

Product datasheet for RC402897

BRCA1 (NM_007294) Human Mutant ORF Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	Mutant ORF Clones	
Product Name:	BRCA1 (NM_007294) Human Mutant ORF Clone	
Mutation Description:	Q356X	
Affected Codon#:	356	
Affected NT#:	1066	
Nucleotide Mutation:	BRCA1 Mutant (Q356X), 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:	1065 bp	
Restriction Sites:	Sgfl-Mlul	

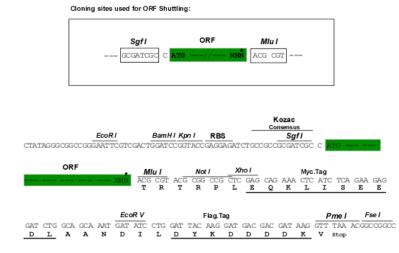


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	BRCA1 (NM_007294) Human Mutant ORF Clone – RC402897
ORF Nucleotide Sequence:	<pre>>RC402897 representing NM_007294 Red=Cloning site Blue=ORF Green=Tags(s)</pre>
	TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC GCC <mark>GCGATCGC</mark> C
	ATGGATTTATCTGCTCTTCGCGTTGAAGAAGTACAAAATGTCATTAATGCTATGCAGAAAATCTTAGAGT GTCCCATCTGTCTGGAGTTGATCAAGGAACCTGTCTCCACAAAGTGGACCACATATTTTGCAAATTTTG CATGCTGAAACTTCTCAACCAGAAGAAAGGGCCTTCACAGTGTCCTTTATGTAAGAATGATATAACCAAA AGGAGCCTACAAGAAAGTACGAGATTTAGTCAACTTGTTGAAGAGCTATTGAAAATCATTTGTGCTTTC AGCTTGACACAGGTTTGGAGTATGCAAACAGCTATAATTTTGCAAAAAGGAAAATAACTCTCTCGAACA TCTAAAAGATGAAGTTTCTATCATCCACAGTATGGGCTACAGAAAACGGCACAAAGACTCTCACAGAGT GAACCCGAAAATCCTTTCCTT
Protein Sequence	<pre>Red=Cloning site Green=Tags(s) MDLSALRVEEVQNVINAMQKILECPICLELIKEPVSTKCDHIFCKFCMLKLLNQKKGPSQCPLCKNDITK</pre>
	RSLQESTRFSQLVEELLKIICAFQLDTGLEYANSYNFAKKENNSPEHLKDEVSIIQSMGYRNRAKRLLQS EPENPSLQETSLSVQLSNLGTVRTLRTKQRIQPQKTSVYIELGSDSSEDTVNKATYCSVGDQELLQITPQ GTRDEISLDSAKKAACEFSETDVTNTEHHQPSNNDLNTTEKRAAERHPEKYQGSSVSNLHVEPCGTNTHA SSLQHENSSLLLTKDRMNVEKAEFCNKSKQPGLARSQHNRWAGSKETCNDRRTPSTEKKVDLNADPLCER KEWNK
	SGPTRTRRLEQKLISEEDLAANDILDYKDDDDKV
Restriction Sites:	Sgfl-Mlul

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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 customport@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. <u>More info</u>

- **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:	<u>NP 009225</u>
RefSeq Size:	1065 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

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STATES BRCA1 (NM_007294) Human Mutant ORF Clone – RC402897
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MW:

Gene Summary:

39.1 kDa

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]

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