

Product datasheet for RC224281L2V

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Lunatic Fringe (LFNG) (NM_001040167) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Lunatic Fringe (LFNG) (NM 001040167) Human Tagged ORF Clone Lentiviral Particle

Symbol: Lunatic Fringe

Synonyms: SCDO3

Mammalian Cell None

Selection:

Vector:

pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_001040167

ORF Size: 1137 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC224281).

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 001040167.1

 RefSeq Size:
 2384 bp

 RefSeq ORF:
 1140 bp

 Locus ID:
 3955

 UniProt ID:
 Q8NES3

Cytogenetics: 7p22.3

Protein Families: Transmembrane

Protein Pathways: Notch signaling pathway





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MW: 41.77 kDa

Gene Summary:

This gene is a member of the glycosyltransferase 31 gene family. Members of this gene family, which also includes the MFNG (GeneID: 4242) and RFNG (GeneID: 5986) genes, encode evolutionarily conserved glycosyltransferases that act in the Notch signaling pathway to define boundaries during embryonic development. While their genomic structure is distinct from other glycosyltransferases, these proteins have a fucose-specific beta-1,3-N-acetylglucosaminyltransferase activity that leads to elongation of O-linked fucose residues on Notch, which alters Notch signaling. The protein encoded by this gene is predicted to be a single-pass type II Golgi membrane protein but it may also be secreted and proteolytically processed like the related proteins in mouse and Drosophila (PMID: 9187150). Mutations in this gene have been associated with autosomal recessive spondylocostal dysostosis 3. [provided by RefSeq, May 2018]