

Product datasheet for RC212746L4V

OriGene Technologies, Inc.

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WNT9B (NM_003396) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: WNT9B (NM_003396) Human Tagged ORF Clone Lentiviral Particle

Symbol: WNT9B

Synonyms: WNT14B; WNT15

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

ACCN: NM_003396 **ORF Size:** 1071 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC212746).

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 003396.1

RefSeq Size: 1464 bp
RefSeq ORF: 1074 bp
Locus ID: 7484

 UniProt ID:
 O14905

 Cytogenetics:
 17q21.32

Protein Families: Secreted Protein, Transmembrane



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Protein Pathways: Basal cell carcinoma, Hedgehog signaling pathway, Melanogenesis, Pathways in cancer, Wnt

signaling pathway

MW: 38.97 kDa

Gene Summary: The WNT gene family consists of structurally related genes that encode secreted signaling

proteins. These proteins have been implicated in oncogenesis and in several developmental processes, including regulation of cell fate and patterning during embryogenesis. This gene is a member of the WNT gene family. Study of its expression in the teratocarcinoma cell line NT2 suggests that it may be implicated in the early process of neuronal differentiation of NT2 cells induced by retinoic acid. This gene is clustered with WNT3, another family member, in the chromosome 17q21 region. Alternative splicing results in multiple transcript variants.

[provided by RefSeq, Feb 2016]