

## Product datasheet for **RC400713**

### BRCA2 (NM\_000059) Human Mutant ORF Clone

#### Product data:

Product Type:	Mutant ORF Clones
Product Name:	BRCA2 (NM_000059) Human Mutant ORF Clone
Mutation Description:	L2776X
Affected Codon#:	2776
Affected NT#:	8327
Nucleotide Mutation:	BRCA2 Mutant (L2776X), Myc-DDK-tagged ORF clone of Homo sapiens breast Cancer, early onset (BRCA2) as transfection-ready DNA
Effect:	Breast cancer
Symbol:	BRCA2
Synonyms:	BRCC2; BROVCA2; FACD; FAD; FAD1; FANCD; FANCD1; GLM3; PNCA2; XRCC11
E. coli Selection:	Kanamycin (25 ug/mL)
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
Tag:	Myc-DDK
ACCN:	NM_000059
ORF Size:	8325 bp
Restriction Sites:	SgfI-RsrII
ORF Nucleotide Sequence:	>RC400713 representing NM_000059 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**GCGATCGCC**

ATGCCTATTGGATCCAAAGAGAGGCCAACATTTTTGAAATTTTTAAGACACGCTGCAACAAAGCAGATT  
TAGGACCAATAAGTCTTAATTGGTTTGAAGAACTTTCTTCAGAAGCTCCACCCTATAATTCTGAACCTGC  
AGAAGAATCTGAACATAAAAAACAACAATTACGAACCAACCTATTTAAAACCTCCACAAAGGAAACCATCT  
TATAATCAGCTGGCTTCAACTCCAATAATTTCAAAGAGCAAGGGCTGACTCTGCCGCTGTACCAATCTC  
CTGTAAGAAGATTAGATAAATTCAAATTAGACTTAGGAAGGAATGTTCCCAATAGTAGACATAAAGTCT  
TCGCACAGTGAAGAACTAAAATGGATCAAGCAGATGATGTTTCTGTCCACTTCTAAATCTTGTCTTAGT  
GAAAGTCTGTTGTTCTACAATGTACACATGTAACACCACAAAGAGATAAGTCAGTGGTATGTGGGAGTT



[View online »](#)

TGTTTCATACACCAAAGTTTGTGAAGGGTCGTCAGACACCAAAACATATTTCTGAAAGTCTAGGAGCTGA  
GGTGGATCCTGATATGTCTTGGTCAAGTTCTTTAGCTACACCACCACCCTTAGTTCTACTGTGCTCATA  
GTCAGAAATGAAGAAGCATCTGAAACTGTATTTCTCATGATACTACTGCTAATGTGAAAAGCTATTTTT  
CCAATCATGATGAAAGTCTGAAGAAAAATGATAGATTTATCGTCTCTGTGACAGACAGTGA AACACAAA  
TCAAAGAGAAGCTGCAAGTCATGGATTTGGAAAAACATCAGGGAATTCATTTAAAGTAAATAGCTGCAAA  
GACCACATTGGAAAGTCAATGCCAAATGTCTAGAAGATGAAGTATATGAAACAGTTGTAGATACCTCTG  
AAGAAGATAGTTTTTTCATTATGTTTTCTAAATGTAGAACAAAAATCTCAAAAAAGTGAAGAACTAGCAA  
GACTAGGAAAAAATTTCCATGAAGCAACGCTGATGAATGTGAAAAATCTAAAAACCAAGTGA AAGAA  
AAATACTCATTTGTATCTGAAGTGAACCAATGATACTGATCCATTAGATTCAATGTAGCAAATCAGA  
AGCCCTTTGAGAGTGAAGTGACAAAATCTCCAAGGAAGTTGTACCGTCTTTGGCCTGTGAATGGTCTCA  
ACTAACCTTTAGGCTCAAATGGAGCCAGATGGAGAAAAATACCCTATTGCATATTTCTTCATGTGAC  
CAAAATATTTAGAAAAAGACCTATTAGACACAGAGAACAAAAGAAAGAAAGATTTTCTACTTCAGAGA  
ATTCTTTGCCACGTATTTCTAGCCTACCAAAATCAGAGAAGCCATTAAATGAGGAAACAGTGGTAAATAA  
GAGAGATGAAGAGCAGCATCTTGAATCTCATACAGACTGCATTCTTGCAAGTAAAGCAGGCAATATCTGGA  
ACTTCTCCAGTGGCTTTCATTTAGGGTATCAAAAAGTCTATATTCAGAATAAGAGAATCACCTAAAG  
AGACTTCAATGCAAGTTTTTCCAGGTATATGACTGATCCAAACTTTAAAAAAGAAACTGAAGCCTCTGA  
AAGTGGACTGGAATACATACTGTTTGCTCACAGAAGGAGGACTCCTTATGTCCAAATTTAATTGATAAT  
GGAAGCTGGCCAGCCACCACACAGAATTCTGTAGCTTTGAAGAATGCAGGTTAATATCCACTTTGA  
AAAAGAAAAAATAAGTTTATTTATGCTATACATGATGAAACATCTTATAAAGGAAAAAATACCGAA  
AGACCAAAAATCAGAATAATTAAGTGTTCAGCCAGTTTGAAGCAATGCTTTTGAAGCACCATTACA  
TTTGCAATGCTGATTCAGGTTTATTGCATTCTCTGTGAAAAGAAGCTGTTACAGAATGATTCGAAG  
AACCAATGCTCCTTAACTAGCTCTTTGGGACAATCTGAGGAAATGTTCTAGAAATGAAACATGTTT  
TAATAATACAGTAATCTCTCAGGATCTTGATTATAAAGAAGCAAAATGTAAATAAGGAAAAACTACAGTTA  
TTTATTACCCAGAAGCTGATTCTGTGATGCCTGCAGGAAGGACAGTGTGAAAAATGCTCAAAAAGCA  
AAAAAGTTTCCAGATATAAAGAAGAGGTCTGGCTGCAGCATGTACCCAGTACAACATTCAAAAGTGG  
ATACAGTGATACTGACTTTCAATCCCAGAAAAGTCTTTTATATGATCATGAAAATGCCAGCACTCTTATT  
TTAACTCTACTTCCAAGGATGTTCTGTCAAACCTAGTCATGATTTCTAGAGGCAAGAATCATACAAAA  
TGTCAGACAAGCTCAAAGGTAACAATTATGAATCTGATGTTGAATTAACCAAAAATATCCCATGGAAAA  
GAATCAAGATGTATGTGCTTTAAATGAAAATTA AAAACGTTGAGCTGTTGCCACCTGAAAAATACATG  
AGAGTAGCATCACCTTCAAGAAAGGTACAATTCACCAAAAACACAATCTAAGAGTAATCAAAAAAATC  
AAGAAGAACTACTTCAATTTCAAAAATAACTGTCAATCCAGACTCTGAAGAATTTTCTCAGACATGA  
GAATAATTTGTCTTCCAAGTAGCTAATGAAAGGAATAATCTTGCTTTAGGAAATACTAAGGAATTCAT  
GAAACAGACTTGACTTGTGTAACGAACCCATTTTCAAGAACTCTACCATGGTTTTATATGGAGACACAG  
GTGATAAACAAGCAACCAAGTGTCAATTA AAAAAGATTTGGTTTATGTTCTTGCAAGGAGAACAAAA  
TAGTGTAAAGCAGCATATAAAAATGACTCTAGGTCAAGATTTAAAATCGGACATCTCCTTGAATATAGAT  
AAAATACCAGAAAAAATAATGATTACATGAACAAATGGGCAGGACTCTTAGGTCCAATTTCAAATCACA  
GTTTTGGAGGTAGCTTCAAGACAGCTTCAAATAAGGAAATCAAGCTCTCTGAACATAACATTAAGAAGAG  
CAAAATGTTCTTCAAAGATATTGAAGAACAATATCCTACTAGTTTAGCTTGTGTTGAAATGTAATACC  
TTGGCATTAGATAATCAAAAAGAACTGAGCAAGCCTCAGTCAATTAATACTGTATCTGCACATTTACAGA  
GTAGTGTAGTTGTTTCTGATTGTAAAAATAGTCATATAACCCCTCAGATGTTATTTTCCAAGCAGGATTT  
TAATTCAAACCATAATTTAACACCTAGCCAAAAGGCAGAAATTACAGAATTTTCTACTATATTAGAAGAA  
TCAGGAAGTCAGTTTGAATTTACTCAGTTTAGAAAACCAAGCTACATATTGCAGAAGAGTACATTTGAAG  
TGCCTGAAAACAGATGACTATCTTAAAGACCCTTCTGAGGAATGCAGAGATGCTGATCTTCATGTCAT  
AATGAATGCCCATCGATTGGTCAGGTAGACAGCAGCAAGCAATTTGAAGGTACAGTTGAAATTAACGG  
AAGTTTGTGCTGGCCTGTTGAAAAATGACTGTAACAAAAGTCTTCTGGTTATTTAACAGATGAAAAAG  
TGGGGTTTAGGGCTTTTATTCTGCTCATGGCACAAAACCTGAATGTTTCTACTGAAGCTCTGCAAAAAGC  
TGTGAAACTGTTTAGTGATATTGAGAATATTAGTGAGGAACTCTGCAGAGGTACATCCAATAAGTTTA  
TCTTCAAGTAAATGTCATGATTCTGTTGTTCAATGTTTAAAGATGAAAAATCATAATGATAAACTGTAA  
GTGAAAAAATAATAATGCCAACTGATATTACAAAATAATTTGAAATGACTACTGGCACTTTTGTGGA  
AGAAATTAAGTAAAAATCAAGAGAAATGAAAAATGAAGATAACAAATATACTGCTGCCAGTAGAAAT  
TCTCATAACTTAGAATTTGATGGCAGTGATTCAAGTAAAAATGATACTGTTTGTATTATAAAGATGAAA  
CGGACTTGCTATTTACTGATCAGCACAACATATGTCTTAAATTAATCTGGCCAGTTTATGAAGGAGGGAAA

CACTCAGATTAAGAAGATTTGTCAAGTTAACTTTTTGGAAGTTGCGAAAGCTCAAGAAGCATGTCAT  
 GGTAATACTTCAATAAAGAACAGTTAACTGCTACTAAAACGGAGCAAAATATAAAGATTTTGAGACTT  
 CTGATACATTTTTTTCAGACTGCAAGTGGGAAAAATATTAGTGTGCGCCAAAGAGTCATTTAATAAAATTTG  
 AAATTTCTTTGATCAGAAACCAGAAGAAATGCATAACTTTTCCTAAATTTCTGAATTACATTCTGACATA  
 AGAAAGAACAAAATGGACATTCTAAGTTATGAGGAAACAGACATAGTTAAACACAAAATCTGAAAGAAA  
 GTGTCCAGTTGGTACTGGAATCAACTAGTGACCTCCAGGGACAACCCGAACGTGATGAAAAGATCAA  
 AGAACCCTACTCTATTGGGTTTTTCATACAGCTAGCGGGAAAAAAGTTAAATTTGCAAAGGAATCTTTGGAC  
 AAAGTGAAAAACCTTTTTGATGAAAAAGCAAGGTAAGTAAATCACCAGTTTTAGCCATCAATGGG  
 CAAAGACCCTAAAGTACAGAGAGGCTGTAAAGACCTGAATTAGCATGTGAGACCATTGAGATCACAGC  
 TGCCCCAAAGTGTAAAGAAATGCAGAATCTCTCAATAATGATAAAAACTTGTTTCTATTGAGACTGTG  
 GTGCCACCTAAGCTCTAAGTGATAATTTATGTAGACAACTGAAATCTCAAAACATCAAAAAGTATCT  
 TTTTGAAGTTAAAGTACATGAAAATGTAGAAAAAGAACAGCAAAAAGTCTGCAACTTGTACACAAA  
 TCAGTCCCCTTATTCAGTCATTGAAAATTCAGCCTTAGCTTTTTACACAAGTTGTAGTAGAAAACTTCT  
 GTGAGTCAGACTTCATTACTGGAAGCAAAAAATGGCTTAGAGAAGGAATATTTGATGGTCAACCAGAAA  
 GAATAAATACTGCAGATTATGTAGGAAATTTTGTATGAAAATTAATCAAACAGTACTATAGCTGAAA  
 TGACAAAAATCATCTCTCCGAAAAACAAGATACTTATTTAAGTAACAGTAGCATGTCTAACAGCTATTCC  
 TACCATTCTGATGAGGTATATAATGATTCAGGATATCTCTCAAAAAATAAACTTGATTCTGGTATTGAGC  
 CAGTATTGAAGAATGTTGAAGATCAAAAAACACTAGTTTTTCCAAAGTAATATCCAATGTAAGAGATGC  
 AAATGCATACCCACAACTGTAATGAAGATAATTTGCGTTGAGGAACTTGTGACTAGCTCTTACCCTGC  
 AAAAAATAAAATGCAGCCATTAATTTGCCATATCTAATAGTAATAATTTTGGAGTAGGGCCACCTGCAT  
 TTAGGATAGCCAGTGGTAAATCGTTTTGTGTTTCATGAAACAATTAAGGAAAGTAAAGACATATTTAC  
 AGACAGTTTCAGTAAAGTAATTAAGGAAAAACAGGAGAATAAATCAAAAAATTTGCCAAACGAAAATATG  
 GCAGTTGTTACGAGGCATTGGATGATTCAGAGGATTTCTTCATAACTCTAGATAAGTAAAGTGA  
 GCACGCATTCACATAAGGTTTTTGTGACATTCAGAGTGAAGAATTTTACAACATAACCAAAATATGTC  
 TGGATTGGAGAAAGTTTCTAAAAATACCTTGTGATGTTAGTTTGGAACTTCAGATATATGAAATGT  
 AGTATAGGGAAGCTTCATAAGTCAGTCTCATCTGCAAATACTTGTGGGATTTTTAGCACAGCAAGTGAA  
 AATCTGTCCAGGTATCAGATGCTTATTACAAAACGCAAGACAAGTGTCTTGAATAGAAAGATAGTAC  
 CAAGCAAGTCTTTTCCAAAGTATTGTTTAAAAGTAACGAACATTCAGACCAGCTCACAAGAGAAGAAAAT  
 ACTGCTATACGTACTCCAGAACATTTAATATCCAAAAAGGCTTTTCATATAATGTGGTAAATTCATCTG  
 CTTTCTCTGGATTTAGTACAGCAAGTGGAAAGCAAGTTCCATTTTGAAGGTTCTTACACAAAGTTAA  
 GGGAGTGTAGAGGAATTTGATTTAATCAGAAGTGGAGTACTTCACTATTCACCTACGTCTAGACAA  
 AATGTATCAAAAAATCTCCTCGTGTGATAAGAGAAACCCAGAGCACTGTGAAACTCAGAAATGGAAA  
 AAACCTGCAGTAAAGAATTTAAATATCAAATAACTTAAATGTTGAAGGTGGTCTTCAGAAAATATCA  
 CTCTATTAAGTTTCTCCATATCTCTCAATTTCAACAAGACAAAACAACAGTTGGTATTAGGAACCAAA  
 GTGTCACTTGTGAGAACATTCATGTTTTGGGAAAAAGAACAGGCTTCACTAAAAACGTAATAATGGAAA  
 TTGGTAAAACGAACTTTTTCTGATGTTCTGTGAAAACAAATAGAAAGTTGTTCTACTTACTCCAA  
 AGATTCAGAAAACACTTTGAAACAGAAGCAGTAGAAATGCTAAAGCTTTTATGGAAGATGATGAAGT  
 ACAGATCTAAACTGCCAAGTCATGCCACATTTCTTTTTACATGTCCCGAAAATGAGGAAATGGTTT  
 TGTCAAATCAAGAATTGGAAAAAGAAGAGGAGAGCCCTTATCTTAGTGGGAGAACCCCAATCAAAAAG  
 AAACCTTAAATGAATTTGACAGGATAATAGAAAAATCAAGAAAAATCCTTAAAGGCTTCAAAAAGCACT  
 CCAGATGGCACATAAAAAGATCGAAGATTGTTTATGCATCATGTTTCTTTAGAGCCGATTACCTGTGTAC  
 CCTTTCGCACAACATAAGGAACGTCAAGAGATACAGAAATCCAAATTTTACCACCTGGTCAAGAATTTCT  
 GTCTAAATCTCATTTGTATGAACATCTGACTTTGGAAAAATCTTCAAGCAATTTAGCAGTTTCAGGACAT  
 CCATTTTATCAAGTTTCTGCTACAAGAAATGAAAAATGAGACACTTGATTACTACAGGCAGACCAACCA  
 AAGTCTTTGTTCCACCTTTTAAACTAAATCACATTTTACAGAGTTGAACAGTGTGTTAGGAATATTA  
 CTTGGAGGAAAACAGACAAAAGCAAAACATTGATGGACATGGCTCTGATGATAGTAAAAATAAGATTAAT  
 GACAATGAGATTCATCAGTTTAACAAAAACAACCTCAATCAAGCAGCAGCTGAACTTTACAAAAGTGTG  
 AAGAAGAACCTTTAGATTTAATTACAAGTCTTCAAGATGCCAGAGATATACAGGATATGCGAATTAAGAA  
 GAAACAAGGCAACGCGTCTTTCCACAGCCAGGCACTGTATCTTGAAAAACATCCACTCTGCCTCGA  
 ATCTCTCTGAAAGCAGCAGTAGGAGGCCAAGTTCCCTCTGCGTGTCTCATAAACAGCTGTATACGTATG  
 GCGTTTCTAAACATTGCATAAAAAATTAACAGCAAAAAATGCAGAGTCTTTTTCAGTTTACACTGAAGATTA  
 TTTTGGTAAGGAAAGTTTATGGACTGAAAAAGGAATACAGTTGGCTGATGGTGGATGGCTCATACCCTCC

AATGATGGAAAGGCTGGAAAAGAAGAATTTTATAGGGCTCTGTGTGACACTCCAGGTGTGGATCCAAAGC  
TTATTTCTAGAATTTGGGTTTATAATCACTATAGATGGATCATATGGAACTGGCAGCTATGGAATGTGC  
CTTTCCTAAGGAATTTGCTAATAGATGCCTAAGCCCAGAAAGGGTGCTTCTTCAACTAAAATACAGATAT  
GATACGGAAATTGATAGAAGCAGAAGATCGGCTATAAAAAAGATAATGGAAAGGGATGACACAGCTGCAA  
AAACACTTGTTCTCTGTGTTTCTGACATAATTTTCATTGAGCGCAAATATATCTGAACTTCTAGCAATAA  
AACTAGTAGTGCAGATACCCAAAAAGTGGCCATTATTGAACTTACAGATGGGTGGTATGCTGTAAAGGCC  
CAGTTAGATCCTCCCCTCTTAGCTGTCTTAAAGAATGGCAGACTGACAGTTGGTCAGAAGATTATTCTTC  
ATGGAGCAGAACTGGTGGGCTCTCCTGATGCCTGTACACCTCTTGAAGCCCAGAATCTTTATG

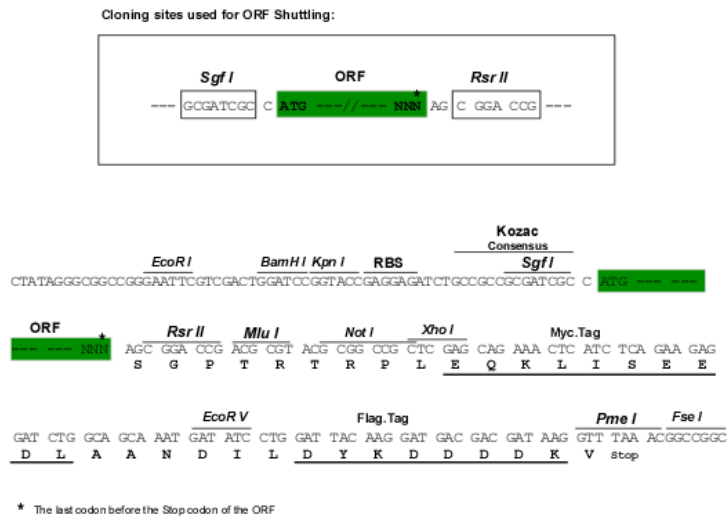
AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC  
TGGATTACAAGGATGACGACGA TAAGGTTTAA

**Protein Sequence:** >RC400713 representing NM\_000059  
 Red=Cloning site Green=Tags(s)

MPIGSKERPTFFEIFKTRCNKADLGPISLWNWFELSSSEAPPYNSEPAEESSEHKNNNYEPNLFKTPQRKPS  
 YNQLASTPIIFKEQGLTLPLYQSPVKELDKFKLDLGRNVNPSRHKSLRTVKTMDQADDVSCPLLNSCLS  
 ESPVVLQCTHTVTPQRDKSVVCGSLFHTPKFVKGRQTPKHISESLGAEVDPDMSWSSSLATPTLLSSTVLI  
 VRNEEASETVFPHDTTANVKSYFSNHDESLKKNDRFIAVTDSENTNQREAASHGFGKTSNGNSFKVNSCK  
 DHIGKSMPNVLEDEVYETVVDTSEEDSFLCFSKCRTKNLQKVRTSKTRKKIFHEANADECEKSKNQVKE  
 KYSFVSEVEPNDDPLDSNVANQKPFESGSDKI SKEVVVPSLACEWSQLTSLGNGAQMEKIPLLHSSCD  
 QNISEKDLLDTENKRKDFLTSENSLPRISLSEKPLNEETVVNKRDEEQHLESHTDCILAVKQAIISG  
 TSPVASSFQGIKKSIFRIRESPKETFNASFSGHMTDPNFKKETEASESGLEIHTVCSQKEDSLCPNLIDN  
 GSWPATTTQNSVALKNAGLISTLKKTNKFIYAIHDETSYKGGKIPKQKSELINCSAQFEANAFEAPLT  
 FANADSGLLHSSVKRSCSQNDSEPTLSLTSSFGTILRKCSRNETCSNNTVISQDLDYEAKCNKEKLQL  
 FITPEADSLSCLQEQCENDPKSKKVSIDIKEEVLAAACHPVQHSKVEYSDTDFQSQKSLLYDHENASTLI  
 LTPTSKDVL SNLVMISRGKESYKMSDKLKGNNYSDVELTKNIPMEKNQDVCALNENYKNVELLPPEKYM  
 RVASPSRKVQFNQNTNLRVIQKQNEETTSISKITVNPDSEELFSDNENNFVQVANERNLALGNTKELH  
 ETDLTCVNEPIFKNSTMVLVYGDGDKQATQVSIKDLVYVLAENKNSVKQHIKMTLGGDLKSDISLNI  
 KIPEKNNDYMNKWAAGLLGPISNHSFGGSFRASNKEIKLSEHNKIKSKMFFKDIIEEQYPTSLACVEIVNT  
 LALDNQKLSKPKQSINTVSAHLQSSVVVSDCKNSHITPQMLFSKQDFNSNHLTPSQAEITELSTILEE  
 SGSQFEFTQFRKPSYILQKSTFEVPEQMILKTTSEECRDADLHVIMNAPSIGQVDSKQFEGTVEIKR  
 KFAGLLKNDCKNSASGYLTDENEVGRGFYSAHGTLNVSTEALQKAVKLFSDIENISEETSAEVHPISL  
 SSKKCHDSVVSFMFKIENHNDKTVSEKNNKQQLILQNNIEMTTGTVEEITENYKRNTENEDNKYTAASRN  
 SHNLEFDGSDSSKNDTVCIHKDETDLLFTDQHNICLKLSGQFMKEGNTQIKEDLSDLTFLEVAQAQEACH  
 GNTSNKEQLTATKTEQNIKDFETSDTFFQTASGKNISVAKESFNKIVNFFDQKPEELHNFSLNSELHSDI  
 RKNKMDILSYEETDIVKHKILKESVPVGTGNQLVTFQGGPERDEKIKEPTLLGFHTASGKVKIAKESLD  
 KVKNLFDKEQGTSEITFSHQWAKTLKYREACKDLELACETIEITAAPKCKEMQNSLNNDKNLVSIVTV  
 VPPKLLSDNLCRQTNELKTSKIFLKVKHENVEKETAKSPATCYTNQSPYSVIENSALAFYTCSRKTS  
 VSQTSLLAEAKWVREGIFDQPERINTADYVGNLYENNSNSTIAENDKNHLSEKQDYLNSSSMSNSYS  
 YHSDEVYNDSGYL SKNKLDSGIEPVLKNVEDQKNTSFSKVISNVKDANAYPQTVNEDICVEELVTSSPC  
 KNKNAAIKLSISNSNNEVGGPPAFRIASGKIVCVSHETIKKVKDIFTDSFSKVIKENNENKSKICQTKIM  
 AGCYEALDSEDILHNSLDNDECSTHSHKVFADIQSEEILQHNQNMGLKVKISPCDVSLETSDICKC  
 SIGKLVHKSVAANTCGIFSTASGKSVQVSDASLQARQVFSEIEDSTKQVFSKVLFKSNEHSDQLTREEN  
 TAIRTPHELISQKGFYNVNVSSAFSGFSTASGKQVSILESSLHKVKGVLVEEFDLIRTEHSLHYSPTS  
 NVSKILPRVDRNPEHCVNSEMEKTCSEFKLSNNLNVEGGSSENNHSIKVSPYLSQFQDQKQLVLGK  
 VSLVENIHLVKGQASPKNVKMEIGKTEFTSDVPVKTNIEVCSTYSKDSSENYFETEAVEIAKAFMEDDEL  
 TDSKLP SHATHSLFTCPENEEMVLSNSRIGKRRGEPLILVGEPSIKRNLNDFDRIIENQEKSLKASKST  
 PDGTIKDRRLFMHVSLEPITCVPFRTTKERQEQNPNTAPGQEFLLSKSHLYEHLTLEKSSNLAVSGH  
 PFYQVSATRNEKMRHLITTRPTKVFVPPFKTKSHFHRVEQCVRNINLEENRQKQKQIDGHGSDSKNKIN  
 DNEIHQFNKNNNSNQAAAVFTFKCEEEPLDLITSLQNARDIQDMRIKKKQRQVFPQPGSLYLAKTSTLPR  
 ISLKAAVGGQVPSACSHKQLYTYGVSKHCIKINSKNAESFQFHTEDYFGKESLWTGKGIQLADGGWLLPS  
 NDGKAGKEEFYRALCDTPGVDPKLSRIWVYNHYRWIWKLAAMECAFPEKANRCLSPERVLLQLKYRY  
 DTEIDRSRRSAIKKIMERDDTAAKTLVLCVSDIISLSANISETSSNKTSSADTQKVAIIELTDGWYAVKA  
 QLDPPLLAVLKNLGRLLTVGQKIIILHGAELVGPDACTPLEAPESLM

SGPTRRRLEQKLISEEDLAANDILDYKDDDDKV

**Restriction Sites:** Sgfl-RsrII

**Cloning Scheme:**

**OTI Disclaimer:**

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

**OTI Annotation:**

This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**Components:**

The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

**RefSeq:**

[NP\\_000050](#)

**RefSeq Size:**

8325 bp

**RefSeq ORF:**

10257 bp

**Locus ID:**

675

**Cytogenetics:**

13q13.1

**Protein Families:**

Druggable Genome

**Protein Pathways:**

Homologous recombination, Pancreatic cancer, Pathways in cancer

**MW:**

305.3 kDa

**Gene Summary:**

Inherited mutations in BRCA1 and this gene, BRCA2, confer increased lifetime risk of developing breast or ovarian cancer. Both BRCA1 and BRCA2 are involved in maintenance of genome stability, specifically the homologous recombination pathway for double-strand DNA repair. The largest exon in both genes is exon 11, which harbors the most important and frequent mutations in breast cancer patients. The BRCA2 gene was found on chromosome 13q12.3 in human. The BRCA2 protein contains several copies of a 70 aa motif called the BRC motif, and these motifs mediate binding to the RAD51 recombinase which functions in DNA repair. BRCA2 is considered a tumor suppressor gene, as tumors with BRCA2 mutations generally exhibit loss of heterozygosity (LOH) of the wild-type allele. [provided by RefSeq, May 2020]