

Product datasheet for RC217569L3V

OriGene Technologies, Inc.

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WDR4 (NM_033661) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: WDR4 (NM_033661) Human Tagged ORF Clone Lentiviral Particle

Symbol: WDR4

Synonyms: GAMOS6; hWH; MIGSB; TRM82; TRMT82; Wuho

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

 Tag:
 Myc-DDK

 ACCN:
 NM_033661

 ORF Size:
 1236 bp

ORF Nucleotide

OTI Disclaimer:

1230 50

Sequence:

The ORF insert of this clone is exactly the same as(RC217569).

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 033661.3

 RefSeq Size:
 1524 bp

 RefSeq ORF:
 1239 bp

 Locus ID:
 10785

 UniProt ID:
 P57081

 Cytogenetics:
 21q22.3

 Domains:
 WD40

 MW:
 45.3 kDa







Gene Summary:

This gene encodes a member of the WD repeat protein family. WD repeats are minimally conserved regions of approximately 40 amino acids typically bracketed by gly-his and trp-asp (GH-WD), which may facilitate formation of heterotrimeric or multiprotein complexes. Members of this family are involved in a variety of cellular processes, including cell cycle progression, signal transduction, apoptosis, and gene regulation. This gene is excluded as a candidate for a form of nonsyndromic deafness (DFNB10), but is still a candidate for other disorders mapped to 21q22.3 as well as for the development of Down syndrome phenotypes. Alternatively spliced transcript variants have been found for this gene. [provided by RefSeq, May 2012]