

Product datasheet for RC217860L4V

OriGene Technologies, Inc.

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RAD51C (NM_002876) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: RAD51C (NM_002876) Human Tagged ORF Clone Lentiviral Particle

Symbol: RAD51C

Synonyms: BROVCA3; FANCO; R51H3; RAD51L2

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_002876

ORF Size: 405 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC217860).

Sequence:
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.

RefSeg: NM 002876.2

 RefSeq Size:
 607 bp

 RefSeq ORF:
 408 bp

 Locus ID:
 5889

 UniProt ID:
 043502

 Cytogenetics:
 17q22

Protein Families: Druggable Genome

Protein Pathways: Homologous recombination





ORIGENE

MW: 14.7 kDa

Gene Summary:

This gene is a member of the RAD51 family. RAD51 family members are highly similar to bacterial RecA and Saccharomyces cerevisiae Rad51 and are known to be involved in the homologous recombination and repair of DNA. This protein can interact with other RAD51 paralogs and is reported to be important for Holliday junction resolution. Mutations in this gene are associated with Fanconi anemia-like syndrome. This gene is one of four localized to a region of chromosome 17q23 where amplification occurs frequently in breast tumors. Overexpression of the four genes during amplification has been observed and suggests a possible role in tumor progression. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2013]