCTG GTTCTTCAGAGAAATAGACTTACTGGCCAGTGATCCTCATGAGGCTTTAATATGTAAAA S S E K I D L L A S D P H E A L I C K S |1341 |1361 l1371 GTGAAAGAGTTCACTCCAAATCAGTAGAGAGTAATATTGAAGACAAAATATTTGGGAAAA ERVHSKSVESNIEDKIFGKT CCTATCGGAAGAAGCCACCCCCAACTTAAGCCATGTAACTGAAAATCTAATTATAG Y R K K A S L P N L S H V T E N L I I G

|951

A F V T E P Q I I Q E R P L T N K L K R GTAAAAGGAGACCTACATCAGGCCTTCATCCTGAGGATTTTATCAAGAAAGCAGATTTGG K R R P T S G L H P E D F I K K A D L A |511 |521 l 1581  ${\tt CAGTTCAAAAGACTCCTGAAATGATAAATCAGGGAACTAACCAAACGGAGCAGAATGGTC}$ V Q K T P E M I N Q G T N Q T E Q N G Q |531 |1651 |1661 |1681 AAGTGATGAATATTACTAATAGTGGTCATGAGAATAAAACAAAAGGTGATTCTATTCAGA V M N I T N S G H E N K T K G D S I Q N ATGAGAAAATCCTAACCCAATAGAATCACTCGAAAAAGAATCTGCTTTCAAAACGAAAG E K N P N P I E S L E K E S A F K T K A |571 |581 CTGAACCTATAAGCAGCAGTATAAGCAATATGGAACTCGAATTAAATATCCACAATTCAA E P I S S S I S N M E L E L N I H N S K |591 

AAGCACCTAAAAAGAATAGGCTGAGGAGGAAGTCTTCTACCAGGCATATTCATGCGCTTG
A P K K N R L R R K S S T R H I H A L E

l 1851

I 1861

| 1991 | 2001 | 2011 | 2021 | 2031 | 2041

- GAAACCTACAACTCATGGAAGGTAAAGAACCTGCAACTGGAGCCAAGAAGAGTAACAAGC
  N L Q L M E G K E P A T G A K K S N K P | 671 | 681
- |2111 |2121 |2131 |2141 |2151 |2161 ATGCACCTGGTTCTTTTACTAAGTGTTCAAATACCAGTGAACTTAAAGAATTTGTCAATC 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 CTAGCCTTCCAAGAGAAAAAAGAAGAAGAAACTAGAAACAGTTAAAGTGTCTAATAATG S L P R E E K E E K L E T V K V S N N A |731 |741

- |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
- |2531 |2541 |2551 |2561 |2571 |2581 GCATAGAAATGGAAGAAAGTGAACTTGATGCTCAGTATTTGCAGAATACATTCAAGGTTT

- I E M E E S E L D A Q Y L Q N T F K V S |851 | |861

- |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

- | 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

|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

|3311 |3321 |3331 |3341 |3351 |3361 AGCATCCTGAAATAAAAAGCAAGAATATGAAGAAGTAGTTCAGACTGTTAATACAGATT H P E I K K Q E Y E E V V Q T V N T D F |1111 | 1121

|3491 |3501 |3511 |3521 |3531 |3541 GTTTTGCTGAAAATGACATTAAGGAAAGTTCTGCTGTTTTTAGCAAAAGCGTCCAGAAAG F A E N D I K E S S A V F S K S V Q K G |1171 | 1181

| 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 | 1221

367	1		136	81		36	91		- 1	370	1		37	11		37	21		
CCTG	CTT	CCA	ACA	CTT	GTT/	ATTT	GG.	ΓΑΑ	AGT.	AAA	CAA	TAT	ACC	TTC'	TCAC	TCT	AC.	ΓAG	GC
С	F	Q	Н	L	L	F		K	V	N	N	Ι	P	S	Q		T	R	Н
						12	31									12	41		
1373			37		<b>aa .</b>									71		37			~ .
ATAG	CAC	CGT	TGC	TAC	CGA											ATTA	TC		GA
S	T	V	Α	T	Ε	C	L	S	K	N	T	Ε	Ε	N	L	L	S	L	K
						12	51									12	61		
379			38											31		38			
AGAA	TAG	CTT	AAA'	TGA	CTG											CAG	GA/	ACA'	TC
N	S	L	N	D	С	S  12		Q	V	Ι	L	A	K	A	S	Q  12		Н	Н
385	1		38	61		38	71		1.	388	1		38	91		39	01		
ACCT	TAG	TGA	GGA										ттс	ACA	GTGC	CAGT	GA	ΑТТ	GG
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						112	31									110	ΟI		
391	1		139	21		39	31		- 1.	394	1		39	51		39	61		
AAGA	CTT	GAC	TGC.	AAA	TAC	AAAC	AC	CCA	GGA'	TCC	TTT	CTT	GAT	TGG'	TTC	TCC	AAA	ACA.	AA
	L				Т											S			М
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						110										110	21		
397			39			39								11		140			
TGAG	GCA	TCA	GTC'	TGA	AAG(	CCAG	GG	AGT"	TGG'	TCT	GAG	TGA	CAA	GGA.	ATTO	GTT	TCI	AGA'	TG
R	Η	Q	S	Ε	S	Q	G	V	G	L	S	D	K	E	L	V	S	D	D
						13	31									13	41		
1403	1		140	41		140	51		- 1	406	1		40	71		140	81		
ATGA	AGA	AAG	AGG.	AAC	GGG	CTTG	GA	AGA.	AAA	TAA	TCA	AGA	AGA	GCA.	AAG	CATG	GA:	ГТС	AA
Ε	Ε	R	G	T	G	L	E	E	N	N	Q	Ε	E	Q	S	M	D	S	N
						13	51									13	61		
						•													
1409	1																		
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atgt																		ctg	ct
5-		5		55										3		33		J	
tctt																			ca

gctaggacgtcatctttgactgaatgagctttaacatcctaattactggtggacttactt
tgtgac
Exon 11   Start: 126952   End: 127040   Length: 88
4101
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$

gcctacacatacactgcctagctcattgtagcatactaaatactgattttaatgaataag
ctaaaccttcgaaacccatttgctaatcc
Exon 12   Start: 135409   End: 135580   Length: 171
4191
4251
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$

tgttggccaaacactgatatcttaagcaaaattctttccttcc
agagtaaggacctagctccaacattttatgatccttgctcagcacatgggtaattatgga
Exon 13   Start: 141370   End: 141496   Length: 126
tcagggccaaggaatatagattttttttttcagccttgtctcagctgggtgtctttattt
${\tt actctgtcttaaagtgttccttttattatcattatttttttaatcattgaattccatt}$
4361  4371  4381  4391  4401  4411 CAGTATTAACTTCACAGAAAAGTAGTGAATACCCTATAAGCCAGAATCCAGAAGGCCTTT
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

$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
tcaaggc
Exon 14   Start: 143463   End: 143653   Length: 190
4491

C	т∕тт				45  AAAC														
		Q				Y								K	٧		D		
GGA E			GCT	GGA.	46 AGAG E  15	TC: S	[GG	GCC	ACA	CGA	TTT	GAC	GGA		ATC S	TTA	CTT( L	GCC.	
			AGg		tatt														tga
atc	cta	.cat	aaa	gat	atto	tgg	gtt	aac	caa	ctt	tta	gat	gta	.cta	gtc	tat	cat	gga	cac
					ttaa														tta
					aact													aat	gat
					gtct														tgg
ctc	acg	cct	gt																
Exo	n 1	5	St	art	: 14	<u>1</u> 674	16	l E	nd:	14	705	6	Le	ngt	h:	310			
					ttcc														gat
					gcat														
					cttt														

tcat	gta	ccca	attt	tt	ctc	ttaa	acc	taa	ctt	tatt	ggtc	tttt	taa	ttc	tta	aca	gag	ac
caga	act	ttgt	aat	tc	aaca						aatta						ttc	ag
	•	681	DED 4.								471							
											CTTCT							
G			Y	L	E	S	G	Τ	S	L	F S		Ъ	Р	E	S	D	Ρ
	1	561									157	1						
	14	741		- [-	475	1		1476	61		477	1	1.	478	1		47	91
CTTC	TGA	AGAC	CAG <i>I</i>	AGC	CCC	AGA(	GTC	AGC'	TCG'	TGT]	rggca.	ACAT	ACC	ATC'	TTC	AAC	CTC'	TG
S	E	D	R	Α	P	Ε	S	Α	R	V	G N	I	P	S	S	T	S	Α
	1	581									159	1						
											483							
											rgccc.							
L		V	P	Q	L	K	V	Α	E	S	A Q		Р	Α	Α	A	H	T
	16	601									161	1						
	48	361		- [-	487	1		1488	31		1489	1	1.	490	1		49	11
CTAC	TGA	ГАСТ	[GC]	rgg	GTA:	ΓΑΑΊ	ГGС	AAT	GGA.	AGA <i>I</i>	AGTG	TGAG	CAG	GGA	GAA(	GCC.	AGA.	ΑТ
Т	D	Т	Α	G	Y	N	Α	М	Ε	E	s v	S	R	Ε	K	P	Ε	L
	16	621									163	1						
	149	921		1.	493 <sup>-</sup>	1		1494	41		495	1	1.	496	1		149	71
TGAC			ACA								TCCA							
Т											S M							
-		641	•	_	10	·	••	••	10	••	1165		•	٥	ŭ	_	-	•
	•	981																
			[gte	gag	tgta	atco	cat	atg	tat	ctc	cctaa	tgac	taa	gac	tta	aca	aca	tt
Ε	E																	
	16	661																
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ctgg	aaa	gagt	ttt	cat	gta	ggta	att	gtca	aat <sup>.</sup>	taat	caacc <sup>.</sup>	taga	ıgga	aga	aat	cta	gaa	aa
caat	caca	agtt	cte	gtg	taa	ttta	aat	ttc	gat	tact	taatt	tctg	gaaa	att <sup>.</sup>	tag	atc <sup>.</sup>	tag	at
aaag	ctat	tagt	gtg	gga	tta	tttt	tat	gtat	tat	ttad	cttga	gaaa	ata	att	atta	aaa	tat	ta
ot.com	· aaa:	aoct	:a+s		tt o		a+ m	· atat	tao	σact	ttcg	aa++		a++·	· ttc:	c++·	· tc+	a+
5 556	uuu	~5 · ·	- 400		2 2 E	5000	~ ೮	aua	وعد	5401	Jucg	السال	65ª	400	550		300	40

· · · · ctgtaaaagca

Exon 16 | Start: 150289 | End: 150376 | Length: 87  $\tt gagacttcaggtgtcttagaattttttaaatgtaccctttctgagaaaaacagagactta$  $\tt gtattctgagctgtgtgctagaggtaactcatgataatggaatatttgatttaatttcag$ 4991 |5001 |5011 |5021 |5031 15041  ${\tt ATGCTCGTGTACAAGTTTGCCAGAAAACACCACATCACTTTAACTAATCTAATTACTGAA}$ MLVYKFARKHHITLTNLITE 1671 |5071 . . . . |5051 |5061  ${\tt GAGACTACTCATGTTATGAAAACAGgtataccaagaacctttacagaataccttgca}$ ETTHVVMKTD |1691 . . . •  $\verb|tctgctgcataaaaccacatgaggcgaggcacggtggcgcatgcctgtaatcgcagcact|$  $\verb|ttgggaggccgaggcagatcacgagattaggagatcgagaccatcctggccagcat|\\$  $\tt ggtgaaaccccgtctctactaaaaaaataaaaaattagctgggtgtggtcgcgtgcgcct\\$  $\tt gtagtcccagctactcgtgaggctgaggcaggagaatcacttgaaccggggagatggagg$  ttgcagtgagccgagatcatgccactgc Exon 17 | Start: 154033 | End: 154110 | Length: 77 tata atggagat ctatagctagccttggcgtctagaagatgggtgttgagaagagggagt $\tt ggacagatatttcctctggtcttaacttcatatcagcctcccctagacttccaaatatcc$  $\verb|atacctgctggttata| attagtggtgttttcagcctctgattctgtcaccaggggtttta|$ ga at cata a at ccagatt gat ctt ggg ag tgt a aaa aactg ag gct ctt tag ctt ctt ag $\tt gacag cacttcct gatttt gtttt caacttct aat ccttt gag tgttttt cattct g cag$ |5081 |5091 |5101 |5111 |5121 |5131  $\tt ATGCTGAGTTTGTGTGAACGGACACTGAAATATTTTCTAGGAATTGCGGGAGGAAAAT$ A E F V C E R T L K Y F L G I A G G K W 1701 |5141 |5151 . . . . . .  $\tt GGGTAGTTAGCTATTTCTgtaagtataatactatttctcccctcctcctttaacacctc$ V V S Y F W . . . . . . • . aga att g cattttta caccta acgttta acaccta acgtttt ttgctgatgctgagtctga $\tt gttaccaaaaggtctttaattgtaatactaaactacttttatctttaatatcactttgtt$  ${\tt cagata} a {\tt gctggtgatgctgggaaaatgggtctcttttataactaataggacctaatctg}$ 

 $\verb|ctcctag| catagram| cat$ 

gcagcaggcaaacttata

Exon	18	ı	Sta	rt:	15	461	1	En	.d:	154	651	ı	Len	gth	1: 4	10			
ataa	cta	ata	ıgga	cct	aat	ctg	cto	cta	.gca	atg	tta	gca	tat	gag	cta	agg	gati	ttat	tt
· aata	· gtc	ggc	agg	aat	сса	tgt	gca	· igca	ggc.	aaa	.ctt	ata	· atg	ttt	aaa	· att	aaa	cato	:aa
	0 0	00 -	00			-6-	0	-6	- 00				0						
•	•		•										•						
ctct	gtc	tcc	aga	agg	aaa	ctg	ctg	gcta	.caa	.gcc	tta	tta	aag	ggc	tgt	gg	ctti	taga	ıgg
						_													
gaag	gac	ctc	tcc	tct	gtc	att	ctt	cct	gtg	ctc	ttt	tgt	gaa	tcg	cte	gac	ctct	tcta	tc
	+ ~~	222			~++		c+c			«+ »		+ ~+		++	+ > +	•	+ c+	•	· ~ ~
tccg	uga	aac	ıgag	cac	guu		عات	getg	lau	gua	acc	ugu	, С		lai	ga			ag
			161			517			51				191						
GGGT													Ggt	aag	gtac	ctt	gat	gtta	ıca
V	T	•	S .721	Ι	K	E	R	K	M	L	N	E	721						
		11	. / 21									1	.731						
aact	aac	cag	gaga	tat	tca	ttc	agt	cat	ata	gtt	aaa	aat	gta	ttt	gct	tc	ctt	ccat	ca
atgc	· acc	act	ttc	ctt	aac	aat	gca	· icaa	att	ttc	cat	gat	aat	gae	rgat	.ca	tcaa	· agaa	tt
8-							0					0		06				-6	
•																			
atgc	agg	cct	gca	ctg	tgg	ctc	ata	cct	ata	atc	cca	gcg	gctt	tgg	gag	ggc	tgag	ggcg	gct
						_													
tgga	tca	cct	gat	gtc	ggg	agt	tca	aga	.cca	.gcc	tga	cca	aca	tgg	gaga	aaa	ccc	gtt	tc
												<b>.</b>							
tact	aaa	aat	aca	aaa	ιτa	gcc	ggg	gctt	ggt	ggc	act	ugo	C						

 $\verb|ccacgcccaactaatttttgtatttttagtagagatgaggtttcaccatgttggtcaga|\\$  $\verb|ctggtgtcgaactcctgacctcaagtgatctgcctcgctctcagtctcccaaagtgctagga|\\$  $\verb|ttacaggggtgagccactgcgcctgaatgccttaaatatgacgtgtctgctccac|$ |5201 |5211 |5221 |5231 |5241  ${\tt CATGATTTGAAGTCAGAGGAGATGTGGTCAATGGAAGAACCACCAAGGTCCAAAGCGA}$ H D F E V R G D V V N G R N H Q G P K R 1741 |1751 |5261 |5271 . . . . . . .  $\tt GCAAGAGAATCCCAGGACAGAAAGgtaaagctccctccctcaagttgacaaaaatctcac$ A R E S Q D R K  $\verb|cccaccactctgtattccactcccctttgcagagatgggccgcttcattttgtaagactt|\\$  $\verb|attacatacatacacagtgctagatactttcacacaggttcttttttcactcttccatcc|\\$  ${\tt caaccacataaataagtattgtctctactttatgaatgataaaactaagagatttagaga}$  $\tt ggctgtgtaatttggattcccgtctcgggttcagatcttagctgataagtggaagagctg$ . . ggactttaagcagatgagaatcta

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

5281  5291  5301  5311  5321  5331 . ATCTTCAGGGGGCTAGAAATCTGTTGCTATGGGCCCTTCACCAACATGCCCACAGgtaag I F R G L E I C C Y G P F T N M P T D  1761   1771
agcctgggagaaccccagagttccagcaccagcctttgtcttacatagtggagtattata
tctctaccactctccaaacaaaaaaacagcacctaaatgttatcctatggcaaaaaaaa
taccttgtcccccttctcaagagcatgaaggtggttaatagttaggattcagtatgttat

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

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

attaatggaaattagatctttgatttttttttttttctttc
5341   5351   5361   5371   5381   539 ATCAACTGGAATGGATGGTACAGCTGTGTGGTGCTTCTGTGGTGAAGGAGCTTTCATCAT Q L E W M V Q L C G A S V V K E L S S F   1781   1791
agaaatgatgggct

Exon 22 | Start: 170281 | End: 170341 | Length: 60

tgacagttccagtagtcctactttgacactttgaatgctctttccttcc
5411
g

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

taaaaatacaaa	 aattagctggg	 tgtgatggca	 tgtgcctgta	 attccagct	actcagga
ggcagagacagg	 agaattgcttg	· · · aacccaggag	 gcggaggttg	aatgagccg	agattgcg
ccatcacactct	 agcctcggcga	· · cagagcaaga	 ctccgtctca	 aaaaaaaaa	aaaaaaaa
ttagcttctacc	 tcattaatcct	 aagaactcat	 acaaccagga	 .ccctggagt	cgattgat
tagagcctagtc	caggagaatga	 attgacacta	 atctctgctt	 gtgttctct	gtctccag
5471 CAATTGGGCAGA' I G Q M	C E A	CCTGTGGTGA			CAGTGTAG
CACTCTACCAGT	Q E L		TGATACCCCA		CAGCCACT
5591 ACTGACTGCAGC	+11 CAGCCACAGGT				
+61 GGCCTTTCCAGG	+71 CCCTGGGAGCT				
+121 AAATATTTTATG	+131 TACATCAGCCT				
+181 TTTTCTGCTTGA		+201 GAAATCTGCC			
+241 ACCTGAGAAGAT	+251  TTAAAACCAT	+261 TTAAACGCCA	+271 CCAATTGAGC	+281 AAGATGCTG	+291 ATTCATTA
+301 TTTATCAGCCCT	+311 ATTCTTTCTAT	+321 TCAGGCTGTT	+331 GTTGGCTTAG	+341 GGCTGGAAG	+351 CACAGAGT
+361	+371	+381	+391	+401	+411

GGCTTGGCCTCAA	GAGAATAGCT	GGTTTCCCTA	AGTTTACTTC	TCTAAAACCC	TGTGTTC
+421	l+431	+441	+451	+461	+471
ACAAAGGCAGAGA	GTCAGACCCT	TCAATGGAAG	GAGAGTGCTT	GGGATCGATT	TATGTGAC
+481	+491	l+501	l+511	l+521	l+531
TTAAAGTCAGAAT	AGTCCTTGGG	CAGTTCTCAA	ATGTTGGAGT	GGAACATTGG	GGAGGAA
			+571		
ATTCTGAGGCAGG	TATTAGAAAT	GAAAAGGAAA	CTTGAAACCT	GGGCATGGTG	GCTCACG
l+601					
CCTGTAATCCCAG	CACTTTGGGA	GGCCAAGGTG	GGCAGATCAC	TGGAGGTCAG	GAGTTCG
			l+691		
AAACCAGCCTGGC	CAACATGGTG	SAAACCCCATC	CTCTACTAAAA	ATACAGAAAT	TAGCCGG
+721					
TCATGGTGGTGGA	CACCTGTAAT	CCCAGCTACT	CAGGTGGCTA	AGGCAGGAGA	ATCACTT
			+811		
CAGCCCGGGAGGT	GGAGGTTGCA	IGTGAGCCAAG	FATCATACCAC	GGCACTCCAG	CCTGGGT
•	•	•	+871	•	•
GACAGTGAGACTG	TGGCTCAAAA	AAAAAAAAAA	AAAAAGGAAA	ATGAAACTAG	AAGAGAT
+901					
TTCTAAAAGTCTG	AGAIAIAIII	GCIAGAIIIC	JIAAAGAAIGI	GIICIAAAAC	AGCAGAA
			+991		
GATTTTCAAGAAC					
+1021 TGTTTATTGTTGT			+1051		
+1081 TGAATATTTCATA			+1111		
IGAATATITOATA	ICIAIAAAAI	GACAGATOCC	CACCAGGAAGG	MAGCIGIIGC	,1110111
+1141 GAGGTGATTTTT			+1171		
+1201 GCTTGCTGAAGGA			+1231 ACCCATTATO		

+1291

+1301

+1311

|+1281

 $\tt GTTGGAAGGACTAGGTCTTCCCTAGCCCCCCCAGTGTGCAAGGGCAGTGAAGACTTGATT$ 

+1271

+1261

GTAC.	+1321 AAAATAC					
CACT"	+1381 TCCAcca				ttctc	
	tgtgctt					
	tattgag					
	ggtcaga					
	aaggtgt					
acta	agat					

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