

## Product datasheet for **RC403033**

### **BRCA1 (NM\_007294) Human Mutant ORF Clone**

#### Product data:

Product Type:	Mutant ORF Clones
Product Name:	BRCA1 (NM_007294) Human Mutant ORF Clone
Mutation Description:	Q1200X
Affected Codon#:	1200
Affected NT#:	3598
Nucleotide Mutation:	BRCA1 Mutant (Q1200X), Myc-DDK-tagged ORF clone of Homo sapiens breast Cancer, early onset (BRCA1), transcript variant 1 as transfection-ready DNA
Effect:	Breast cancer
Symbol:	BRCA1
Synonyms:	BRCAI; BRCC1; BROVCA1; FANCS; IRIS; PNCA4; PPP1R53; PSCP; RNF53
E. coli Selection:	Kanamycin (25 ug/mL)
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
Tag:	Myc-DDK
ACCN:	NM_007294
ORF Size:	3597 bp
Restriction Sites:	Sgfl-Mlul
ORF Nucleotide Sequence:	>RC403033 representing NM_007294 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**GCGATCGCC**

ATGGATTATCTGCTCTTCGCGTTGAAGAAGTACAAAATGTCATTAATGCTATGCAGAAAATCTTAGAGT  
GTCCCATCTGTCTGGAGTTGATCAAGGAACCTGTCTCCACAAAGTGTGACCACATATTTTGCAAATTTTG  
CATGCTGAAACTTCTCAACCAGAAGAAAGGGCCTTCACAGTGTCTTTATGTAAGAATGATATAACCAA  
AGGAGCCTACAAGAAAGTACGAGATTTAGTCAACTTGTGAAGAGCTATTGAAAATCATTGTGCTTTTC  
AGCTTGACACAGGTTTGGAGTATGCAACAGCTATAATTTTGCAAAAAGGAAAATAACTCTCCTGAACA  
TCTAAAAGATGAAGTTTCTATCATCAAAGTATGGGCTACAGAAACCGTCCAAAAGACTTCTACAGAGT  
GAACCCGAAAATCCTTCTTGCAGGAAACAGTCTCAGTGTCCAACCTCTAACCTTGGAACTGTGAGAA



[View online »](#)

CTCTGAGGACAAAGCAGCGGATACAACCTCAAAAGACGTCTGTCTACATTGAATTGGGATCTGATTCTTC  
TGAAGATACCGTTAATAAGGCAACTTATTGCAGTGTGGGAGATCAAGAATTGTTACAAATCACCCCTCAA  
GGAACCAGGGATGAAATCAGTTTGGATTCTGCAAAAAGGCTGCTTGTGAATTTCTGAGACGGATGTA  
CAAATACTGAACATCATCAACCCAGTAATAATGATTTGAACACCACTGAGAAGCGTGCAGCTGAGAGGCA  
TCCAGAAAAGTATCAGGGTAGTTCTGTTCAAACCTGCATGTGGAGCCATGTGCCACAAATACTCATGCC  
AGCTCATTACAGCATGAGAACAGCAGTTTATTACTCACTAAAGACAGAATGAATGTAGAAAAGGCTGAAT  
TCTGTAATAAAAAGCAAACAGCCTGGCTTAGCAAGGAGCCAACATAACAGATGGGCTGGAAGTAAGGAAAC  
ATGTAATGATAGGCGGACTCCCAGCACAGAAAAAAGGTAGATCTGAATGCTGATCCCCTGTGTGAGAGA  
AAAGAATGGAATAAGCAGAACTGCCATGCTCAGAGAATCCTAGAGATACTGAAGATGTTCTTGGATAA  
CACTAAATAGCAGCATTAGAAAAGTAAATGAGTGGTTTTCCAGAAGTGAAGTGTAGGTTCTGATGA  
CTCACATGATGGGGAGTCTGAATCAAATGCCAAAGTAGCTGATGATTGGACGTTCTAAATGAGGTAGAT  
GAATATTCTGGTTCTTCCAGAGAAAATAGACTTACTGGCCAGTGCCTCATGAGGCTTTAATATGTA  
GTGAAAGAGTTCACTCCAATCAGTAGAGATAATATTGAAGACAAAATATTTGGGAAAACCTATCGGAA  
GAAGGCAAGCCTCCCAACTTAAGCCATGTAAGTAAAATCTAATTATAGGAGCATTGTTACTGAGCCA  
CAGATAATACAAGAGCGTCCCCTCACAATAAATTAAGCGTAAAAGGAGACCTACATCAGGCTTCATC  
CTGAGGATTTTATCAAGAAAGCAGATTTGGCAGTTCAAAAGACTCCTGAAATGATAAATCAGGGAAC  
CCAAACCGAGCAGAATGGTCAAGTGAATATTACTAATAGTGGTCATGAGAATAAAACAAAAGGTGAT  
TCTATTAGAATGAGAAAAATCCTAACCAATAGAATCACTCGAAAAAGAATCTGCTTTCAAACGAAAG  
CTGAACCTATAAGCAGCAGTAAAGCAATATGGAAGTGAATTAATAATCCACAATCAAAGCAGCTAA  
AAAGAATAGGCTGAGGAGGAAGTCTTCTACCAGGCATATTCATGCGCTTGAAGTGTAGTGTAGTAA  
CTAAGCCACCTAATTGTACTGAATTGCAAAATGATAGTTGTTCTAGCAGTGAAGAGATAAAGAAAA  
AGTACAACCAATGCCAGTCAAGCAGCAGAAAACCTACAACCTCATGGAAGGTAAGAAAGTGAAGTGA  
AGCCAAAGAGTAACAAGCCTAAGTGAACAGCAAGTAAAGACATGACAGCGATACTTTCCAGAGCTG  
AAGTTAACAAATGCACCTGGTTCTTTTACTAAGTGTTCAAATACCAAGTGAAGTAAAGAATTTGCAATC  
CTAGCCTTCCAAGAGAAGAAAAAGAGAAAAGTAAAGTGTCTAATAATGCTGAAGACCC  
CAAAGTCTCATGTTAAGTGGAGAAAGGTTTTGCAAAGTGAAGATCTGTAGAGAGTAGCAGTATTTCA  
TTGGTACTGTTACTGATTATGGCACTCAGGAAAGTATCTCGTTACTGGAAGTGAAGTGTAGGAAAG  
CAAAAACAGAACCAATAAATGTGTGAGTCAAGTGTGCAGCATTGAAAACCCCAAGGACTAATTCATG  
TTGTTCCAAAGATAATAGAAATGACACAGAAGGCTTTAAGTATCCATTGGGACATGAAGTAAACCAAGT  
CGGAAACAAGCATAGAAATGGAAGAAAGTGAAGTGTGCTCAGTATTTGCAGAATACATTCAAGGTTT  
CAAAGCGCAGTCAATTTGCTCCGTTTTCAAATCCAGGAAATGCAGAAGAGGAATGTGCAACATTCTG  
CCTCTGGGTCCTTAAAGAAACAAAGTCCAAAAGTCACTTTTGAATGTGAACAAAAGGAAGAAAATCAA  
GGAAAGAATGAGTCTAATATCAAGCCTGTACAGACAGTAAATCACTGCAGGCTTTCTGTGGTTGGTC  
AGAAAGATAAGCCAGTTGATAATGCCAAATGTAGTATCAAAGGAGGCTCTAGGTTTTGTCTATCATCTCA  
GTTCAGAGGCAACGAACTGGACTCATTACTCCAAATAAATGGAAGTAAAGTAAAGTAAAGTAAAGTAA  
CCACCCTTTTCCCATCAAGTCAATTTGTTAAACTAAATGTAAGAAAAATCTGCTAGAGGAAAACCTTTG  
AGGAACATTCAATGTACCTGAAAGAGAAATGGGAAATGAGAACATCCAAGTACAGTGAACACAATTAG  
CCGTAATAACATTAGAGAAAATGTTTTAAAGAAGCCAGCTCAAGCAATTAATGAAGTAGGTTCCAGT  
ACTAATGAAGTGGCTCCAGTATTAATGAAATAGGTTCCAGTGTGAAAACATTCAAGCAGAAGTGAAGTA  
GAAACAGAGGGCCAAAATGAAATGCTATGCTTAGATTAGGGTTTTGCAACCTGAGGCTATAAAACAAAG  
TCTTCTGGAAGTAATTGTAAGCATCCTGAAATAAAAAGCAAGAATATGAAGAAGTGTTCAGACTGTT  
AATACAGATTTCTCTCCATCTGATTTAGATAAATAGAACAGCCTATGGGAAAGTGTGATGATCTC  
AGGTTTGTCTGAGACACCTGATGACCTGTTAGATGATGGTGAATAAAGGAAGATACTAGTTTGTGTA  
AAATGACATTAAGGAAAGTTCTGCTGTTTTAGCAAAGCGTCCAGAAAGGAGAGCTTAGCAGGAGTCT  
AGCCCTTTCACCCATACACATTTGGCT

AGCGGACCGACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC  
TGGATTACAAGGATGACGACGA TAAGGTTTAA

**Protein Sequence:** >RC403033 representing NM\_007294  
 Red=Cloning site Green=Tags(s)

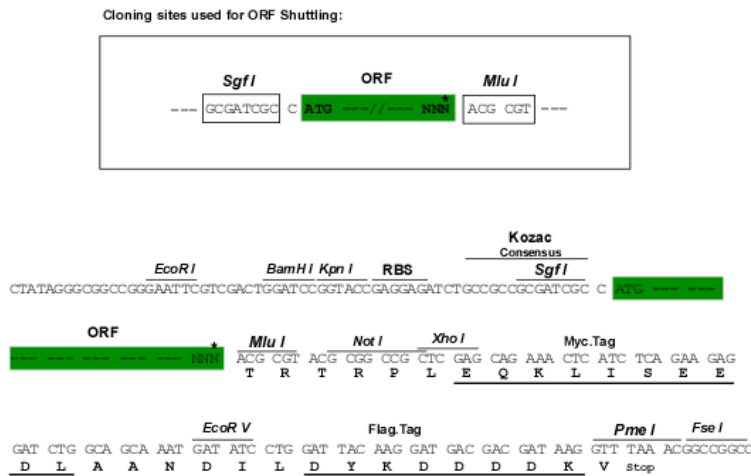
MDLSALRVEEVQNVINAMQKILECPICLELIEKPVSTKCDHIFCKFCMLKLLNQKKGPSQCPLCKNDITK  
 RSLQESTRFSQLVEELLKIICAFQLDTGLEAYNSYNFAKKENNSPEHLKDEVSI IQSMGYRNRARLLQS  
 EPENPSLQETSLSVQLSNLGTVRTLRKQRIQPQKTSVYIELGSDSSEDTVNKATYCSVGDQELLQITPQ  
 GTRDEISLDSAKKAACEFSETDVTNTEHHQPSNNDLNTTEKRAAERHPEKYQGSSVSNLHVPCGTNTHA  
 SSLQHENSLLLLTKDRMNVEKAFCNKSKQPGLARSQHNRWAGSKETCNDRRTPSTEKKVDLNADPLCER  
 KEWNNKQKLPCCSENPRDTEVPWITLNSSIQKVNEWFSRSEDELLGSDSDHGESESNKAVADVLDVLENEVD  
 EYSGSSEKIDLLASDPHEALICKSERVHKSVESNIEDKIFGKTYRKKASLPNLSHVTENLIIGAFVTEP  
 QIIQERPLTNKLRKRRTSGLHPEDFIKKADLAVQKTPEMINQGTNQTQEQNGQVMNITNSGHENKTKGD  
 SIQNEKNPNPIESLEKESAFKTKAEPISSSISNMELELNIHNSKAPKKNRLRRKSSTRHIALELVVSRN  
 LSPNCTELQIDSCSSSEI KKKKYNQMPVHRNRLQLMEGKEPATGAKKSNKPNEQTSKRHDSDTFPEL  
 KLTNAPGSFTKCSNTSELKEFVNPSLPREEKEELETVKVSNNAEADPKDMLSGERVLQTERSVESSSIS  
 LVPGTGYGTQESISLLEVSTLGAKTEPNKCVSQCAAFENPKGLIHGCSKDNRNDEGFKYPLGHEVNH  
 RETSIEMEESELDAQYLQNTFKVSKRQSFAPFSNPNAEEECATFSAHSGSLKKQSPKVTFECEQKEENQ  
 GKNESNIKPVQTVNITAGFPVVGQKDKPVDNAKCSIKGGRFCLSSQFRGNETGLITPNKHGLLQNPYRI  
 PPLFPKISFVKTKCKKNLLEENFEHSMSPEREMGNENIPSTVSTISRNNIRENVFKEASSSININEVGSS  
 TNEVGSSINEIGSSDENIQAELGRNRGPKLNAMLRLGVLQPEVYKQSLPGSNCKHPEIKKQEYEEVVQTV  
 NTDFSPYLISDNLEQPMGSSHASQVCSETPDDLDDGEIKEDTSAFENDIKESSAVF SKSVQK GELSRSP  
 SPFTHTHLA

SGP TRRRLEQKLI SEEDLAANDILDYKDDDDKV

**Restriction Sites:**

SgfI-MluI

**Cloning Scheme:**



\* The last codon before the Stop codon of the ORF

**OTI Disclaimer:** Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at [custsupport@origene.com](mailto:custsupport@origene.com) or by calling 301.340.3188 option 3 for pricing and delivery.

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\\_009225](#)

**RefSeq Size:** 3597 bp

**RefSeq ORF:** 5592 bp

**Locus ID:** 672

**Cytogenetics:** 17q21.31

**Domains:** BRCT, RING

**Protein Families:** Druggable Genome, Transcription Factors

**Protein Pathways:** Ubiquitin mediated proteolysis

**MW:** 131.9 kDa

**Gene Summary:**

This gene encodes a 190 kD nuclear phosphoprotein that plays a role in maintaining genomic stability, and it also acts as a tumor suppressor. The BRCA1 gene contains 22 exons spanning about 110 kb of DNA. The encoded protein combines with other tumor suppressors, DNA damage sensors, and signal transducers to form a large multi-subunit protein complex known as the BRCA1-associated genome surveillance complex (BASC). This gene product associates with RNA polymerase II, and through the C-terminal domain, also interacts with histone deacetylase complexes. This protein thus plays a role in transcription, DNA repair of double-stranded breaks, and recombination. Mutations in this gene are responsible for approximately 40% of inherited breast cancers and more than 80% of inherited breast and ovarian cancers. Alternative splicing plays a role in modulating the subcellular localization and physiological function of this gene. Many alternatively spliced transcript variants, some of which are disease-associated mutations, have been described for this gene, but the full-length natures of only some of these variants has been described. A related pseudogene, which is also located on chromosome 17, has been identified. [provided by RefSeq, May 2020]