

Product datasheet for **RC400674**

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:	Q2491X
Affected Codon#:	2491
Affected NT#:	7471
Nucleotide Mutation:	BRCA2 Mutant (Q2491X), Myc-DDK-tagged ORF clone of Homo sapiens breast Cancer, early onset (BRCA2) as transfection-ready DNA
Effect:	Breast and/or ovarian 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:	7470 bp
Restriction Sites:	SgfI-RsrII
ORF Nucleotide Sequence:	>RC400674 representing NM_000059 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGCC**

ATGCCTATTGGATCCAAAGAGAGGCCAACATTTTTTGAATTTTTAAGACACGCTGCAACAAAGCAGATT
TAGGACCAATAAGTCTTAATTGGTTTGAAGAACTTTCTTCAGAAGCTCCACCCTATAATTCTGAACCTGC
AGAAGAATCTGAACATAAAAAACAACAATTACGAACCAACCTATTTAAACTCCACAAAGGAAACCATCT
TATAATCAGCTGGCTTCAACTCCAATAATTTCAAAGAGCAAGGGCTGACTCTGCCGCTGTACCAATCTC
CTGTAAGAATTAGATAAATTCAAATTAGACTTAGGAAGGAATGTTCCCAATAGTAGACATAAAGTCT
TCGCACAGTGAAGAACTAAATGGATCAAGCAGATGATGTTTCTGTCCACTTCTAAATCTTGTCTTAGT
GAAAGTCTGTTGTTCTACAATGTACACATGTAACACCACAAAGAGATAAGTCAGTGGTATGTGGGAGTT



[View online »](#)

TGTTTCATACACCAAAGTTTGTGAAGGGTCGTCAGACACCAAAACATATTTCTGAAAGTCTAGGAGCTGA
GGTGGATCCTGATATGTCTTGGTCAAGTTCTTTAGCTACACCACCACCCTTAGTTCTACTGTGCTCATA
GTCAGAAATGAAGAAGCATCTGAAACTGTATTTCTCATGATACTACTGCTAATGTGAAAAGCTATTTTT
CCAATCATGATGAAAGTCTGAAGAAAAATGATAGATTTATCGTCTCTGTGACAGACAGTGA AACACAAA
TCAAAGAGAAGCTGCAAGTCATGGATTTGGAAAAACATCAGGGAATTCATTTAAAGTAAATAGCTGCAAA
GACCACATTGGAAAGTCAATGCCAAATGTCTAGAAGATGAAGTATATGAAACAGTTGTAGATACCTCTG
AAGAAGATAGTTTTTTCATTATGTTTTTCTAAATGTAGAACAAAAATCTACAAAAAGTGAAGAAAGCA
GACTAGGAAAAAATTTTCCATGAAGCAACGCTGATGAATGTGAAAAATCTAAAAACCAAGTGAAAGAA
AAATACTCATTTGTATCTGAAGTGGAACCAATGATACTGATCCATTAGATTCAAATGTAGCAAATCAGA
AGCCCTTTGAGAGTGGAAGTGACAAAATCTCCAAGGAAGTTGTACCGTCTTTGGCCTGTGAATGGTCTCA
ACTAACCTTTTCAAGTCTAAATGGAGCCAGATGGAGAAAAATACCCTATTGCATATTTCTTTCATGTGAC
CAAAATATTTTCAAGAAAAGACCTATTAGACACAGAGAACAAAAGAAAGAAAGATTTTCTTACTTCAGAGA
ATTCTTTGCCACGTATTTCTAGCCTACCAAAATCAGAGAAGCCATTAAATGAGGAAACAGTGGTAAATAA
GAGAGATGAAGAGCAGCATCTTGAATCTCATACTGACTGCATTCTTGCAGTAAAGCAGGCAATATCTGGA
ACTTCTCCAGTGGCTTTCATTTCAAGGTATCAAAAAGTCTATATTCAGAATAAGAGAATCACCTAAAG
AGACTTTCAATGCAAGTTTTTCAAGTCAATGACTGATCCAAACTTTAAAAAAGAAACTGAAGCCTCTGA
AAGTGGACTGGAATACATACTGTTTGCTCACAGAAGGAGGACTCCTTATGTCCAAATTTAATTGATAAT
GGAAGCTGGCCAGCCACCACCACAGAAATCTGTAGCTTTGAAGAATGCAGGTTAATATCCACTTTGA
AAAAGAAAAACAAATAAGTTTATTTATGCTATACATGATGAAACATCTTATAAAGGAAAAAATACCGAA
AGACCAAAAAATCAGAATAATTAAGTGTTCAGCCAGTTTGAAGCAATGCTTTTGAAGCACCATTACA
TTTGCAAAATGCTGATTCAGGTTTATTGCATTCTCTGTGAAAAGAAGCTGTTTACAGAAATGATTTGAAG
AACCAATGCTGATTCAGGTTTATTGCATTCTCTGTGAAAAGAAGCTGTTTACAGAAATGAAACATGTTT
TAATAATACAGTAATCTCTCAGGATCTTGATTATAAAGAAGCAAAATGTAAATAAGGAAAAACTACAGTTA
TTTATTACCCAGAAGCTGATTCTGTGATGCCTGCAGGAAGGACAGTGTGAAAAATGCTCAAAAAAGCA
AAAAAGTTTCAAGATATAAAGAAGAGGTCTGGCTGCAGCATGTACCCAGTACAACATTCAAAAGTGGAA
ATACAGTGATACTGACTTTCAATCCCAGAAAAGTCTTTTATATGATCATGAAAAATGCCAGCACTCTTATT
TTAACTCTACTTCCAAGGATGTTCTGTCAAACCTAGTCATGATTTCTAGAGGCAAGAATCATACAAAA
TGTCAGACAAGCTCAAAGGTAACAATTATGAATCTGATGTTGAATTAACCAAAAAATTTCCCATGGAAAA
GAATCAAGATGTATGTGCTTTAAATGAAAATTAAAAAACGTTGAGCTGTTGCCACCTGAAAAATACATG
AGAGTAGCATCACCTTCAAGAAAGGTACAATTCACCAAAAACACAATCTAAGAGTAATCAAAAAAATC
AAGAAGAACTACTTCAATTTCAAAAATAACTGTCAATCCAGACTCTGAAGAATTTTCTCAGACATGA
GAATAATTTTGTCTTCCAAGTAGCTAATGAAAGGAATAATCTTGCTTTAGGAAATACTAAGGAATTCAT
GAAACAGACTTGACTTGTGTAACCGAACCCATTTTCAAGAATCTACCATGGTTTTATATGGAGACACAG
GTGATAAACAAGCAACCCAAGTGTCAATTA AAAAAGATTTGGTTTATGTTCTTGCAGAGGAGAACAAAA
TAGTGTAAAGCAGCATATAAAAAAGACTCTAGGTCAAGATTTAAATCGGACATCTCCTTGAATATAGAT
AAAATACCAGAAAAAATAATGATTACATGAACAAATGGGCAGGACTCTTAGGTCCAATTTCAAATCACA
GTTTTGGAGGTAGCTTCAAGACAGCTTCAAATAAGGAAATCAAGCTCTCTGAACATAACATTAAGAAGAG
CAAAATGTTCTTCAAAGATATTGAAGAACAATATCCTACTAGTTTAGCTTGTGTTGAAATGTAATACC
TTGGCATTAGATAATCAAAAGAACTGAGCAAGCCTCAGTCAATTAATACTGTATCTGCACATTTACAGA
GTAGTGTAGTTGTTTCTGATTGTAAAAATAGTCATATAACCCCTCAGATGTTATTTTCCAAGCAGGATTT
TAATTCAAACCATAATTTAACACCTAGCCAAAAGGCAGAAATTACAGAATTTTCTACTATATTAGAAGAA
TCAGGAAGTCAGTTTGAATTTACTCAGTTTAGAAAACCAAGCTACATATTGCAGAAGAGTACATTTGAAG
TGCCTGAAAACCAAGATGACTATCTTAAAGACCCTTCTGAGGAATGCAGAGATGCTGATCTTCATGTCAT
AATGAATGCCCATCGATTGGTCAGGTAGACAGCAGCAAGCAATTTGAAGGTACAGTTGAAATTAACGG
AAGTTTGTGCTGGCCTGTTGAAAAATGACTGTAACAAAAGTCTTCTGGTTATTTAACAGATGAAAAATGAAG
TGGGGTTTAGGGCTTTTATTCTGCTCATGGCACAAAAGTGAATGTTTCTACTGAAGCTCTGCAAAAAGC
TGTGAAACTGTTTAGTGATATTGAGAATATTAGTGAGGAACTCTGCAGAGGTACATCCAATAAGTTTA
TCTTCAAGTAAATGTCATGATTCTGTTGTTCAATGTTTAAAGATGAAAAATCATAATGATAAACTGTAA
GTGAAAAAATAATAATGCCAACTGATATTACAAAATAATTTGAAATGACTACTGGCACTTTTGTGGA
AGAAATTAAGTAAAAATCAAGAGAAATACTGAAAATGAAGATAACAAATATACTGCTGCCAGTAGAAAT
TCTCATAACTTAGAATTTGATGGCAGTGATTCAAGTAAAAATGATACTGTTTGTATTATAAAGATGAAA
CGGACTTGCTATTTACTGATCAGCACAACATATGTCTTAAATTTCTGGCCAGTTTATGAAGGAGGGAAA

CACTCAGATTAAGAAGATTTGTCAAGTTAACTTTTTGGAAGTTGCGAAAGCTCAAGAAGCATGTCAT
GGTAATACTTCAATAAAGAACAGTTAACTGCTACTAAAACGGAGCAAAATATAAAAGATTTTGAGACTT
CTGATACATTTTTTTCAGACTGCAAGTGGGAAAAATATTAGTGTGCGCCAAAGAGTCATTTAATAAAATTTG
AAATTTCTTTGATCAGAAACCAGAAGATTGCATAACTTTTCTTAAATTTCTGAATTACATTCTGACATA
AGAAAGAACAAAATGGACATTCTAAGTTATGAGGAAACAGACATAGTTAAACACAAAATCTGAAAGAAA
GTGTCCAGTTGGTACTGGAATCAACTAGTGACCTCCAGGGACAACCCGAACGTGATGAAAAGATCAA
AGAACCCTACTCTATTGGTTTTTCATACAGCTAGCGGGAAAAAAGTTAAATTTGCAAAGGAATCTTTGGAC
AAAGTGAAAAACCTTTTTGATGAAAAAGCAAGGTAAGTAAATCACCAGTTTTAGCCATCAATGGG
CAAAGACCCTAAAGTACAGAGAGCCTGTAAAGACCTGAATTAGCATGTGAGACCATTGAGATCACAGC
TGCCCCAAAGTGTAAAGAAATGCAGAATTCTCTCAATAATGATAAAAACTTGTTTCTATTGAGACTGTG
GTGCCACCTAAGCTCTAAGTGATAATTTATGTAGACAACTGAAATCTCAAAACATCAAAAAGTATCT
TTTTGAAAGTTAAAGTACATGAAATGTAGAAAAAGAACAGCAAAAAGTCTGCAACTTGTACACAAA
TCAGTCCCCTTATTCAGTCATTGAAAATTCAGCCTTAGCTTTTTACACAAGTTGTAGTAGAAAACTTCT
GTGAGTCAGACTTCATTACTGGAAGCAAAAAATGGCTTAGAGAAGGAATATTTGATGGTCAACCAGAAA
GAATAAATACTGCAGATTATGTAGGAAATTTTGTATGAAAATTAATCAAACAGTACTATAGCTGAAAA
TGACAAAAATCATCTCTCCGAAAAACAAGATACTTATTAAGTAAACAGTAGCATGTCTAACAGCTATTCC
TACCATTCTGATGAGGTATATAATGATTGAGGATATCTCTCAAAAAATAAACTTGATTCTGGTATTGAGC
CAGTATTGAAGAATGTTGAAGATCAAAAAACACTAGTTTTTCCAAAGTAATATCCAATGTAAGAGATGC
AAATGCATACCCACAACTGTAATGAAGATATTTGCGTTGAGGAACTTGTGACTAGCTCTTCCACCTGC
AAAAATAAAAAATGCAGCCATTAATTTGCCATATCTAATAGTAATAATTTTGAGGTAGGGCCACCTGCAT
TTAGGATAGCCAGTGGTAAATCGTTTTGTGTTTCATGAAACAATTAAGGAAAGTAAAGACATATTTAC
AGACAGTTTCAGTAAAGTAATTAAGGAAAAACAGGAGAATAAATCAAAAAATTTGCCAAACGAAAATATG
GCAGTTGTTACGAGGCATTGGATGATTGAGGATATCTTTCATAACTCTAGATAATGATGAATGTA
GCACGCATTCACATAAGGTTTTTGTGACATTCAGAGTGAAGAAATTTTACAACATAAACAAAATATGTC
TGGATTGGAGAAAGTTTCTAAAAATCACCTTGTGATGTTAGTTTGGAACTTCAGATATATGAAATGT
AGTATAGGGAAGCTTCATAAGTCAGTCTCATCTGCAAATCTTGTGGGATTTTTAGCACAGCAAGTGAA
AATCTGTCCAGGTATCAGATGCTTATTACAAAACGCAAGACAAGTGTCTTGAATAGAAAGATAGTAC
CAAGCAAGTCTTTTCCAAAGTATTGTTTAAAAGTAACGAACATTCAGACCAGCTCACAAGAGAAGAAAAT
ACTGCTATACGTACTCCAGAACATTTAATATCCAAAAAGGCTTTTTCATATAATGTGGTAAATTCATCTG
CTTTCTCTGGATTTAGTACAGCAAGTGGAAAGCAAGTTCCATTTTGAAGGTTCTTACACAAAGTTAA
GGGAGTGTAGAGGAATTTGATTTAATCAGAAGTGGGATAGTCTTCACTATTCACCTACGTCTAGACAA
AATGTATCAAAAATACCTCCTCGTGTGATAAGAGAAACCCAGAGCACTGTGAAACTCAGAAATGGAAA
AAACCTGCAGTAAAGAATTTAAATTATCAAATAACTTAAATGTTGAAGGTGGTTCTTCAGAAAATATCA
CTCTATTAAGGTTTCTCCATATCTCTCAATTTCAACAAGACAAAACAACAGTTGGTATTAGGAACCAAA
GTGTCACTTGTGAGAACATTCATGTTTTGGGAAAAAGAACAGGCTTCCACTAAAAACGTAATAATGGAAA
TTGGTAAAACTGAAACTTTTTCTGATGTTCTGTGAAAACAAATAGAAAGTTTGTCTACTTACTCCAA
AGATTCAGAAAACACTTTGAAACAGAAGCAGTAGAAATGCTAAAGCTTTTATGGAAGATGATGAAGT
ACAGATCTAAACTGCCAAGTCATGCCACATTTCTTTTTACATGTCCGAAAATGAGGAAATGGTTT
TGTCAAATTCAGAATTTGAAAAAGAGAGGAGAGCCCTTATCTTAGTGGGAGAACCCCAATCAAAAAG
AACTTATTAATGAATTTGACAGGATAATAGAAAATCAAGAAAATCCTTAAAGGCTTCAAAAAGCACT
CCAGATGGCACATAAAAAGATCGAAGATTGTTTATGCATCATGTTTCTTTAGAGCCGATTACCTGTGTAC
CCTTTCGCACAACATAAGGAACGTCAAGAGATACAGAAATCCAAATTTTACCACCTGGTCAAGAAATTTCT
GTCTAAATCTCATTTGTATGAACATCTGACTTTGGAAAAATCTTCAAGCAATTTAGCAGTTTTCAGGACAT
CCATTTTATCAAGTTTCTGCTACAAGAAATGAAAAATGAGACACTTGATTACTACAGGCAGACCAACCA
AAGTCTTTGTTCCACCTTTTAAACTAAATCACATTTTACAGAGTTGAACAGTGTGTTAGGAATATTA
CTTGGAGGAAAACAGACAAAAGCAAAACATTGATGGACATGGCTCTGATGATAGTAAAAATAAGATTAAT
GACAATGAGATTCATCAGTTTAAACAAAACAACCTCAATCAAGCAGCAGCTGTAACCTTTCACAAAGTGTG
AAGAAGAACCTTTAGATTTAATTACAAGTCTTCAGAATGCCAGAGATATA

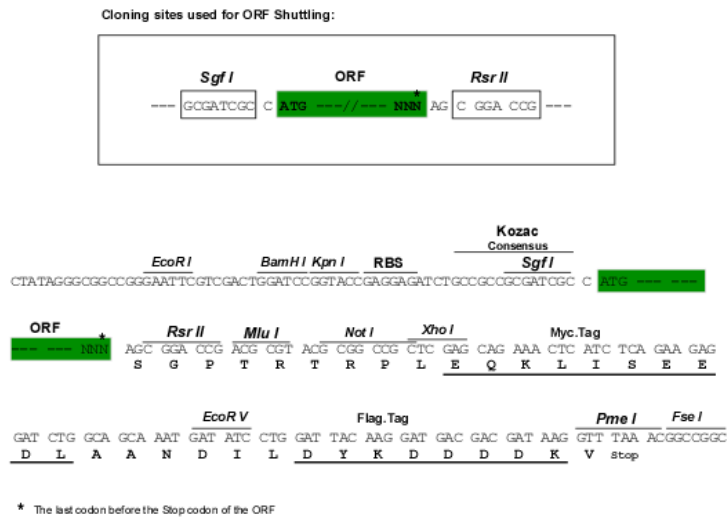
AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAAAACCTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC
TGGATTACAAGGATGACGACGA TAAGGTTTAA

Protein Sequence: >RC400674 representing NM_000059
 Red=Cloning site Green=Tags(s)

MPIGSKERPTFFEIFKTRCNKADLGPISLWNWFEELESSEAPPYNSEPAEESEHKNNNYEPNLFKTPQRKPS
 YNQLASTPIIFKEQGLTLPLYQSPVKELDKFKLDLGRNVNPSRHKSLRTVKTMDQADDVSCPLLNSCLS
 ESPVVLQCTHVTPOQDKSVVCGSLFHTPKFVKGRQTPKHISESLGAEVDPDMSWSSSLATPPTLSTVLI
 VRNEEASETVFPHDTTANVKSYFSNHDESLKKNDRFIASVTDSENTNQREAASHGFGKTSGNSFKVNSCK
 DHIGKSMPNVLEDEVYETVVDTSEEDSFSLCFSKCRTKNLQKVRTSKTRKKIFHEANADECEKSKNQVKE
 KYSFVSEVEPNDDPLDSNVANQKPFESGSDKI SKEVVVPSLACEWSQLTSLGNGAQMEKIPLLHSSCD
 QNISEKDLLDTENKRKDFLTSENSLPRISLSEKPLNEETVVNKRDEEQHLESHTDCILAVKQAIISG
 TSPVASSFQGIKKSIFRIRESPKETFNASFSGHMTDPNFKKETEASESGLEIHTVCSQKEDSLCPNLIDN
 GSWPATTTQNSVALKNAGLISTLKKTNKFIYAIHDETSYKGGKIPKQKSELINCSAQFEANAFEAPLT
 FANADSGLLHSSVKRSCSQNDSEPTLSLTSSFGTILRKCSRNETCSNNTVISQDLDYEAKCNKEKLQL
 FITPEADSLSCLQEGQCENDPKSKKVSIDIKEEVLAAACHPVQHSKVEYSDTDFQSQKSLLYDHENASTLI
 LTPTSKDVL SNLVMISRGKESYKMSDKLKGNNYSDVELTKNIPMEKNQDVCALNENYKNNVELLPPEKYM
 RVASPSRKVQFNQNTLNRVIQKNQEETTSISKITVNPDEELFSDNENNFVQVANERNLALGNTKELH
 ETDLTCVNEPIFKNSTMVLVYGDGDKQATQVSIKKDLVYVLAENKNSVKQHIKMTLGQDLKSDISLNI
 KIPEKNNDYMNKWAAGLLGPISNHSFGGSFRASNKEIKLSEHNKIKSKMFFKDIIEEQYPTSLACVEIVNT
 LALDNQKLSKPKQSINTVSAHLQSSVVVSDCKNSHITPQMLFSKQDFNSNHLTPSQKAEITELSTILEE
 SGSQFEFTQFRKPSYILQKSTFEVPEQMILKTTSEECRDLHVMNAPSIGQVDSKQFEGTVEIKR
 KFAGLLKNDCKNSASGYLTDENEVGRGFYSAHGKLVNSTEALQKAVKLFSDIENISEETSAEVHPISL
 SSKKCHDSVVSFMFKIENHNDKTVSEKNNKQQLILQNNIEMTTGTFVEEITENYKRNTENEDNKYTAASRN
 SHNLEFDGSDSSKNDTVCIHKDETDLLFTDQHNI CLKLSGQFMKEGNTQIKEDLSLTFLEVAKAQEACH
 GNTSNKEQLTATKTEQNIKDFETSDTFFQTASGKNISVAKESFNKIVNFFDQKPEELHNFSLNSELHSDI
 RKNKMDILSYEETDIVKHKILKESVPVGTGNQLVTFQGGPERDEKIKEPTLLGFHTASGKVKIAKESLD
 KVKNLFDKEQGTSEITSFSHQWAKTLKYREACKDLELACETIEITAAPKCKEMQNSLNDKNLVSIEYV
 VPPKLLSDNLCRQTEMLKTSKIFLKVKHENVEKETAKSPATCYTNQSPYSVIENSALAFYTCSRKTS
 VSQTSLLLEAKKWLREGIFDQPERINTADYVGNLYENNSNSTIAENDKNHLSEKQDYLSSSSMSNSYS
 YHSDEVYNDSGYL SKNKLDSGIEPVLKNVEDQKNTSFSKVISNVKDANAYPQTVNEDICVEELVTSSPC
 KNKNAAIKLSISNSNFEVGPFAFRIASGKIVCVSHETIKKVKDIFTDSFSKVIKENNENKSKICQTKIM
 AGCYEALDDSEDILHNSLDNDECSTHSHKVFADIQSEEILQHNQNMGLKVKISPCDVSLETSDICKC
 SIGKLVHSSVSSANTCGIFSTASGKSVQVSDASLQARQVFSEIEDSTKQVFSKVLFKSNEHSDQLTREEN
 TAIRTPHEHLISQKGFYNVNVSSAFSGFSTASGKQVSILESSLHKVKGVL EEFDLIRTEHSLHYSPTRQ
 NVSKILPRVDRNPEHCVNSEMEKTCSEFKLSNNLNVEGGSSENNHSIKVSPYLSQFQQDKQLVLGK
 VSLVENIHVLGKEQASPKNVKMEIGKTEFTSDVPVKTNIEVCSTYSKDSYFETEAVEIAKAFMEDDEL
 TDSKLP SHATHSLFTCPENEEMVLSNSRIGKRRGEPLILVGEPSIKRLLNEFDRIIENQEKSLKASKST
 PDGTIKDRRLFMHVSLEPITCVPFRTTKERQEQNPNTAPGQEFLLSKSHLYEHLTLEKSSSNLAVSGH
 PFYQVSATRNEKMRHLITTRPTKVFVPPFKTKSHFHRVEQCVRNINLEENRQKQNIIDGHGSDSSKNKIN
 DNEIHQFNKNNNSNQAAAVFTKCEEEPLDLITSLQNARDI

SGPTRRRLEQKLI 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:

7470 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: 273.9 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]