

# Product datasheet for TG314439

# BRCA2 Human shRNA Plasmid Kit (Locus ID 675)

## **Product data:**

#### **Product Type:** shRNA Plasmids **Product Name:** BRCA2 Human shRNA Plasmid Kit (Locus ID 675) Locus ID: 675 BRCC2; BROVCA2; FACD; FAD; FAD1; FANCD; FANCD1; GLM3; PNCA2; XRCC11 Synonyms: pGFP-V-RS (TR30007) Vector: E. coli Selection: Kanamycin Mammalian Cell Puromycin Selection: Format: **Retroviral plasmids Components:** BRCA2 - Human, 4 unique 29mer shRNA constructs in retroviral GFP vector(Gene ID = 675). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-V-RS Vector, TR30013, included for free. <u>NM 000059</u>, <u>NM 000059.1</u>, <u>NM 000059.2</u>, <u>NM 000059.3</u>, <u>BC026160</u>, <u>BC047568</u>, <u>NM 000059.4</u> RefSeq: **UniProt ID:** P51587 Inherited mutations in BRCA1 and this gene, BRCA2, confer increased lifetime risk of Summary: 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 13g12.3 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 20201 shRNA Design: These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact techsupport@origene.com. If you need a special design or shRNA sequence, please utilize our custom shRNA service.



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### **GRIGENE** BRCA2 Human shRNA Plasmid Kit (Locus ID 675) – TG314439

Performance Guaranteed: OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).

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