Gene: BRCA1 - Sequence: NG_005905.2 Transcript: NM_007294.3 - Protein: NP_009225.1 Date : March 2, 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
-229 -219 -209 -199 -189 -179 GTACCTTGATTTCGTATTCTGAGAGGCTGCTTGCTTAGCGGTAGCCCCTTGGTTTCCGTGC
-169 -159 -149 -139 -129 -119 CAACGGAAAAGCGCGGGAATTACAGATAAATTAAAACTGCGACTGCGCGGGGTGAGCTC
-109 -99 -89 -79 -69 -59 CTGAGACTTCCTGGACGGGGGACAGGCTGTGGGGTTTCTCAGATAACTGGGCCCCTGCGG
-49 -39 -29

Exon 2	Sta:	rt: 9	3869	End:	9396	7 Le	ength	: 98			
aggctaco	· cacca	·	ccggtc	agtca	ctcct	ctgta	· gcttt	ctctt	tcttgg	agaa	ag
· · · gaaaagad	cccaa		tggcag	· caata	tgtgaa	aaaaat	ttcag	aattt	atgttg	tcta	at
· · · tacaaaaa	agcaa	· cttct	agaatc	tttaa	aaata	• aagga	cgttg	tcatt	agttct	ttgg	tt
 tgtattat	tctaa	· aaacc			aaatti				tgataa	aatg	aa
gttgtcat	ttta	· taaac	ctttta	• aaaag	atata1	tatata	atgtt	tttct	aatgtg	ttaa	ag
-19 TTCATTGO				ATTTA	11 TCTGCT	rcttc(GCGTT	GAAGA	31 AGTACA V Q 11		41 GT V
CATTAATO I N <i>F</i>			61 AAATCT I L 21	TAGAG			Ggtaa	• gtcag	 cacaag	agtg	· ta
ttaattto	gggati	tccta	tgatta	tctcc	tatgc	aaatga	aacag	aattg	 acctta	cata	ct
• agggaaga	aaaaga	• acatg	tctagt	aagat	taggc	cattg!	taatt	gctga	 ttttct	taac	tg
aagaactt	ctaaaa		• agaaaa					actct	 gcctct	ccca	ct
cctctcct	Ettte	· aacac	aaatcc	tgtgg	tccgg	gaaaga	acagg	gactc	 tgtctt	gatt	gg •
ttctgcad	ctggg	gcagg	aatcta	gttta	gatta	actgg	• C				

Exon 3 Start: 102205 End: 102258 Length: 53	
	l
	;
	ſ
	:
	ſ
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	ı
cacctaactttatagaagctttactttcttcaattaagtgagaacgaaaaatccaactc	;
	ſ
gtaaagttttgacatatattatctttttttttttttttt	ſ

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
201 211
ttatgggacatctgccttatacaggtattagaaacttactgcctttctctaatgcttcta
taatcttgttttatatt

Exon 5 Start: 113028 End: 113116 Length: 88	
	Ictgtgga
	• tgacaaa
	gattata
	atttcag
221 231 241 251 261 GAGCCTACAAGAAAGTACGAGATTTAGTCAACTTGTTGAAGAGCTATTGAAAA S L Q E S T R F S Q L V E E L L K I 81	TCATTTG
281 291 301	
	· · · · · · · · · · · · · · · · · · ·
	· · · itgtgctt
tctttcttagtgatacagaaaataatagt	

Exon																						
· taca										tgaa										ctt	agt	gt
taag										taad												CC
tagg	agt	aa	.aa	tti						ttta											ggt	aa
cctt										aaag											ctt	tg
atta	taa	ıtt	ca	ta						acto												ag
ATGC A	AAA N		GC				ГТG		AA.	AAA(K		AAA N		СТС		СТ				AAA		
AAGT V	TTC S			AT(AA(GΤΑ	TG	GG	CTA(Y	CAGA	AAA N		TGC		AA			41 TCT L	ACA		421 TG E 141
AACC P	CG <i>P</i> E		AT	CC:		CT:				acca											ctt	• tt
cttt	ttt	tt	tt	cti	tt	ttt	tt	tti	tg		Egga							ccc	agg	cta	gaa	• gc
agto									ag		ctgg	gga	tta	cag	ggc	ac	gcc	gcc	acc	atg	cca	gg
ctaa																						
ctcc										ccto												
agcc																						

Exc	n	7	St	tart	: 1	181	04	I	End:	11	820	9	Len	ngth	n: 1	105				
act	ac	tac	ctat	ctat	ttt	gta	Igag	act	Eggg	tct	cac	tct	gttg	gctt	tato	gct	ggt	· ctt	gaac	:
· tcc	ctg	· gc	ctca	· aagc	agt	cct	.gct	CC	agcc	tcc	caa	agt	gctg	idda	atta	ata		atga	agct	
acc	cgc	· tco	cca	gccc	cag	aca	ıttt	tag	gtgt	gta	aat	tcc	tggg	gcat	cttt	ctt	cca	ggca	atca	L
tac	cat	gtt	ago	ctga	ctg	atg	_T atg	gto	caat	tta	ttt	tgt	ccat	ggt	igto	caa	gtt	tcto	cttc	:
· agg	gag	gaa	aaaq	gcac	aga	act	.ggc	caa	acaa	ttg:	ctt	gac	tgtt	ctt	ctad	cca	tac	tgtt	taç	ſ
CAG Q	GGA E	AAC T	CCA(451 GTCT(L 151	CAG S	TGI	461 CCA Q	AC:	ГСТС	47: TAA(N	CCT		48 AACI T 16	GT(V	GAG <i>i</i> R		491 TCT L			501
AAG K			GGA:	511 ΓΑCΑ Ω 171	ACC P					53: TGT(V	СТА	CAT I	54 TGAA E 18	ATT(L	GGgt G	caa	.	tcto	cago	Г
ttt	tt	taa	agta	· attt	aat	aat	.aat	tgo	ctgg	att	cct	tat	ctta	ıtaç	gttt	tg	cca	aaaa	atct	:
tgg	gtc	ata	aatt	ctgt:	att	tgt	.ggt	ago	gcag	ctt	tgg	• gaa	gtga	att	ctta	atg	agc	ccta	atgg	ſ
tga	ıgt	tat	aaa	aaaa	tgt	aaa	laga	cg	cagt	tcc	cac	ctt	gaaç	gaat	cctt	cac	ttt	aaaa	aagg	ſ
· gaç	gca	.aaa	agag	ggcc	agg	cat	.ggt	gg	ctca	cac	ctg	· taa	tccc	cago	cact	tt		aggo	ccaa	L
agt	gg	gtg	ggat	cac	ctg	agg	ıtcg	gga	agtt	cga	gac	cag	ccta	ıgco	ca					

Exon	8	Star	t:	1206	595	Er	nd:	120	740	I	engt	th:	45		
tagaa															aataat
ccacc															· :ggcga
														ttgta	acctgo
														agtad	catttt
tttaa														aattt	:tttag
GATCT	'GAT'	CTTC S S	TGA	AGA:	TACC(GTTA	AAT <i>I</i>	AAGG	CAAC	TTA	TTG	CAG		gtcaa	• aagaga
														gatgt	ittctc
														agttt	:gttat
														cgggt	tactaa
· gcata															ettect
ccctc	:aagt	aggc	tgg	tgtt	·	ccaq	gact	agaa	atca	tgg	rtati	tgg			

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

Exon	10		Sta	rt:	12.	3124	Ei	nd:	126	549)	Len	gth	: 3	425			
agtt	ttc	tga	tgg	cca	atc	· tgct	ctta	atto	cact	.ctt	aga	.cgt	tag	aga	aata	ggtg	rtgg	ſ
tttc	tgc	ata	ggg	aaa	att	ctga									• ggaa	ataa	ıtct	
aggt	• aaa	tag	• gaa	tta	aat		gagt						gta	tac	ttgg	tagt	tta	L
tgag	gtt	agt	· ttc	tct	aat	atag	ccag	ttgg	gttg	gatt	tcc	acc	tcc	aag	gtgt	atga	ıagt	
atgt	att	ttt	tta	atg	aca	attc	agtt	tttg	gagt	acc	ttg	rtta	ttt	ttg	tata	tttt	cag	ſ
671 CTGC		$T \subset X$	68		TCN	69:		 T				71		тсл	72		. א יד א	
A	C	E	F	S	E		V C		N	Т	E	Н	Н	Q		S N		
731		~ ~ ~	74				l					77		.	78			
ATGA D	L	gaa N	T	T	E E	JAAG(K 1 25:	R A			R R	H	P	AGA E	.aaa K	Y 26	Q G		
791			80		007		1					83		maa	84			
GTTC S	V	S S	N N	L	GCA:	V 1	E P		G	T	N N	Т	Н	A A	S 28	S I		
851 AGCA		~ 7. 7.	186			87:		 				189		7. (2.7.	190		ית הי	,
H H	E	gaa N	S		L		L T			R	M	N	.1G1 V	E E		A E		
911			92			93:			941			195			196			
TCTG																		
С	N	K	S	K	Q	P (A	R	S	Q	Н	N	R	W 132	A G 1	; S	,

```
| 991 | 1001 | 1011 | 1021
     | 981
GTAAGGAAACATGTAATGATAGGCGGACTCCCAGCACAGAAAAAAAGGTAGATCTGAATG
 K E T C N D R R T P S T E K K V D L N A
               |331
                                       1341
|1031 |1041
              |1051 |1061
                             |1071
                                      |1081
CTGATCCCCTGTGTGAGAGAAAGAATGGAATAAGCAGAAACTGCCATGCTCAGAGAATC
 D P L C E R K E W N K Q K L P C S E N P
               1351
                                       1361
11091
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              |1111
                      |1121
                             |1131
                                      11141
CTAGAGATACTGAAGATGTTCCTTGGATAACACTAAATAGCAGCATTCAGAAAGTTAATG
 R D T E D V P W I T L N S S I Q K V N E
               |371
                                       1381
|1151
      |1161
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                              |1191
                                      |1201
AGTGGTTTTCCAGAAGTGATGAACTGTTAGGTTCTGATGACTCACATGATGGGGAGTCTG
 W F S R S D E L L G S D D S H D G E S E
               |391
                                       1401
|1211 |1221
               11231
                      11241
                             |1251
                                      |1261
AATCAAATGCCAAAGTAGCTGATGTATTGGACGTTCTAAATGAGGTAGATGAATATTCTG
 S N A K V A D V L D V L N E V D E Y S G
               |411
|1271
      |1281
              |1291
                      |1301
                             11311
                                      11321
GTTCTTCAGAGAAAATAGACTTACTGGCCAGTGATCCTCATGAGGCTTTAATATGTAAAA
S S E K I D L L A S D P H E A L I C K S
               |431
                                       1441
                                      |1381
11331
      |1341
              |1351
                      |1361 |1371
GTGAAAGAGTTCACTCCAAATCAGTAGAGAGTAATATTGAAGACAAAATATTTGGGAAAA
 ERVHSKSVESNIEDKIFGKT
               |451
                                      |461
|1391 |1401
              |1411
                      |1421
                             |1431
                                      |1441
CCTATCGGAAGAAGCCACCCCAACTTAAGCCATGTAACTGAAAATCTAATTATAG
 Y R K K A S L P N L S H V T E N L I I G
               |471
                                      |481
              11471
11451
      11461
                     |1481
                              11491
A F V T E P Q I I Q E R P L T N K L K R
               1491
                                       1501
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| 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
                                         1521
|1571 |1581
               11591
                       |1601
                               |1611
                                         11621
CAGTTCAAAAGACTCCTGAAATGATAAATCAGGGAACTAACCAAACGGAGCAGAATGGTC
 V Q K T P E M I N Q G T N Q T E Q N G Q
                1531
       |1641
               11651
                        |1661
                                |1671
11631
                                         11681
AAGTGATGAATATTACTAATAGTGGTCATGAGAATAAAACAAAAGGTGATTCTATTCAGA
V M N I T N S G H E N K T K G D S I Q N
                |551
                                         1561
|1691
      |1701
                |1711
                       |1721
                                |1731
                                         |1741
ATGAGAAAATCCTAACCCAATAGAATCACTCGAAAAAGAATCTGCTTTCAAAACGAAAG
 E K N P N P I E S L E K E S A F K T K A
                |571
                                         1581
|1751 |1761
                |1771
                        11781
                               |1791
                                         11801
CTGAACCTATAAGCAGCAGTATAAGCAATATGGAACTCGAATTAAATATCCACAATTCAA
 E P I S S S I S N M E L E L N I H N S K
                |591
|1811
       |1821
               |1831
                        |1841
                                11851
                                        11861
AAGCACCTAAAAAGAATAGGCTGAGGAGGAAGTCTTCTACCAGGCATATTCATGCGCTTG
 A P K K N R L R R K S S T R H I H A L E
                1611
11871
       |1881
                |1891
                        |1901
                                |1911
                                         11921
AACTAGTAGTCAGTAGAAATCTAAGCCCACCTAATTGTACTGAATTGCAAATTGATAGTT
 L V V S R N L S P P N C T E L Q I D S C
                |631
|1931 |1941
                |1951
                        |1961
                               |1971
                                         |1981
GTTCTAGCAGTGAAGAGATAAAGAAAAAAAGTACAACCAAATGCCAGTCAGGCACAGCA
 S S S E E I K K K K Y N Q M P V R H S R
                |651
                                         |661
       12001
               12011
                       12021
11991
                                12031
GAAACCTACAACTCATGGAAGGTAAAGAACCTGCAACTGGAGCCAAGAAGAGTAACAAGC
 N L Q L M E G K E P A T G A K K S N K P
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                                          1681
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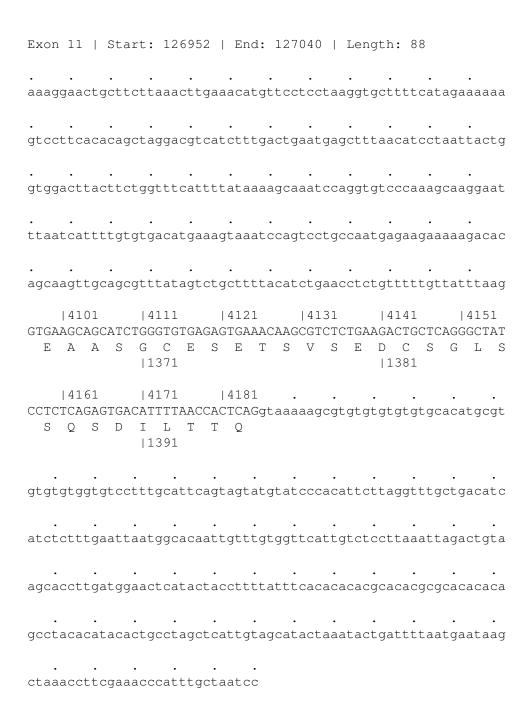
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N E	Q	Т	S K	R H 691	D	S	D	Τ	F	Р	Ε	L	K L 701	Τ	N
2111		1212	1	2131		1	214	1		121	51		12161		
ATGCAC				•								AGA	,		TC
A P	G	S	F T	K C	S	N	Τ	S	Ε	L	K	Ε	F V	N	Р
				711									721		
2171		218	1	2191			220	1		122	11		2221		
CTAGCC	TTCC	AAGA	GAAGA	AAAAGA	AGA	GAA.	ACT	AGA	AAC	AGT	TAA.	AGT	GTCTAA	AATA	TG
S L	Р	R	E E	K E 731	Ε	K	L	Ε	Τ	V	K	V	S N 741	N	A
2231		224	1	2251	-	1	226	1		22	71		2281		
CTGAAG.	ACCC	Caaa													TG
E D	P	K	D L	M L	S	G	Ε	R	V	L	Q	Τ	E R	S	V
				751									761		
2291		1230	1	2311			232	1		123	31		2341		
TAGAGA	GTAG	CAGT	ATTTC									TCA	GGAAAG	TAT	СТ
E S	S	S	I S	L V 771	Р	G	Τ	D	Y	G	Τ	Q	E S 781	I	S
2351		236	1	2371		-	238	1		23	91		2401		
CGTTAC	TGGA	AGTT	AGCAC	TCTAGG	GAA	.GGC	AAA	AAC	AGA	ACC	AAA	TAA	ATGTGI	GAG	TC
L L	E	V	S T	L G	K	Α	K	Τ	Ε	Р	N	K	C V	S	Q
				791									801		
2411		1242	1	2431		1	244	1		124	51		12461		
AGTGTG	CAGC	ATTT	GAAAA									CAA	AGATAA	TAG	AA
C A	A	F	E N	P K	G	L	I	Н	G	С	S	K	D N	R	N
				811									821		
2471		1248	1	2491			250	1		25	11		2521		
ATGACA	CAGA	AGGC	TTTAA	GTATCC	ATT	GGG.	ACA	TGA	AGT	TAA	CCA	CAG'	TCGGGA	AAC	AA
D T	Ε	G	F K	Y P 831	L	G	Н	Ε	V	N	Н	S	R E 841	Τ	S
2531		1254	1	2551		J	256	1		125	71		2581		
GCATAG.															ТТ
ΙE			E S	E L								Τ	F K	V	S
				851									861		

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|2591 |2601 |2611 |2621 |2631
                                      12641
CAAAGCGCCAGTCATTTGCTCCGTTTTCAAATCCAGGAAATGCAGAAGAGGAATGTGCAA
 K R Q S F A P F S N P G N A E E E C A T
               871
                                       1881
|2651 |2661
               |2671 |2681
                             |2691
                                      12701
CATTCTCTGCCCACTCTGGGTCCTTAAAGAAACAAAGTCCAAAAGTCACTTTTGAATGTG
 F S A H S G S L K K Q S P K V T F E C E
               1891
                                       1901
12711
       |2721
              |2731
                      12741
                              12751
                                      12761
Q K E E N Q G K N E S N I K P V Q T V N
                |911
                                       1921
|2771
      12781
               |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
               1931
                                       1941
|2831 |2841
               12851
                      12861
                              12871
                                      12881
GTAGTATCAAAGGAGGCTCTAGGTTTTGTCTATCATCTCAGTTCAGAGGCAACGAAACTG
 S I K G G S R F C L S S Q F R G N E T G
               |951
                                       1961
|2891
      |2901
              |2911
                      |2921
                              12931
                                      12941
GACTCATTACTCCAAATAAACATGGACTTTTACAAAACCCATATCGTATACCACCACTTT
L I T P N K H G L L Q N P Y R I P P L F
               1971
                                       1981
12951
      |2961
               |2971
                      12981
                              |2991
                                      13001
TTCCCATCAAGTCATTTGTTAAAACTAAATGTAAGAAAAATCTGCTAGAGGAAAACTTTG
 PIKSFVKTKCKKNLLEENFE
               1991
                                       11001
|3011 |3021
               |3031
                      13041
                             |3051
                                      |3061
AGGAACATTCAATGTCACCTGAAAGAGAAATGGGAAATGAGAACATTCCAAGTACAGTGA
 E H S M S P E R E M G N E N I P S T V S
               11011
                                       11021
13071
      13081
              13091
                      |3101
                             13111
                                      13121
GCACAATTAGCCGTAATAACATTAGAGAAAATGTTTTTAAAGAAGCCAGCTCAAGCAATA
 T I S R N N I R E N V F K E A S S S N I
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                                       11041
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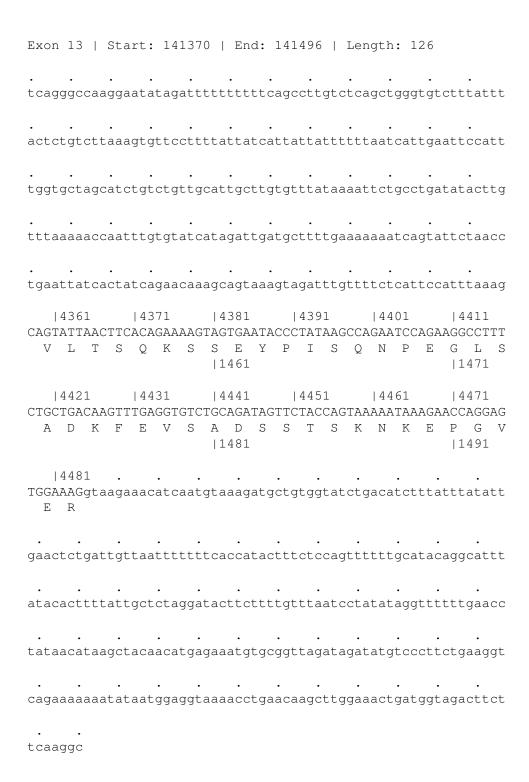
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                                         11061
|3191 |3201
               |3211 |3221
                               |3231
                                        13241
GTGATGAAAACATTCAAGCAGAACTAGGTAGAAACAGAGGGCCAAAATTGAATGCTATGC
 D E N I Q A E L G R N R G P K L N A M L
                11071
                                         11081
13251
       |3261
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                       |3281
                               |3291
TTAGATTAGGGGTTTTGCAACCTGAGGTCTATAAACAAAGTCTTCCTGGAAGTAATTGTA
 R L G V L Q P E V Y K Q S L P G S N C K
                |1091
                                         |1101
|3311
       |3321
               |3331 |3341 |3351
                                        |3361
AGCATCCTGAAATAAAAAAGCAAGAATATGAAGAAGTAGTTCAGACTGTTAATACAGATT
 H P E I K K Q E Y E E V V Q T V N T D F
                11111
                                         11121
|3371 |3381
               13391
                       13401
                               13411
                                        13421
{\tt TCTCTCCATATCTGATTTCAGATAACTTAGAACAGCCTATGGGAAGTAGTCATGCATCTC}
 S P Y L I S D N L E Q P M G S S H A S Q
                |1131
13431
      |3441
              |3451
                       |3461
                               13471
                                        13481
AGGTTTGTTCTGAGACACCTGATGACCTGTTAGATGATGGTGAAATAAAGGAAGATACTA
V C S E T P D D L L D D G E I K E D T S
                |1151
                                         11161
                       |3521 |3531
13491
       |3501
               |3511
                                        |3541
GTTTTGCTGAAAATGACATTAAGGAAAGTTCTGCTGTTTTTTAGCAAAAGCGTCCAGAAAG
 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
               13631
13611
       13621
                       13641
                                13651
                                        13661
GAGGGGCCAAGAAATTAGAGTCCTCAGAAGAGAACTTATCTAGTGAGGATGAAGAGCTTC
 G A K K L E S S E E N L S S E D E E L P
                11211
                                         11221
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3671 3681 3691 3701 3711 CCTGCTTCCAACACTTGTTATTTGGTAAAGTAAACAATATACCTTCTCA	
C F Q H L L F G K V N N I P S Q 1231	S T R H
3731 3741 3751 3761 3771 ATAGCACCGTTGCTACCGAGTGTCTGTCTAAGAACACAGAGGAGAATTT	
S T V A T E C L S K N T E E N L	L S L K
3791 3801 3811 3821 3831 AGAATAGCTTAAATGACTGCAGTAACCAGGTAATATTGGCAAAGGCATC	
N S L N D C S N Q V I L A K A S	
3851 3861 3871 3881 3891	
ACCTTAGTGAGGAAACAAAATGTTCTGCTAGCTTGTTTTCTTCACAGTG L S E E T K C S A S L F S S O C	
LSEETKCSASLFSSQC 1291	S E L E 1301
3911 3921 3931 3941 3951	3961
AAGACTTGACTGCAAATACAAACACCCAGGATCCTTTCTTGATTGGTTC	CTTCCAAACAAA
DLTANTNTQDPFLIGS 1311	S K Q M 1321
3971 3981 3991 4001 4011	4021
TGAGGCATCAGTCTGAAAGCCAGGGAGTTGGTCTGAGTGACAAGGAATT	
R H Q S E S Q G V G L S D K E L 1331	V S D D 1341
4031 4041 4051 4061 4071	4081
ATGAAGAAGAGGGAACGGGCTTGGAAGAAATAATCAAGAAGAGCAAAG	GCATGGATTCAA
EERGTGLEENNQEEQS 1351	M D S N 1361
4091	gaagtgagctaa
	laaggaactgct
tettaaaettgaaaeatgtteeteetaaggtgetttteatagaaaaaag	

		•			•			•			•
gctag	gacgt	catct	ttgac	tgaat	gagct	ttaac	atcct	aatta	.ctggt	ggact	tactt
											ttttg
tataa	.C										



Ex	on	12	5	tar	t:	135	409	-	End	l: 1	355	80	I	eng	th:	17	1		
ta	ttt	ttt										ccaa						aca	ataa
ca	tca	agt										ccc							aata
at	ttc	atg	ggc	att								ntta							cttg
· ta	gtt	cca										• naaa							tttc
at	tta	atg	· gaa	agc								cttg							gaag
CA	GAG	41 GGA																	4241 AGAA
Q	R	D	Τ	M		H 401	N	L	Ι	K	L	Q	Q	Ε		A 411	Ε	L	E
GC	TGT		51 'AGA																4301 TGAC
		L			Н							S			S		I		D
TC	TTC											43 AAG					AGgt	tgt	gtat
S		А			D							S			Ε	K 451	_	_	
tg	ttg	gcc	aaa	cac	tga	tat	· ctt	aag	• caa	aat	tct	ttc	ctt	· ccc	ctt	tat	· ctc	stt	ctga
ag	agt	• aag	gac	· :cta	gct	cca	• aca	ttt	· tat	gat	cct	tgc	tca	• .gca	cat	ggg.	· taat	ta	• tgga
												cag							gtca
												itgc							gata
												jcta							

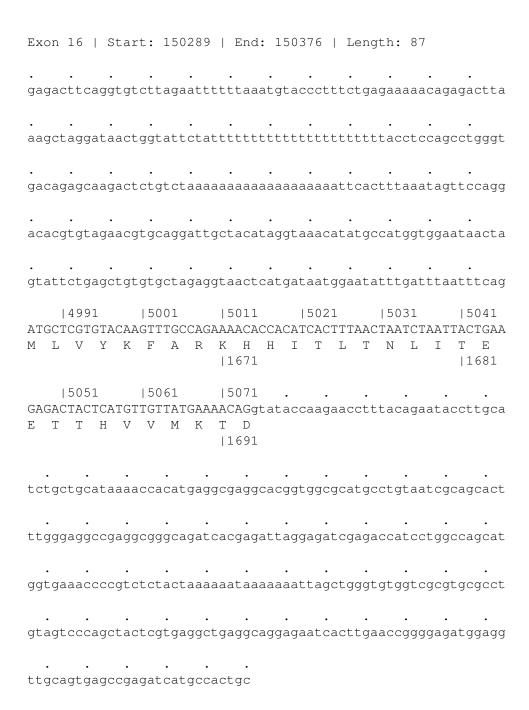


Exor	n 1	4	St	art	: 1	434	63	E	End:	: 14	365	3	L€	engt	h:	190			
agc																		ctg	rctt
														ccat			tgt	caa	ıccc
tgad																	tct	ttt	ttg
ccag																	gcc	cag	rcaa
gtat	.ga	ıttt	gtc	ctt [.]	tca					gatç				cttc	cat	tta	tct	ttc	tag:
		44	91		4	501			451	.1		45	21		4	531		1	4541
GTC	ATC																		TGG
S	S	Р	S	K		P 501		L	D	D	R	W	Y	М		S 511		S	G
		45	51		4	561			457	71		45	81		4	591		ı	4601
GAG	ГСТ	TCA	.GAA	TAG	AAA	СТА	.CCC	CATO	CTCF	AAGA	GGA	GCI	CAI	TAA	GGT	TGT	TGA	TGT	GGA
S	L	Q	N	R		Y 521		S	Q	Ε	Ε	L	Ι	K		V 531		V	Е
		46	11		4	621			463	31		46	341		4	651		ı	4661
GGA	GCA	ACA	.GCT	GGA	AGA	.GTC	TGG	GCC	CACA	ACGA	TTT	'GAC	GGI	AAAC	ATC	TTA	СТТ	GCC	CAAG
E	Q	Q	L	Ε		S 541		Ρ	Н	D	L	Τ	Ε	Τ		Y 551		Р	R
		46	71																
GCAZ	AGA			taa	tat	ttc	atc								ttg	att	tta	ctc	tga
Q	D	L	Ε																
· atco	•													ac+ a					
acco	- L C	icat	aaa	yatı	acc	cty	gtt	aac	LCac	1000	.сса	.yaı	.y.c	ıcta	.gcc	Lat	cat	yya	icac
•																			
tttt	gt	tat	act	taa	tta	agc	сса	ctt	tag	gaaa	aat	ago	ctca	aagt	gtt	aat	caa	ggt	tta
ctt																			

ctcacgcctgt

Exon	. 15	Sta	rt:	14	6/4	6	Εn	d:	14/	056	ı	Leng	gth	: 3.	10			
tcta	aaatta	atac	tat	tcc	tat	gac	taa	acc	ttt	gcat	tat	atct	ttt	tat	ctc	cct	agg	at
atat	ttctaa	aaac	tag	cat						taaa			gtt	aag	gtg	ttt	gct	ac
ataa	tgccat	catt	tcc	ttt						actt				cac	caa	cac	tgt	at
tcat	gtacco	catt	ttt	ctc	tta	acc	taa	ctt	tat	tggt	ct	tttt	caa	ttc	tta	aca	gag	ac
caga	acttt	gtaa	ttc	aac	att	cat	cgt	tgt	gta	aatt	caa	actt	cct	CCC	att	cct	ttc	ag
AGGG G	4683 AACCCC T P 1563	CTTA Y					AAT			CTT(F		TGAT	ΓGA	472: CCC' P	ГGA			ТС
CTTC S	4743 TGAAGA E D 1583	ACAG R	AGC	475 CCC P	AGA		AGC	TCG		TGG(CAA		ACC	ATC'	ГТС	AAC	CTC	ΤG
CATT L	4803 GAAAG3 K V 1603	TTCC P	CCA		GAA		TGC	AGA	ATC S	TGC(CCA	S	rcc.	AGC'	TGC'	TGC	TCA	ТА
CTAC T	4863 TGATA(D T 1623	CTGC A				TGC A		GGA	.AGA E	AAG: S					GAA			ΑT
TGAC T	4921 AGCTT(A S	CAAC T		493 .AAG R		CAA N	49 .CAA K		AAT M	GTC(S	951 CAT M 651	GGT		496: GTC: S			49 GAC T	

4981
gtggaaaagctatactttgggtatgatataggactttcgaattggaattttcctttctat
Etgtaaaagca



Exon	17	1	Sta	rt:	15	4033	1	End:	15	4110)	Len	gth	: 7	7		
· tataa	atg	gaç	gatc	tat	agct	tagc	cti	tggcç	gtct	agaa	agat	ggg	tgt	tgaç	gaaq	gago	ggagt
· ggaca	• agat	tat	ttc	ctc	tggt	tctt	aa	cttca	ntat	cago	ccto	•	tag	actt	· ccc	aaat	atcc
· atac	ctg	cto	ggtt	ata	atta	agtg	gt	gtttt	.cag	cct	ctga	attc	tgt	caco	cag	gggt	ttta
gaat	cata	aaa	atcc	aga	ttga	atct	tg	ggagt	:gta	• aaaa	aact	• :gag	gct	cttt	cago	cttc	cttag
· gaca	gca	ctt	cct	gat				aactt						tttt	ccat	ttct	gcag
	ΓGΑ	GT1	TGT	GTG		ACGG	AC		AAAT Y	ATTI	ГТСТ	AGG	AAT	TGC	GGG2	AGG <i>I</i>	5131 AAAAT K W 1711
	AGT:	ГАС		TTT				taata								aaca	· acctc
agaat	tg	cat	ttt	tac	acct	taac	gti	ttaac	cacci	taag	ggtt	ttt	gct	gato	gct	gagt	ctga
gtta	ccaa	• aaa	aggt	ctt				tacta						aata	atca	actt	tgtt
caga	caaq	gct	.ggt	gat	gct	ggga	aa	atggg	· jtct	cttt	tat	aac	taa	tago	gac	ctaa	itctg
ctcc	cago	caa	atgt	tag	cata	atga	gci	taggg	gatti	tatt	taa	atag	tcg	gcaq	gga	atco	catgt
gcag	cago	gca	aaac	tta	ta												

Exon	18		Sta	rt:	15	4611		End	d:	15	465	1	L	eng	th:	40			
ataa	ctaa	ata				ctgc												ttta	ıttt
aata	gtcc	dac	• agga	aato	ccat	tgtg	ca	· gca	ggc	caa	act	tat	·	tgt	tta	aat	taa	acat	caa
ctct	gtct	cc	• agaa	agga	aaa	ctgc	tg	cta	caa	igc	ctt	att	·	agg	gct	gtg	Igct	ttag	jagg
• gaag	gaco	ctc	· :tcc1	tct	gtca	attc	tt	cct	gtg	jct(tto		aat		tga	icct	ctct	atc
tccg	tgaa	aaa	• .gag	cac	gtto	cttc	tg	ctg	tat	gta	aac	ctç	gtc	ttt	tct	atg	ratc	tctt	tag
GGGT(CCA Q		TAT	ГАА	5171 AGAA E	AG	AAA	AAT		ГGА	AT(gta	agt	act	• .tga [.]	tgtt	aca
aact	• aacc	cag	· agat	tatt	cat	ttca	gt	cata	at <i>a</i>	igti	taa	aaa	• atg	tat	ttg	ctt	.cct	tcca	ıtca
atgc						aatg													att
atgc	• aggc	cct				ctca													egct
tgga	tcac	cct	gat	gtc	ggg			aga							gga	gaa	·	ccgt	ttc
tacta	aaaa	aat	· acaa	aaat	taq		gg:	ctt	ggt	gg	cac	ttg	gcc						



Exon	20	Sta	art:	166	867	E	End:	166	921		Leng	th:	54			
cttg	gcctç	gatto	ggtga	acaa	aagt	• .gaq	gatg	ctca	ıgtcc	tt	gaat	· gac	aaa	gaat	gcctg	t
• agag	tgcag	Igtca	aacta						ıgatc						ıtgttc	t
· ggac	attgg	Jacto	gctt						Icaga		tcat	·	gtg	gtga	· lacaga	a
· gaaa	aagaa	· laago	ctctt	cct	tttt	• gaa	aagt	ctgt	.tttt	tg	aata	· .aaa	gcc	aata	ıttctt	t
tata	actaç	gatti	ctcct	tct	ctcc	att	ccc	ctgt	.ccct	ct	ctct	· tcc	tct	cttc	ttcca	g
ATCT	5281 TCAGG R 1761	GGGG	CTAGA	AAAT	CTGI	TGC	CTAT	GGGC G F	5311 CCCTT F 1771	CA T	CCAA	532 CAT M	GCC	CACA	5331 Ggtaa D	
agcc	tggga	· Igaa	·	agag	ttcc	ago	·	agco	:tttg	tc	ttac	ata	gtg	gagt	attat	• а
agca	agato	· cca	cgato	gggg	gtto	cto	caga	ttgc	tgaa	at	gttc	· tag	agg	ctat	tctat	t
tctc	tacca	ictct	ccaa	aaca	.aaac	• ago	cacc	taaa	ıtgtt	at	ccta	tgg	caa	aaaa	laaact	а
tacc	ttgtc		cttct	caa	gago	atq	gaag	gtgg	ŗttaa	ta	gtta	• .gga ⁻	ttc	agta	ıtgtta	t
gtgt	tcaga	itgg	cgtt	gagc	tgct	gtt	tagt	gcca	lacat	• gt	tagt	gag	aaa	atat	.C	

Exon 21	. 5	Start	:: 1	16879	0	Enc	d: 1	688	863		Len	gth	: 7	3			
attaatg															gaga	acta	ıt
caaacct																	.g
gcaaatt	gact	ctaaa	aato	ccata	ccc	ctac	ctat	ttt	• :aag	gac	cat	tgt	cct	ttg	gag	caga	ıg
agacaga																	ıg
agggcct													aat	gtc	cati	ttta	ıg
ATCAACI Q L	'GGAZ E		ATGO		GCT.	GTGI	GGT	'GCI	TCI	GT V		GAA K	GGA		TTC	ATCA	5391 AT F
TCACCCT T L	TGG(agtat													et
· cctgtga				cacct													
tcttctc	· :tgc	ccaca	atao	cctgt	gcc	aaaa	igac	· tcc	cato	ctg	ŗtaa	ggg.	atg	ggt	• aag	gatt	.t
• gagaact				aatat													ca
tatgtcc	· ctc	cccct	cct	· cctct											aat	cato	ca
agaaatg	· atg	ggct															

Exon 22 Start: 170281 End: 170341 Length: 60
tgacagttccagtagtcctactttgacactttgaatgctctttccttcc
5411 5421 5431 5441 5451 5461 GGTGTCCACCCAATTGTGGTTGTGCAGCCAGATGCCTGGACAGAGGACAATGGCTTCCAT G V H P I V V V Q P D A W T E D N G F H 1811 1821
· g

Exon 23 Start: 172182 End: 173689 Length: 1507	
	cagga
tagagcctagtccaggagaatgaattgacactaatctctgcttgtgttctctgtct	
~ ~	
5591 *11 *21 *31 *41 ACTGACTGCAGCCACAGGTACAGAGCCACAGGACCCCAAGAATGAGCTTACA	
*61 *71 *81 *91 *101 GGCCTTTCCAGGCCCTGGGAGCTCCTCTCACTCTTCAGTCCTTCTACTGTCCTGGC	
$ \star121 \star131 \star141 \star151 \star161 $ AAATATTTTATGTACATCAGCCTGAAAAGGACTTCTGGCTATGCAAGGGTCCCTTA	
*181 *191 *201 *211 *221 TTTTCTGCTTGAAGTCTCCCTTGGAAATCTGCCATGAGCACAAAATTATGGTAATT	
*241 *251 *261 *271 *281 ACCTGAGAAGATTTTAAAACCATTTAAACGCCACCAATTGAGCAAGATGCTGATTC	

```
| *341
TTTATCAGCCCTATTCTTTCTATTCAGGCTGTTGTTGGCTTAGGGCTGGAAGCACAGAGT
   | *401
                                        | * 411
GGCTTGGCCTCAAGAGAATAGCTGGTTTCCCTAAGTTTACTTCTCTAAAACCCTGTGTTC
   | *421
         | * 431 | * 441 | * 451
                                | *461
                                        | * 471
ACAAAGGCAGAGTCAGACCCTTCAATGGAAGGAGTGCTTGGGATCGATTATGTGAC
   | *481 | *491 | *501 | *511 | *521
                                        1 * 531
{\tt TTAAAGTCAGAATAGTCCTTGGGCAGTTCTCAAATGTTGGAGTGGAACATTGGGGAGGAA}
   ATTCTGAGGCAGGTATTAGAAATGAAAAGGAAACTTGAAACCTGGGCATGGTGGCTCACG
   | * 601
          | * 611
                 | * 621
                         | * 631
                                 | *641
\verb|CCTGTAATCCCAGCACTTTGGGAGGCCAAGGTGGGCAGATCACTGGAGGTCAGGAGTTCG|\\
         | * 671 | * 681
   | * 661
                         | * 691
                                | *701
                                        | *711
AAACCAGCCTGGCCAACATGGTGAAACCCCATCTCTACTAAAAATACAGAAATTAGCCGG
   l *721
         | <del>*</del> 731 | <del>*</del> 741 | <del>*</del> 751
                                | *761
                                        | *771
TCATGGTGGTGGACACCTGTAATCCCAGCTACTCAGGTGGCTAAGGCAGGAGAATCACTT
   l * 781
         {\tt CAGCCCGGGAGGTGGAGGTTGCAGTGAGCCAAGATCATACCACGGCACTCCAGCCTGGGT}
   | * 901
          | * 911
                 | * 921
                         | * 931
                                | * 941
TTCTAAAAGTCTGAGATATATTTGCTAGATTTCTAAAGAATGTGTTCTAAAACAGCAGAA
                         | * 991
   | * 961
         |*971 |*981
                                | *1001
                                        | *1011
GATTTTCAAGAACCGGTTTCCAAAGACAGTCTTCTAATTCCTCATTAGTAATAAGTAAAA
   | *1021 | *1031 | *1041 | *1051 | *1061
                                        | *1071
TGTTTATTGTTGTAGCTCTGGTATATAATCCATTCCTCTTAAAATATAAGACCTCTGGCA
          | * 1091 | * 1101
                         | * 1111 | * 1121
| *1141 | *1151 | *1161 | *1171 | *1181 | *1191
```

*120) 1	* 1211	7	*1221		*1231		*1241		*1251
GCTTGCTGAA	AGGAAGA	AAAAAGT	rgttt:	TCATA	AACC	CATTA	TCCA6	GACTG	TTTAT	TAGCT
*120	61 ACTAGGI									
*132	21 ACGTTT									
*138	31 . catgaat	igactgt	ctcttq	gagact	tagg	· ccagc	cgact	·	• agago	ccttt
tcactgtgct										
gatttattga										
gcaaggtcag										
· aataaaggto	•	•								•
· · · actaaqat	5 2 2 3 9 0 0	5 5 5 6 5 6	- 5 - 0 0 0			5 - 5 - 5	55500		5 5 6 6	9 9
uctaayat										

LRG 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