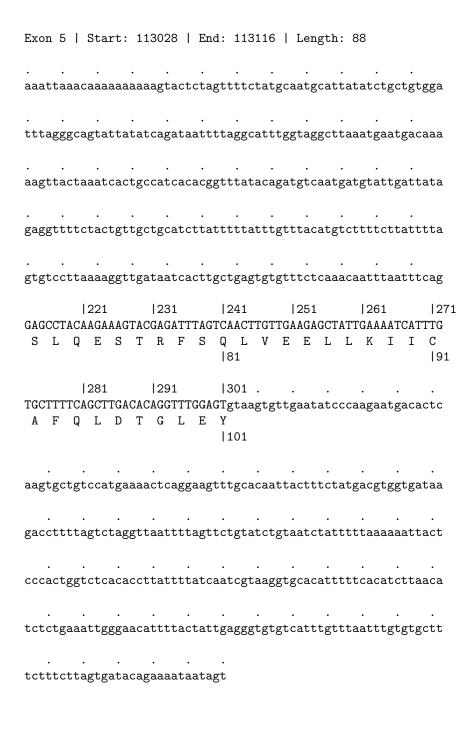
Gene: BRCA1 - Sequence: $NG_005905.2$ Transcript: $NM_007294.3$ - Protein: $NP_009225.1$ Date: February 25, 2015

 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 $\verb|tacttatatttaccgaaactggagacctccattagggcggaaagagtgggggattgggac| \\$ $\verb|ctcttcttacgactgctttggacaataggtagcgattctgaccttcgtacagcaattact|\\$ $\tt gtgatgcaataagccgcaactggaagagtagaggctagaggcaaggcactttatggcaaa$ $\verb|ctcaggtagaattcttcctcttccgtctcttttccttttacgtcatccgggggcagactgg|$. |-219 |-209 |-199 |-229 |-189 GTACCTTGATTTCGTATTCTGAGAGGCTGCTGCTTAGCGGTAGCCCCTTGGTTTCCGTGG |-169 |-129 |-119 CAACGGAAAAGCGCGGGAATTACAGATAAATTAAAACTGCGACTGCGCGGCGTGAGCTCG l-109 l-99 I-89 l-79 1-69 $\tt CTGAGACTTCCTGGACGGGGGACAGGCTGTGGGGTTTCTCAGATAACTGGGCCCCTGCGC$ |-39 |-29 ${\tt TCAGGAGGCCTTCACCCTCTGCTCTGGGTAAAGgtagtagagtcccgggaaagggacagg}$ $\tt gggcccaagtgatgctctggggtactggcgtgggagagtggatttccgaagctgacagat$ $\tt gggtattctttgacggggggtagggcggaacctgagaggcgtaaggcgttgtgaaccct$

ggggaggggggagtttgtaggtcgcgagggaagcgctgaggatcaggaagggggcactg
agtgtccgtgggggaatcctcgtgataggaactggaatatgccttgagggggacactatg
tctttaaaaacgtcggctggtcatgaggtcagg
Exon 2 Start: 93869 End: 93967 Length: 98
tacaaaaagcaacttctagaatctttaaaaaataaaggacgttgtcattagttctttggtt
-19
51 61 71 CATTAATGCTATGCAGAAAATCTTAGAGTGTCCCATCTGgtaagtcagcacaagagtgta I N A M Q K I L E C P I C 21

ttctgcactggggcaggaatctagtttagattaactggc
Exon 3 Start: 102205 End: 102258 Length: 53
aattcgtacgaactattatcaactaatcttttaaatgctgatgatagtatagagtattga
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
cccaggctggagtgcagtggcatgatcttggctcactgcaacctccgcccccg

Exoi	ı 4	:	St	art	: 1	111	.451	1	Eı	nd:	11:	152	8	L	eng	gth	: 7	7			
tatg	gaa	.ta	tat	cat	taa	aat	atg	gcc	ata	atta	aact	ttt [.]	tat	ta	agt	tt	tat	gtg	gato	ata	aca
gtaa	agc	cat	tat	gca	tgt	taa	ıgti	tca	gti	ttto	cata	aga	tca	tt	gct	ta	tgt	agt	tta	ıggt	ttt
tgc1	ta	.tg	cag	cat	cca	aaa	aaa	caa	tta	agga	aaa	cta	ttg	ct	tgt	caa	ttc	aco	tgc	:cat	tac
ttt!	ta	.aa†	tgg	ctc	tta	aag	ggg	cag	ttg	gtga	agat	tta	tct	tt	tca	atg	gct	att	tgo	ctt	ttg
agta	att	cti	ttc	taca	aaa	aag	gaa	agt	aaa	atta	aaat	ttg	ttc	tt	tct	tt	ctt	tat	aat	tta	atag
ATT:	TTG C				AA(I	15 CTT 51	CT(L	CAA N	CC				GGC		TC		GTG	P		ATO	
GAAT N	ΓGA D	20 TAT			AA <i>I</i> F	21 AGg 71	gtat	tat	aat	tttg	ggta	aat;	gat	gc	tag	ggt	tgg	aag	gcaa	ıcca	acag
tagg	gaa	.aaa	agt	agaa	aat	tta	ttt	taa	taa	acat	tago	cgt	tcc	ta	taa	aaa	cca	tto	cato	aga	aaaa
atti	tat	aaa	aag	agt	ttt	tta	ıgca	aca	cag	gtaa	aati	tat	ttc	ca	aag	gtt	att	tto	cctg	gaaa	agtt
ttai	gg	ga	cat	ctg	cct	tta	ita	cag	gta	atta	agaa	aac	tta	ct	gco	ctt	tct	cta	aatg	gctt	ccta
gtgt	taa	.aaa	act [.]	tgc	aga	act	ta†	tgt	aaa	agta	aggį	gct	gta	tc	gco	cgt	gcc	ccc	catt	gto	ctgt
taat	cct	tg	ttt [.]	tta	tat	tt															



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

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taca	ttg	ttg	gtg	tct	tag	ctt	tag	gtga	aatac	agta	ttg	ata	ggc	aaa	ttt	ctt	agt	gt
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taag	gta	gaa	aaca	aag	gac	tct	aaa	itaa	ctttg	atgg	gtct	grg	tat	ττε	gttt	ттв	τττ	CC
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		- 1	311			132	1		331			341			35	1		361
ATGC	AAA	CAG	CTA	ΓΑΑΊ	TTT	TGC	AA <i>A</i>	AAA	GGAAA	ATAA	CTC	TCC	TGA	ACA	TCT	AAA	AGA'	ΤG
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		ı	371			38	1		391			401			41	1		421
AAGT	TTC	TAT	CAT	CCA	AAG	TAT	GGC	CTA	CAGAA	ACCC	TGC	CAA	AAG	ACT	TCT	ACA	GAG'	ΓG
V	S	Ι	Ι	Q	S	М	G	Υ	R N	R	Α	K	R	Τ.	L	Ŋ	S	E
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agtc	ctc	ctg	cct	tag	ссс	cct	tag	gtag	ctggg	atta	ıcag	gca	cgc	gcc	acc	atg	cca	gg
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ctaa	ttt	ttg	tati	ttt	tag	tag	aga	ıcgg		cato	atg	ttg	gcc	agg	gctg	gtc	tcg	aa
ctaa	ttt	ttg	tati	ttt	tag	tag	aga	ıcgg	ggttt	cato	atg	ttg	gcc	agg	gctg	gtc	tcg	aa
ctaa	ttt [.] taa	ttg cct	tat: cagį	ttt [.] gtg:	tag atc	tag cac	aga	ıcgg	ggttt	cato	atg	ttg	gcc	agg	gctg	gtc	tcg	aa

Exon 7 Start: 118104 End: 118209 Length: 105
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
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$
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tggtcataatttgtatttgtggtaggcagctttgggaagtgaattttatgagccctatgg
tgagttataaaaaatgtaaaagacgcagttcccaccttgaagaatcttactttaaaaagg
agtgggtggatcacctgaggtcgggagttcgagaccagcctagcca

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

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taga	aacg	ggg	ggto	ctc	actt	ttgt	tg	gcc	agg	ctg	ggto	ttg	aac	tcct	aac	ctca	aataat
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caca	gtag	gat	gcto	cag	taaa	atat	tt	cta	gtt	gaa	atat	ctg	ttt	ttca	aaca	agta	catttt
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GATC	I'GA'	TTC	TTCI	GA						lGG(JAAC	TTA	TTG	CAGe	gtga	gtca	aagaga
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${\tt gcccagcaaccatttcatttcaactagaagtttctaaaggagagaga$
taaataagattggtcagctttctgtaatcgaaagagctaaaatgtttgatcttggtcatt
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tgacagttctgcatacatgtaactagtgtttcttattaggactctgtcttttccctatag
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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
${\tt GGATTCTGCAAAAAAGGgtaatggcaaagtttgccaacttaacaggcactgaaaagagag}$
D S A K K A
221
${\tt tgggtagatacagtactgtaattagattattctgaagaccatttgggacctttacaaccc}$
${\tt acaaaatctcttggcagagttagagtatcattctctgtcaaatgtcgtggtatggtctga}$
${\tt tagatttaaatggtactagactaatgtacctataataagaccttctgtaactgattgttg}$
$\verb ccctttcgtttttttttttgtttgtttgtttttttttt$
ttgcccaggctggagtg
Exon 10 Start: 123124 End: 126549 Length: 3425
EXON 10 Start. 123124 ENG. 120049 Length. 3425
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671   681   691   701   CTGCTTGTGAATTTTCTGAGACGGATGTAACAAATA	
ACEFSETDVTNT  231	ГЕНН Q Р S N N  241
731  741  751  761	
ATGATTTGAACACCACTGAGAAGCGTGCAGCTGAGA	
D L N T T E K R A A E R  251	R H P E K Y Q G S  261
791  801  811  821	831  841
GTTCTGTTTCAAACTTGCATGTGGAGCCATGTGGCA	
S V S N L H V E P C G T  271	F N T H A S S L Q  281
851  861  871  881 AGCATGAGAACAGCAGTTTATTACTCACTAAAGACA	891  901 AGAATGAATGTAGAAAAGGCTGAAT
HENSSLLLTKD F  291	R M N V E K A E F
291	301  951  961
291   911   921   931   941	301  951  961 AGCCAACATAACAGATGGGCTGGAA
291   911	301  951  961 AGCCAACATAACAGATGGGCTGGAA S Q H N R W A G S  321
291    911   921   931   941  TCTGTAATAAAAGCAAACAGCCTGGCTTAGCAAGGA C N K S K Q P G L A R S   311    971   981   991   1001  GTAAGGAAACATGTAATGATAGGCGGACTCCCAGCA	301  951  961 AGCCAACATAACAGATGGGCTGGAA S Q H N R W A G S  321  1011  1021 ACAGAAAAAAAGGTAGATCTGAATG
291   911	301  951  961 AGCCAACATAACAGATGGGCTGGAA S Q H N R W A G S  321  1011  1021 ACAGAAAAAAAGGTAGATCTGAATG
291    911	301    951   961 AGCCAACATAACAGATGGCTGGAA S Q H N R W A G S   321    1011   1021 ACAGAAAAAAAGGTAGATCTGAATG F E K K V D L N A   341
291    911	301    951   961 AGCCAACATAACAGATGGGCTGGAA S Q H N R W A G S   321    1011   1021 ACAGAAAAAAAAGGTAGATCTGAATG F E K K V D L N A   341    1071   1081 CAGAAAACTGCCATGCTCAGAGAATC
1291	301    951   961 AGCCAACATAACAGATGGGCTGGAA S Q H N R W A G S   321    1011   1021 ACAGAAAAAAAGGTAGATCTGAATG F E K K V D L N A   341    1071   1081 CAGAAACTGCCATGCTCAGAGAATC Q K L P C S E N P   361
1291	301    951   961 AGCCAACATAACAGATGGGCTGGAA S Q H N R W A G S   321    1011   1021 ACAGAAAAAAAGGTAGATCTGAATG T E K K V D L N A   341    1071   1081 CAGAAACTGCCATGCTCAGAGAATC Q K L P C S E N P   361

AGTGGTTTTCCAGAAGTGATGAACTGTTAGGTTCTGATGACTCACATGATGGGGAGTCTG W F S R S D E L L G S D D S H D G E S E 391 1211 1221 1231 1241 1251 11261 AATCAAATGCCAAAGTAGCTGATGTATTGGACGTTCTAAATGAGGTAGATGAATATTCTG S N A K V A D V L D V L N E V D E Y S G 1411 1421 11271 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 |1361 1341 1351 |1371 |1381 GTGAAAGAGTTCACTCCAAATCAGTAGAGAGTAATATTGAAGACAAAATATTTGGGAAAA ERVHSKSVESNIEDKIFGKT 451 1461 1391 11401 11411 1421 1431 11441 CCTATCGGAAGAAGCCAACCTCCCCAACTTAAGCCATGTAACTGAAAATCTAATTATAG Y R K K A S L P N L S H V T E N L I I G 471 481 1451 1471 1481 1461 1491 |1501 A F V T E P Q I I Q E R P L T N K L K R 1491 11511 1521 |1531 1541 l 1551 11561 GTAAAAGGAGACCTACATCAGGCCTTCATCCTGAGGATTTTATCAAGAAAGCAGATTTGG K R R P T S G L H P E D F I K K A D L A |511 |521 11571 l 1581 11591 |1601 |1611 11621  ${\tt CAGTTCAAAAGACTCCTGAAATGATAAATCAGGGAACTAACCAAACGGAGCAGAATGGTC}$ V Q K T P E M I N Q G T N Q T E Q N G Q |531 |541

11181

|1191

1201

11171

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11151

11631

1641

1651

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|1661

AAGTGATGAATATTACTAATAGTGGTCATGAGAATAAAACAAAAGGTGATTCTATTCAGA
V M N I T N S G H E N K T K G D S I Q N

1671

11681

|561

| 1691 | 1701 | 1711 | 1721 | 1731 | 1741 | ATGAGAAAAATCCTAACCCAATAGAATCACTCGAAAAAGAATCTGCTTTCAAAACGAAAG E K N P N P I E S L E K E S A F K T K A | 571 | 581

| 1811 | 1821 | 1831 | 1841 | 1851 | 1861 | AAGCACCTAAAAAGAATAGGCTGAGGAGGAAGTCTTCTACCAGGCATATTCATGCGCTTG | A P K K N R L R R K S S T R H I H A L E | 611 | 621

| 1871 | 1881 | 1891 | 1901 | 1911 | 1921 |
AACTAGTAGTCAGTAGAAATCTAAGCCCACCTAATTGTACTGAAATTGATAGTT
L V V S R N L S P P N C T E L Q I D S C | 641

|2231 |2241 |2251 |2261 |2271 |2281

CTGAAGACCCCAAAGATCTCATGTTAAGTGGAGAAAGGGTTTTGCAAACTGAAAGATCTG E D P K D L M L S G E R V L Q T E R S V  $\mid$  751  $\mid$  761

| 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
- | 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 | 961
- | 2951 | 2961 | 2971 | 2981 | 2991 | 3001 TTCCCATCAAGTCATTTGTTAAAACTAAATGTAAGAAAAATCTGCTAGAGGAAAACTTTG P I K S F V K T K C K K N L L E E N F E | 991 | 1001
- |3011 |3021 |3031 |3041 |3051 |3061 AGGAACATTCAATGTCACCTGAAAGAGAAATGGGAAATGAGAACATTCCAAGTACAGTGA E H S M S P E R E M G N E N I P S T V S |1011 | 1021
- |3071 |3081 |3091 |3101 |3111 |3121 GCACAATTAGCCGTAATAACATTAGAGAAAATGTTTTTAAAGAAGCCAGCTCAAGCAATA T I S R N N I R E N V F K E A S S S N I |1031 |1041
- |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

|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

| 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

|3791 |3801 |3811 |3821 |3831 |3841 AGAATAGCTTAAATGACTGCAGTAACCAGGTAATATTGGCAAAGGCATCTCAGGAACATC N S L N D C S N Q V I L A K A S Q E H H |1271 |1281

391 AAGA					T A C A												961	\ C \ I	۱ ۸
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403 ATGA	AGA	AAG	AGG	AAC	GGGC	CTT	GGA.	AGA/	AAA	ΓΑΑΊ	ΓCA <i>I</i>	AGA.	AGA	GCA	AAG	CATO			
Е	E	R	G	Т	G		E 351	E	N	N	Q	Е	Е	Q	S		D 361	S	N
409 ACTT L		tat	tgg:	aac				gtgt						tta	taga	aagt	tgag	gcta	aa
atgt	tta	tgc	ttti	tgg	ggag	gca	cat [.]	ttta	acaa	aatt	ttc	caa	gta	tag	ttaa	aagg	gaad	ctgo	ct
tctt	aaa		gaa:		tgtt										aagt		ttca	acad	ca
gcta	.gga	cgt	cat	ctt	tgad	etga	aat	gago	ctt1	taad	cato	cct	aat	tac	tggt	Egga	actt	cact	t
ctgg	ttt	cat	ttt:	ata	aaag	gca	aat	ccag	ggtg	gtco	ccaa	aag	caa	gga	attt	caat	tcat	cttt	g
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4161  4171  4181
ctaaaccttcgaaacccatttgctaatcc
Exon 12   Start: 135409   End: 135580   Length: 171

tagttccatactaggtgatttcaattcctgtgctaaaattaatt
${\tt atttaatggaaagcttctcaaagtatttcattttcttggtgccatttatcgtttttgaag}$
4191
1401   1411
4251
1421   1431
4311  4321  4331  4341  4351 . TCTTCTGCCCTTGAGGACCTGCGAAATCCAGAACAAGCACATCAGAAAAAGgtgtgtat
S S A L E D L R N P E Q S T S E K A
1441   1451
catggtatacaacatagctcttgctctattggaagctaagtggaatgggaga
Exon 13   Start: 141370   End: 141496   Length: 126
acticiptic transport to the transport to

tggtgctagcatctgttgttgcattgcttgtgtttataaaattctgcctgatatacttg
4361
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

•	•		•		•	•		•		•	•		•		•	•		•	
tta	aac	ago	tgg	gag	ata	tgg	tgc	ctc	aga	.cca	acc	caa	ccc	cat	gtt	ata	tgt	caa	ссс
tga	cat	att	ggc	agg	caa	· catį	gaa	tcc	aga	· .ctt	cta	ıggo	:tgt	ctt	· gcg	ggc	tct	ttt	ttg
	ort o	++e		ma+		tct	ma c	ata	. a m c	+ a+	.++	. 2++		· c+	++a	·	ac c		caa
cca	guc	aut	, , , ,	gat	CUC	000	gac	aug	agu	ugi	, , , , ,	aut	σαι	gcu	uug	gcu	gcc	cag	caa
•					•			•		•					•				
gta	tga	ittt	gtc	CTT	tca	caa ⁻	ttg	gtg	gcg	ate	ggtt	ttc	tcc	ttc	cat	tta	tct	ttc	tag
		44			•	501			451			45			•	531			4541
GTC.	ATC S		TTC S	TAA K		CCC. P		ATT L		.TG <i>A</i> D		GTG W		CAT M		.CAG S	TTG C	CTC S	TGG G
ъ	۵	Р	۵	ĸ		Р 501	۵	ь	ע	ע	ĸ	W	ĭ	M		.511	C	۵	G
G A G'	тст	45  ידכא		ፐለር		561 CTA			457 TCA			45  GCT		'ፐለለ		:591 'TCT'			4601 GGA
S	L	Q	N	R		Y		S	Q	Е	E	L	I	K		V	D	V	E
					1	521									1	531			
		146	311		14	621		ı	463	1		46	341		14	651		1	4661
GGA	GCA	ACA	GCT	GGA	AGA	GTC'					TTT			AAC					AAG
E	Q	Q	L	E		S 541	G	P	Н	D	L	T	Ε	T		Y .551	L	P	R
					11:	041									ΙI	.551			
		146											•						
GCA. Q	AGA D		'AGg E	taa	tat	ttc	atc	tgc	tgt	att	gga	aca	laac	act	ttg	att	tta	ctc	tga
٩	ב	_	_																
atc	cta	cat	· aaa	.gat	att	ctg	gtt	aac	caa	· .ctt	tta	ıgat	gta	cta	gtc	tat	cat	gga	cac
				J		Ů.	_					J	J		0			50	
· +++	+ cr+	+2+	·	+22	++ >		cc2	c++	+20		tee	. a		a mt	«++		C 2 2	mat	tta
000	ug u	, , ,	acu	vaa	······	ag c	cca		عمان	aac	iaa	age	, , , ,	agu	800	aat	caa	560	oua
					•												4		
CTT	gaa	ıaat	tat	rga	aac	rgt.	ιaa	UCC	atc	tat	att	tta	iatt	aat	ggt	rta	act	aat	gat
ttt	gag	gat	gag	gga	gtc	ttg	gtg	tac	tct	aaa	tgt	att	att	tca	ggc	cag	gca	tag	tgg

Exon	15	5	Star	rt:	14	6746	3	En	d:	147	056		Len	gth	: 3	10			
tcta	aaati	ιατ	cact	tat	rcc.	τατε	gac	taa	acc	cttt	gca	tat	atc	τττ	tat	CTC	CCT	agg	аt
atat	ttcta	aaa	aact	cag	cat	tgtt	zga	.ctg	aaa	agtg	taa	ata	cgt	gtt	aag	gtg	ttt	gct	ac
ataa		.+-							+			++ m			an a		60.6	· + «+	<b>^</b> +
auaa	uguu	160	1661		666	uua	gga	aac	uac	igci	act	uug	gat	uuc	cac	Caa	.cac	ugu	аь
tcat	gtac	cca	attt	ttt	ctc	ttaa	acc	taa	ctt	tat	tgg	tct	ttt	taa	ttc	tta	.aca	gag	ac
	•																	•	
caga	actti	tgt	caat	tca	aac	atto	cat	cgt	tgt	tgta	aat	taa	act	tct	ссс	att	cct	ttc	ag
AGGG	468  AACC		ΓΤΑΟ								'41 CTT(								
G	T I		Y	L	E	S	G	Ι	S	L		S 571		D	P	E	S	D	P
CTTC	474  TGAA										14 [.] TGG:								
S	E I	)									G		Ι					S	
CATTO	480										48								
L	K \  160	J									Α		S						
CTAC	486  TGAT		rgci								48								
Т		Γ				N					S	V 631	S	R		K			
	492			•	493							951							
TGAC	AGCT	ΓC <i>I</i>	AAC/	AGA	AAG	GGT	CAA	CAA	AAC	FAAT	'GTC	CAT	GGT	GGT	GTC	TGG	CCT	GAC	CC
Т	A 5	3 41	T	Ε	R	V	N	K	R	M		M 651		V	S	G	L	T	P

ctgtaaaagca
Exon 16   Start: 150289   End: 150376   Length: 87
4991  5001  5011  5021  5031  5041
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
5051  5061  5071

## 1691 . . $\verb|tctgctgcataaaaccacatgaggcgaggcacggtggcgcatgcctgtaatcgcagcact|$ $\verb|ttgggaggccgaggcgggcagatcacgagattaggagatcgagaccatcctggccagcat|$ $\tt ggtgaaaccccgtctctactaaaaaaataaaaaaattagctgggtgtggtcgcgtgcgcct\\$ $\tt gtagtcccagctactcgtgaggctgaggcaggagaatcacttgaaccggggagatggagg$ ttgcagtgagccgagatcatgccactgc Exon 17 | Start: 154033 | End: 154110 | Length: 77 ${\tt tataatggagatctatagctagccttggcgtctagaagatgggtgttgagaagagggagt}$ $\tt ggacagatatttcctctggtcttaacttcatatcagcctcccctagacttccaaatatcc$ $\verb|atacctgctggttata| attagtggtgttttcagcctctgattctgtcaccaggggtttta|$ ga at cata a at ccagatt gat ctt ggg ag tg ta aa aa act gag gct ctt tag ctt ctt ag tag ag tg ta $\tt gacag cacttcct gatttt gtttt caacttct aat ccttt gag t gttttt cattct g cag$ |5111 |5081 |5091 |5101 |5121 ATGCTGAGTTTGTGTGAACGGACACTGAAATATTTTCTAGGAATTGCGGGAGGAAAAT A E F V C E R T L K Y F L G I A G G K W 1701 |1711 |5141 |5151 . . . . . . . . . . $\tt GGGTAGTTAGCTATTTCTgtaagtataatactatttctcccctcctcctttaacacctc$

ETTHVVMKTD

V V S Y F W

cagcaggcaaacttata
Exon 18   Start: 154611   End: 154651   Length: 40
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$

atgcaggcctgcactgtggctcatacctataatcccagcgctttgggaggctgaggcgct
tactaaaaatacaaaattagccgggcttggtggcacttgcc
Exon 19   Start: 160849   End: 160932   Length: 83
gcattcaaaagattctcctgcctcagcctcccaagtagctgggattacaggtgcctgcc
5261  5271

ggctgtgtaatttggattcccgtctcgggttcagatcttagctgataagtggaagagctg
ggactttaagcagatgagaatcta
Exon 20   Start: 166867   End: 166921   Length: 54
ggacattggactgcttgtccctgggaagtagcagcagaaatcatcaggtggtgaacagaa
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
gtgttcagatggcgttgagctgctgttagtgccaacatgttagtgagaaaatatc

Exon	21	Sta	art:	1687	90	End:	168	863	L	engtl	n: 7	3			
attaa	atgg	gaaatt	cagat	ccttt	gatt	ttttt	ttct	ttca	agc	attti	tatt [.]	tgag	gaga	.ctat	
caaad	cctt	tataco	caagt	eggcc	ttat	ggaga	ictga	taac	cag	agta	catg	gcat	ato	agtg	
gcaaa	attg	gactta	aaaat	ccat	accc	ctact	attt	taag	gacc	attg	tcct	ttgg	gago	agag	
agaca	agad	ctctcc	ccatt	gaga	ggtc	ttgct	ataa	gcct	tca	tccgg	gaga	gtgt	agg	gtag	
agggo	ccte	gggtta	aagta	atgca	gatt	actgo	agtg	attt	tac	atcta	aaat	gtcc	att	ttag	
	ACT( L	5341 GGAATO E W  1781	GGAT( M	GTAC						GTGA V K	AGGA	5381 GCTT L	TCA		
TCACO T	CCTI L	5401  GGCAC  G T  1801	CAgta	aagta	ttgg	gtgco	ctgt	caga	igag	ggagg	gaca	caat	att	ctct	
cctgt	tgag	gcaaga	actgg	gcacc	tgtc	agtco	ctat	ggat	gcc	ccta	ctgt:	agco	ctca	gaag	
tctto	ctct	Egccca	acata	acctg	tgcc	aaaag	gactc	cato	tgt:	aaggg	gatg	ggta	iagg	attt	
gagaa	actg	gcacat	catta	aaata	tact	gaggg	gaaga	.cttt	ttc	cctc1	taac	tctt	ttt	ccca	
tatgt	tcc	ctcccc	cctco	ctctc	tgtg	actgo	ccca	.gcat	act	gtgt1	ttca	acaa	natc	atca	
agaaa	atga	atgggo	ct												

```
Exon 22 | Start: 170281 | End: 170341 | Length: 60
\verb|ccaggagtttgagaccagcctgggcaacatggcaaaaccctgtctctaccaaaaatacaa|\\
aaaaattagccaggggtggtggtacgtgtctgtagttccagctacttaggaggctgagat\\
ggaaggattgcttgagcccaggaggcagaggtggcagttgagcttgagatcacaccactgca\\
              .
|5421 |5431
                        |5441
                                |5451
GGTGTCCACCCAATTGTGGTTGTGCAGCCAGATGCCTGGACAGAGGACAATGGCTTCCAT
G V H P I V V V Q P D A W T E D N G F H
                  1811
{\tt Ggtaaggtgcctgcatgtacctgtgctatatggggtccttttgcatgggtttggtttatc}
      {\tt actcattacctggtgcttgagtagcacagttcttggcacattttaaatatttgttgaatg}
\verb| aatggctaaa | atgtctttttgatgtttttattgttatttgttttatattgtaaa | agtaat | \\
{\tt caaggccgggcatggtggctcatgcctataattccagcactttgggaggtcaagatggag}
g
```

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

taaaaatacaaa	aattagctggg	tgtgatggca	tgtgcctgta	aattccagct	actcagga
ggcagagacagg	agaattgcttg	aacccaggag	gcggaggttg	gaatgagccg	agattgcg
ccatcacactct	agcctcggcga	.cagagcaaga	 ctccgtctca	 aaaaaaaaaa	aaaaaaaa
ttagcttctacc	 tcattaatcct	 aagaactcat	 acaaccagga	accctggagt	cgattgat
tagagcctagtc	caggagaatga	 attgacacta	 atctctgct	 tgtgttctct	gtctccag
5471 CAATTGGGCAGA I G Q M	TGTGTGAGGCA		CCCGAGAGT		
5531 CACTCTACCAGT L Y Q C	GCCAGGAGCTG Q E L	5551 GACACCTACC D T Y L  1851	TGATACCCC		
5591 ACTGACTGCAGC	*11 CAGCCACAGGT				
*61 GGCCTTTCCAGG		*81 CCTCTCACTC			
*121 AAATATTTTATG		*141 GAAAAGGACT			
*181 TTTTCTGCTTGA		*201 GAAATCTGCC			*231 AATTTTTC
*241 ACCTGAGAAGAT		*261 TTAAACGCCA		-	
*301 TTTATCAGCCCT	• -	*321 TCAGGCTGTT		-	*351 CACAGAGT
*361	<b>*</b> 371	<b>*</b> 381	*391	*401	<b> </b> *411

GGCTTGGCCTCAA	GAGAATAGCT	'GGTTTCCCTA	AGTTTACTTC	TCTAAAACC	CTGTGTTC
*421 ACAAAGGCAGAGA			*451 GGAGAGTGCTT		
*481 TTAAAGTCAGAAT			*511 ATGTTGGAGT		
*541 ATTCTGAGGCAGG					
*601 CCTGTAATCCCAG			*631 GGCAGATCAC		
*661 AAACCAGCCTGGC			*691 CTCTACTAAAA		
*721 TCATGGTGGTGGA			*751 CCAGGTGGCTA		
*781 CAGCCCGGGAGGT			*811 GATCATACCAC		
*841 GACAGTGAGACTG	•	-	*871 AAAAAGGAAA	-	-
*901 TTCTAAAAGTCTG			*931 CTAAAGAATGT		
*961 GATTTTCAAGAAC			*991 CCTAATTCCTC		
*1021 TGTTTATTGTTGT					
*1081 TGAATATTTCATA			*1111 CACCAGGAAGG		
*1141 GAGGTGATTTTT			*1171 GAAACCATACA		
*1201 GCTTGCTGAAGGA			*1231 ACCCATTATO		

|*1291

|*1301

|*1311

**|***1281

 $\tt GTTGGAAGGACTAGGTCTTCCCTAGCCCCCCCAGTGTGCAAGGGCAGTGAAGACTTGATT$ 

**|***1271

|*1261

GTAC	*1321 AAAATAC										
CACT	*1381  CCAccat		gactg				gccago				
	tgtgctt				gtaaaa						
	tattgagį									acaaa	ttgtt
gcaaį	ggtcagaa	agtct	gaggt		caactg						
aataa	aaggtgti	tcgca		tgtto		taga		gggt	ccttgo	cagtt	ctagg
actaa	agat										

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