

Product datasheet for **RC400756**

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:	R3128X
Affected Codon#:	3128
Affected NT#:	9382
Nucleotide Mutation:	BRCA2 Mutant (R3128X), 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:	9381 bp
Restriction Sites:	Sgfl-RsrII
ORF Nucleotide Sequence:	>RC400756 representing NM_000059 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGCC**

ATGCCTATTGGATCCAAAGAGAGGCCAACATTTTTGAAATTTTTAAGACACGCTGCAACAAAGCAGATT
TAGGACCAATAAGTCTTAATTGGTTTGAAGAACTTTCTTCAGAAGCTCCACCCTATAATTCTGAACCTGC
AGAAGAATCTGAACATAAAAAACAACAATTACGAACCAACCTATTTAAAACCTCCACAAAGGAAACCATCT
TATAATCAGCTGGCTTCAACTCCAATAATTTCAAAGAGCAAGGGCTGACTCTGCCGCTGTACCAATCTC
CTGTAAGAAGATTAGATAAATCAAATTAGACTTAGGAAGGAATGTTCCCAATAGTAGACATAAAGTCT
TCGCACAGTAAAACTAAAATGGATCAAGCAGATGATGTTTCTGTCCACTTCTAAATCTTGTCTTAGT
GAAAGTCTGTTGTTCTACAATGTACACATGTAACACCACAAAGAGATAAGTCAGTGGTATGTGGGAGTT



[View online »](#)

TGTTCATACACCAAAGTTTGTGAAGGGTCGTCAGACACCAAAACATATTTCTGAAAGTCTAGGAGCTGA
 GGTGGATCCTGATATGTCTTGGTCAAGTTCTTTAGCTACACCACCACCCTTAGTTCTACTGTGCTCATA
 GTCAGAAATGAAGAAGCATCTGAAACTGTATTTCTCATGATACTACTGCTAATGTGAAAAGCTATTTTT
 CCAATCATGATGAAAGTCTGAAGAAAAATGATAGATTTATCGTCTCTGTGACAGACAGTGAACACAAA
 TCAAAGAGAAGCTGCAAGTCATGGATTTGGAAAAACATCAGGGAATTCATTTAAAGTAAATAGCTGCAAA
 GACCACATTGGAAAGTCAATGCCAAATGTCTAGAAGATGAAGTATATGAAACAGTTGTAGATACCTCTG
 AAGAAGATAGTTTTTTCATTATGTTTTCTAAATGTAGAACAAAAATCTACAAAAAGTGAAGCAAGCAA
 GACTAGGAAAAAATTTTCCATGAAGCAACGCTGATGAATGTGAAAAATCTAAAAACCAAGTGAAGAA
 AAATACTCATTTGTATCTGAAGTGAACCAATGATACTGATCCATTAGATTCAAATGTAGCAAATCAGA
 AGCCCTTTGAGAGTGAAGTGACAAAATCTCCAAGGAAGTTGTACCGTCTTTGGCCTGTGAATGGTCTCA
 ACTAACCTTTCAGGTCTAAATGGAGCCAGATGGAGAAAAATACCCTATTGCATATTTCTTCATGTGAC
 CAAAATATTTAGAAAAAGACCTATTAGACACAGAGAACAAAAGAAAGAAAGATTTTCTACTTCAGAGA
 ATTCTTTGCCACGTATTTCTAGCCTACCAAAATCAGAGAAGCCATTAAATGAGGAAACAGTGGTAAATAA
 GAGAGATGAAGAGCAGCATCTTGAATCTCATACTGACTGCATTCTTGAGTAAAGCAGGCAATATCTGGA
 ACTTCTCCAGTGGCTTTCATTTCCAGGTATCAAAAAGTCTATATTCAGAATAAGAGAATCACCTAAAG
 AGACTTTCAATGCAAGTTTTTCCAGGTATATGACTGATCCAAACTTTAAAAAAGAAACTGAAGCCTCTGA
 AAGTGGACTGGAATACATACTGTTTGCTCACAGAAGGAGGACTCCTTATGTCCAAATTTAATTGATAAT
 GGAAGCTGGCCAGCCACCACCACAGAAATCTGTAGCTTTGAAGAATGCAGGTTAATATCCACTTTGA
 AAAAGAAAAACAAATAAGTTTATTTATGCTATACATGATGAAACATCTTATAAAGGAAAAAATACCGAA
 AGACCAAAAATCAGAATAATTAAGTGTTCAGCCAGTTTGAAGCAATGCTTTTGAAGCACCATTACA
 TTTGCAAAATGCTGATTCAGGTTTATTGCATTCTCTGTGAAAAGAAGCTGTTACAGAAATGATTTGAA
 AACCAATACAGTAATCTCTCAGGATCTTGATTATAAAGAAGCAAAATGTAAATAAGGAAAAACTACAGTTA
 TTTATTACCCAGAAGCTGATTCTGTGATGCCTGCAGGAAGGACAGTGTGAAAAATGCTCAAAAAGCA
 AAAAAGTTTCAGATATAAAGAAGAGGTCTGGCTGCAGCATGTACCCAGTACAACATTCAAAAGTGGAA
 ATACAGTGATACTGACTTTCAATCCCAGAAAAGTCTTTTATATGATCATGAAAATGCCAGCACTCTTATT
 TTAACTCTACTTCCAAGGATGTTCTGTCAAACCTAGTCATGATTTCTAGAGGCAAGAATCATAAAAA
 TGTCAGACAAGCTCAAAGGTAACAATTATGAATCTGATGTTGAATTAACCAAAAATATCCCATGGAAAA
 GAATCAAGATGTATGTGCTTTAAATGAAAATTAAAAAACGTTGAGCTGTTGCCACCTGAAAAATACATG
 AGAGTAGCATCACCTTCAAGAAAGGTACAATTCACCAAAAACACAATCTAAGAGTAATCAAAAAAATC
 AAGAAGAACTACTTCAATTTCAAAAATAACTGTCAATCCAGACTCTGAAGAATTTTCTCAGACATGA
 GAATAATTTTGTCTTCCAAGTAGCTAATGAAAGGAATAATCTTGCTTTAGGAAATACTAAGGAATTCAT
 GAAACAGACTTGACTTGTGTAACCGAACCCATTTTCAAGAACTCTACCATGGTTTTATATGGAGACACAG
 GTGATAAACAAGCAACCAAGTGTCAATTA AAAAAGATTTGGTTTATGTTCTTGACAGAGGAGAACAAAA
 TAGTGTAAAGCAGCATATAAAAAAGACTCTAGGTCAAGATTTAAATCGGACATCTCCTTGAATATAGAT
 AAAATACCAGAAAAAATAATGATTACATGAACAAATGGGCAGGACTCTTAGGTCCAATTTCAAATCACA
 GTTTTGGAGGTAGCTTCAAGACAGCTTCAAATAAGGAAATCAAGCTCTCTGAACATAACATTAAGAAGAG
 CAAAATGTTCTTCAAAGATATTGAAGAACAATATCCTACTAGTTTAGCTTGTGTTGAAATGTAATACC
 TTGGCATTAGATAATCAAAAGAACTGAGCAAGCCTCAGTCAATTAATACTGTATCTGCACATTTACAGA
 GTAGTGTAGTTGTTTCTGATTGTAAAAATAGTCATATAACCCCTCAGATGTTATTTTCCAAGCAGGATTT
 TAATTCAAACCATAATTTAACACCTAGCCAAAAGGCAGAAATTACAGAATTTTCTACTATATTAGAAGAA
 TCAGGAAGTCAGTTTGAATTTACTCAGTTTAGAAAACCAAGCTACATATTGCAGAAGAGTACATTTGAAG
 TGCCTGAAAACAGATGACTATCTTAAAGACCCTTCTGAGGAATGCAGAGATGCTGATCTTCATGTCAT
 AATGAATGCCCATCGATTGGTCAGGTAGACAGCAGCAAGCAATTTGAAGGTACAGTTGAAATTAACGG
 AAGTTTGTGGCCTGTTGAAAAATGACTGTAACAAAAGTCTTCTGGTTATTTAACAGATGAAAAATGAAG
 TGGGGTTTAGGGCTTTTATTCTGCTCATGGCACAACCAACTGAATGTTTCTACTGAAGCTCTGCAAAAAGC
 TGTGAAACTGTTTAGTGATATTGAGAATATTAGTGAGGAACTCTGCAGAGGTACATCCAATAAGTTTA
 TCTTCAAGTAAATGTCATGATTCTGTTGTTCAATGTTTAAAGATGAAAAATCATAATGATAAACTGTAA
 GTGAAAAAATAATAATGCCAACTGATATTACAAAATAATTTGAAATGACTACTGGCACTTTTGTGGA
 AGAAATTAAGTAAAAATCAAGAGAAATACTGAAAATGAAGATAACAAATATACTGCTGCCAGTAGAAAT
 TCTCATAACTTAGAATTTGATGGCAGTGATTCAAGTAAAAATGATACTGTTTGTATTATAAAGATGAAA
 CGGACTTGCTATTTACTGATCAGCACAACATATGTCTTAAATTTCTGGCCAGTTTATGAAGGAGGGAAA

CACTCAGATTAAGAAGATTTGTGAGATTTAACTTTTTGGAAGTTGCGAAAGCTCAAGAAGCATGTCAT
 GGTAATACTTCAATAAAGAACAGTTAACTGCTACTAAAACGGAGCAAAATATAAAAGATTTTGAGACTT
 CTGATACATTTTTTTCAGACTGCAAGTGGGAAAAATATTAGTGTGCGCCAAAGAGTCATTTAATAAAATTTG
 AAATTTCTTTGATCAGAAACCAGAAGATTGCATAACTTTTCTTAAATTTCTGAATTACATTCTGACATA
 AGAAAGAACAAAATGGACATTCTAAGTTATGAGGAAACAGACATAGTTAAACACAAAATCTGAAAGAAA
 GTGTCCAGTTGGTACTGGAATCAACTAGTGACCTCCAGGGACAACCCGAACGTGATGAAAAGATCAA
 AGAACCTACTCTATTGGGTTTTTCATACAGCTAGCGGGAAAAAAGTTAAATTTGCAAAGGAATCTTTGGAC
 AAAGTGAAAAACCTTTTTGATGAAAAAGCAAGGTAAGTAAATCACCAGTTTTAGCCATCAATGGG
 CAAAGACCCTAAAGTACAGAGAGGCTGTAAAGACCTGAATTAGCATGTGAGACCATTGAGATCACAGC
 TGCCCCAAAGTGTAAAGAAATGCAGAATTCTCTCAATAATGATAAAAACTTGTTTCTATTGAGACTGTG
 GTGCCACCTAAGCTCTAAGTGATAATTTATGTAGACAACTGAAAATCTCAAAACATCAAAAAGTATCT
 TTTTGAAGTTAAAGTACATGAAAATGTAGAAAAAGAACAGCAAAAAGTCTGCAACTTGTACACAAA
 TCAGTCCCCTTATTCAGTCATTGAAAATTCAGCCTTAGCTTTTTACACAAGTTGTAGTAAAAAAGTCT
 GTGAGTCAGACTTCATTACTGGAAGCAAAAAATGGCTTAGAGAAGGAATATTTGATGGTCAACCAGAAA
 GAATAAATACTGCAGATTATGTAGGAAATTTTGTATGAAAATAATTCAAACAGTACTATAGCTGAAA
 TGACAAAAATCATCTCTCCGAAAAACAAGATACTTATTTAAGTAAACAGTGCATGTCTAACAGCTATTCC
 TACCATTCTGATGAGGTATATAATGATTCAGGATATCTCTCAAAAAATAAACTTGATTCTGGTATTGAGC
 CAGTATTGAAGAATGTTGAAGATCAAAAAACACTAGTTTTTCCAAAGTAAATCCAATGTAAGAGATGC
 AAATGCATACCCACAACTGTAATGAAGATAATTTGCGTTGAGGAACTTGTGACTAGCTCTTACCCTGC
 AAAAAATAAAATGCAGCCATTAATTTGCCATATCTAATAGTAATAATTTTGGAGTAGGGCCACCTGCAT
 TTAGGATAGCCAGTGGTAAATCGTTTTGTGTTTCATGAAACAATTAAGGAAAGTAAAGACATATTTAC
 AGACAGTTTCAGTAAAGTAATTAAGGAAAAACAGGAGAATAAATCAAAAAATTTGCCAAACGAAAATATG
 GCAGTTGTTACGAGGCATTGGATGATTCAGAGGATTTCTTCATAACTCTAGATAAGTAAAGTGA
 GCACGCATTCACATAAGGTTTTTGTGACATTCAGAGTGAAGAATTTTACAACATAACCAAAATATGTC
 TGGATTGGAGAAAGTTTCTAAAAATACCTTGTGATGTTAGTTTGGAACTTCAGATATATGAAATGT
 AGTATAGGGAAGCTTCATAAGTCAGTCTCATCTGCAAATACTTGTGGGATTTTTAGCACAGCAAGTGAA
 AATCTGTCCAGGTATCAGATGCTTATTACAAAACGCAAGACAAGTGTCTTGAATAGAAAGATAGTAC
 CAAGCAAGTCTTTTCAAAGTATTGTTTAAAAGTAACGAACATTCAGACCAGCTCACAAGAGAAGAAAAT
 ACTGCTATACGTACTCCAGAACATTTAATATCCAAAAAGGCTTTTATATAATGTGGTAAATTCATCTG
 CTTTCTCTGGATTTAGTACAGCAAGTGGAAAGCAAGTTCCATTTTGAAGGTTCTTACACAAAGTTAA
 GGGAGTGTAGAGGAATTTGATTTAATCAGAAGTGGGATAGTCTTCACTATTCACCTACGTCTAGACAA
 AATGTATCAAAAAATCTCCTCGTGTGATAAGAGAAACCCAGAGCACTGTGAAACTCAGAAATGGAAA
 AAACCTGCAGTAAAGAATTTAAATTATCAAATAACTTAAATGTTGAAGGTGGTTCTTCAGAAAATATCA
 CTCTATTAAGGTTTCTCCATATCTCTCAATTTCAACAAGACAAAACAACAGTTGGTATTAGGAACCAAA
 GTGTCACTTGTGAGAACATTCATGTTTTGGGAAAAAGAACAGGCTTCACTAAAAACGTAATAATGGAAA
 TTGGTAAAACTGAAACTTTTTCTGATGTTTCTGTGAAAACAAATAGAAAGTTTGTCTACTTACTCCAA
 AGATTCAGAAAAGTACTTTGAAACAGAAGCAGTAGAAATGCTAAAGCTTTTATGGAAGATGATGAAGT
 ACAGATCTAAACTGCCAAGTCAAGCCACATTTCTTTTTACATGTCCGAAAATGAGGAAATGGTTT
 TGTCAAATTCAGAATTTGAAAAAGAAGAGGAGAGCCCTTATCTTAGTGGGAGAACCCCAATCAAAAAG
 AAACCTTAAATGAATTTGACAGGATAATAGAAAAATCAAGAAAAATCCTTAAAGGCTTCAAAAAGCACT
 CCAGATGGCACATAAAAAGATCGAAGATTGTTTATGCATCATGTTTCTTTAGAGCCGATTACCTGTGTAC
 CCTTTCGCACAATAAGGAACGTCAAGAGATACAGAAATCCAAATTTTACCACCTGGTCAAGAAATTTCT
 GTCTAAATCTCATTTGTATGAACATCTGACTTTGGAAAAATCTTCAAGCAATTTAGCAGTTTTCAGGACAT
 CCATTTTATCAAGTTTCTGCTACAAGAAATGAAAAATGAGACACTTGATTACTACAGGCAGACCAACCA
 AAGTCTTTGTTCCACCTTTTAAACTAAATCACATTTTACAGAGTTGAACAGTGTGTTAGGAATATTA
 CTTGGAGGAAAACAGACAAAAGCAAAACATTGATGGACATGGCTCTGATGATAGTAAAAATAAGATTAAT
 GACAATGAGATTCATCAGTTTAACAAAAACAACCTCAATCAAGCAGCAGCTGAACTTTACAAAAGTGTG
 AAGAAGAACCTTTAGATTTAATTACAAGTCTTCAAGATGCCAGAGATATACAGGATATGCGAATTAAGAA
 GAAACAAGGCAACGCGTCTTTCCACAGCCAGGAGTCTGTATCTTGAAAAACATCCACTCTGCCTCGA
 ATCTCTCTGAAAGCAGCAGTAGGAGGCCAAGTTCCCTCTGCGTGTCTCATAAACAGCTGTATACGTATG
 GCGTTTCTAAACATTGCATAAAAATTAACAGCAAAAATGCAGAGTCTTTTTCAGTTTACACTGAAGATTA
 TTTTGGTAAGGAAAGTTTATGGACTGAAAAAGGAATACAGTTGGCTGATGGTGGATGGCTCATACCCTCC

AATGATGGAAAGGCTGGAAAAGAAGAATTTTATAGGGCTCTGTGTGACACTCCAGGTGTGGATCCAAAGC
TTATTTCTAGAATTTGGGTTTATAATCACTATAGATGGATCATATGAAACTGGCAGCTATGGAATGTGC
CTTTCCTAAGGAATTTGCTAATAGATGCCTAAGCCCAGAAAGGGTGCTTCTTCAACTAAAATACAGATAT
GATACGGAAATGATAGAAGCAGAAGATCGGCTATAAAAAAGATAATGAAAGGGATGACACAGCTGCAA
AAACACTTGTCTCTGTGTTTCTGACATAATTTTATTGAGCGCAATATATCTGAACTTCTAGCAATAA
AACTAGTAGTGAGATACCCAAAAAGTGCCATTATTGAACTTACAGATGGGTGGTATGCTGTAAAGGCC
CAGTTAGATCCTCCCCTCTTAGCTGTCTTAAAGAATGGCAGACTGACAGTTGGTCAGAAGATTATCTTC
ATGGAGCAGAACTGGTGGGCTCTCCTGATGCCTGTACACCTCTTGAAGCCCAGAATCTTTATGTTAAA
GATTTCTGCTAACAGTACTCGGCCTGCTCGCTGGTATACCAAACCTGGATTCTTCTGACCCTAGACCT
TTTCTCTGCCCTTATCATCGCTTTTTCAGTGATGGAGGAAATGTTGGTGTGTTGATGTAATTATTCAA
GAGCATACCCTATACAGTGGATGGAGAAGACATCATCTGGATTATACATATTTGCAATGAAAGAGAGGA
AGAAAAGGAAGCAGCAAAATATGTGGAGGCCCAACAAAAGAGACTAGAAGCCTTATTCCTAAAATTCAG
GAGGAATTTGAAGAACATGAAGAAAACACAACAAAACCATATTTACCATCACGTGCACTAACAGACAGC
AAGTTCGTGCTTTGCAAGATGGTGCAGAGCTTTATGAAGCAGTGAAGAATGCAGCAGACCCAGCTTACCT
TGAGGGTTATTTTTCAGTGAAGAGCAGTTAAGAGCCTTGAATAATCACAGGCAAATGTTGAATGATAAGAAA
CAAGCTCAGATCCAGTTGAAATTAGGAAGGCCATGGAATCTGCTGAACAAAAGGAACAAGGTTTATCAA
GGGATGTCACAACCGTGTGGAAGTTGCGTATTGTAAGCTATTCAAAAAAGAAAAGATTCACTTACT
GAGTATTTGGCGTCCATCATCAGATTTTATTCTCTGTTAACAGAAGGAAAGAGATACAGAATTTATCAT
CTTGCAACTTCAAATCTAAAAGTAAATCTGAAAGAGCTAACATACAGTTAGCAGCGACAAAAAACTC
AGTATCAACAACCTACCGGTTTCAGATGAAATTTTATTTTTCAGATTTACCAGCCACGGGAGCCCTTCACTT
CAGCAAATTTTATAGATCCAGACTTTTTCAGCCATCTGTTCTGAGGTGGACCTAATAGGATTTGTCGTTTCT
GTTGTGAAAAAACAGGACTTGCCCTTTTCGTCTATTTGTCAGACGAATGTTACAATTTACTGGCAATAA
AGTTTTGGATAGACCTTAATGAGGACATTATTAAGCCTCATATGTTAATTGCTGCAAGCAACCTCCAGTG
G

AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC
TGGATTACAAGGATGACGACGA TAAGGTTTAA

Protein Sequence: >RC400756 representing NM_000059
 Red=Cloning site Green=Tags(s)

MPIGSKERPTFFEIFKTRCNKADLGPISLNFWEEL SSEAPPYNSEPAEESEHKNNNYEPNLFKTPQRKPS
 YNQLASTPIIFKEQGLTLPLYQSPVKELDKFKLDLGRNVPNSRHKSLRTVKTMDQADDVSCPLLNSCLS
 ESPVVLQCTHTVTPQRDKSVVCGSLFHTPKFVKGRQTPKHISESLGAEVDPDMSWSSSLATPTLSTVLI
 VRNEEASETVFPHDTTANVKSYFSNHDESLKKNDRF IASVTDSSENTNOREAASHGFGKTSNGNSFKVNSCK
 DHIGKSMPNVLEDEVYETVVDTSEEDSFLCF SKCRTKNLQKVRTSKTRKKIFHEANADECEKSKNQVKE
 KYSFVSEVEPNDDPLDSNVANQKPFESGSDKI SKEVVVPSLACEWSQL TLSGLNGAQMEKIPLLHSSCD
 QNISEKDLLDTENKRKDFL TSENSLPRISSLPKSEKPLNEETVVNKRDEEQHLESHTDCILAVKQAI SG
 TSPVASSFQGIKKSIFRIRESPKETFNASFSGHMTDPNFKKETEASESGLEIHTVCSQKEDSLCPNLIDN
 GSWPATTTQNSVALKNAGLISTLKKTNKFIYAIHDETSYKGGKIPKDQKSELINCSAQFEANAFEAPLT
 FANADSGLLHSSVKRSCSQNDSEPTLSL TSSFGTILRKCSRNETCSNNTVISQDLDYEAKCNKEKLQL
 FITPEADSLSCLQEQCENDPKSKKVSIDIKEEVLAAACHPVQHSKVEYSDTDFQSQKSLLYDHENASTLI
 LTPTSKDVL SNLVMISRGKESYKMSDKLKGNNYSDVELTKNIPMEKNQDVCALNENYKNVELLPPEKYM
 RVASPSRKVQFNQNTL RVIQKNQEETTSISKITVNPDSEELFSDNENNFVQVANERNLALGNTKELH
 ETDLTVCNEPIFKNSTMVLYGDTGDKQATQVSIK KDLVYVLAENKNSVKQH IKMTLGQDLKSDISLNI D
 KIPEKNNDYMNK WAGLLGPISNHSFGGSFR TASNKEIKLSEHN IKKSKMFFKDIIEEQYPTSLACVEIVNT
 LALDNQKLSKPKQSINTVSAHLQSSVVVSDCKNSHITPQMLFSKQDFNSNHLTPSQKAEITELSTILEE
 SGSQFEFTQFRKPSYILQKSTFEVPEQM TILKTTSEECRDADLHVIMNAPSIGQVDSKQFEGTVEIKR
 KFAGLLKND CNK SASGYLTDENEVGRGFYSAHGTKLVN STEALQKAVKLFSDIENI SEETS AEVHPISL
 SSSKCHDSVVS MFKIENHNDKTVSEKNNK CQLILQNNIEMTTGTVEEITENYKRNTENEDNKYTAASRN
 SHNLEFDGSDSKNDTVCIHKDETDLLFTDQHNICLKL SGQFMKEGNTQIKEDLSDLTFLEVAQAQEACH
 GNTSNKEQLTATKTEQNIKDFETSDTFFQTASGKNISVAKE SFNKIVNFFDQKPEELHNFSLNSELHSDI
 RKNKMDILSYEETDIVKHKILKESVPVGTGNQLVTFQGGQPERDEKIKEPTLLGFHTASGKVKIAKESLD
 KVKNLFDEKEQGTSEITFSHQWAKTLKYREACKDLELACETIEITAAPKCKEMQNSLNNDKNL VSIETV
 VPPKLLSDNLCRQTNELKTSKSI FLKVKVHENV EKETA KSPATCYTNQSPYSVIENSALAFYTCSRKTS
 VSQTSLL EAKKWLREGIFDGQPERINTADYVGNYL YENNSNSTIAENDKNHLSEKQD TYLSNSSMSNSYS
 YHSDEVYNDSGYL SKNKLDSGIEPVLKNVEDQKNTSFSKVISNVKDANAYPQTVNEDICVEELVTSSPC
 KNKNAAIKLSISNSNFEVGPFAFRIASGKIVCVSHETIKKVKDIFTDSFSKVIKENNENKSKICQTKIM
 AGCYEALDSEDILHNSLDNDECSTHSHKVFADIQSEEILQHNQNM SGLKVKISPCDVSLETSDICKC
 SIGK LHKSVSSANTCGIFSTASGKSVQVSDASLQNAQV FSEIEDSTKQVFSKVLFKSNEHSDQLTREEN
 TAIRTPHELISQKGF SYN NVSSAFSGFSTASGKQVSI ESSLHKVKGVL EEFDLIRTEHSLHYSPTS RQ
 NVSKILPRVDRNPEHCVNSEMEKTCSEFKLSNNLNVEGGSSENNHSIKVSPYLSQFQQDKQQLVLGTK
 VSLVENIHVLGKEQASPKNVKMEIGKTEFTSDVPVKTNIEVCSTYSKDSENYFETEAVEIAKAFMEDDEL
 TDSKLP SHATHSLFTCPENEEMVLSNSRIGKRRGEPLILVGEPSIKRNLNDFDRIIENQEKSLKASKST
 PDGTIKDRRLFMHHVSL EPIITCVPFRTTKERQEI QNPNFTAPGQEF LSKSHLYEHLTLEKSSNLAVSGH
 PFYQVSATRNEKMRHLIT TGRPTKVFVPPFKTKSHFHRVEQCVRNINLEENRQKQNI DGHSDDSKNKIN
 DNEIHQFNKNNSNQAAA VFTFKCEEEPLDLITSLQNARDIQDMRIK KKRQRVFPQPGSLYLAKTSTLPR
 ISLKA AVGGQVPSACSHKQLYTYGVSKHCIKINSKNAESFQFHTEDYFGKESLWTGKGIQLADGGWLIPS
 NDGKAGKEEFYRALCDTPGVDPKLSRIWVYNHYRWI IWKLAAMECAF PKEFANRCLSPERVLLQLKYRY
 DTEIDRSRRSAIKKIMERDDTA AKTLVLCVSDIISLSANISETSSNKTSSADTQKVAIIELTDGWYAVKA
 QLDPPLLAVLKNGR LTVGQKII LHGAELVGSPDACTPLEAPESLMLKISANSTRPARWYTKLGFPPDRP
 FPLPLSSLFSDGGNVGCVDVIIQRAYPIQWMEKTSSGLYIFRNEREEKEAAKYVEAQQKRL EALFTKIQ
 EEFEEHEENTTKPYLPSRALTRQQVRALQDGAELYEAVKNAADPAYLEGYFSEEQLRALNNHRQMLNDKK
 QAQIQLEIRKAMESAEQKEQGLSRDVTTVWKLRIVSYSKKEKDSVILSIWRPSSDLYSLLTEGKRYRIYH
 LATSKSKSERANIQLAATKTKYQQLPVSD EILFQIYQPREPLHF SKFLDPDFQPCSEVDLIGFVVS
 VVKKTGLAPFVYLSDECYNLLAIKFWIDLNEDIKPHMLIAASN LQW

SGPTRRRL EQKLI SEEDLAANDILDYKDDDDKV

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:

9381 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:

344 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]