

Product datasheet for RC210118L3V

OriGene Technologies, Inc.

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Caveolin 3 (CAV3) (NM_001234) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Caveolin 3 (CAV3) (NM 001234) Human Tagged ORF Clone Lentiviral Particle

Symbol: Caveolin 3

Synonyms: LGMD1C; LQT9; MPDT; RMD2; VIP-21; VIP21

Mammalian Cell

Selection:

Puromycin

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

Tag: Myc-DDK
ACCN: NM 001234

ORF Size: 453 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC210118).

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 001234.3

 RefSeq Size:
 1339 bp

 RefSeq ORF:
 456 bp

 Locus ID:
 859

 UniProt ID:
 P56539

 Cytogenetics:
 3p25.3

Protein Families: Druggable Genome, Transmembrane

Protein Pathways: Focal adhesion





MW: 17.3 kDa

Gene Summary:

This gene encodes a caveolin family member, which functions as a component of the caveolae plasma membranes found in most cell types. Caveolin proteins are proposed to be scaffolding proteins for organizing and concentrating certain caveolin-interacting molecules. Mutations identified in this gene lead to interference with protein oligomerization or intracellular routing, disrupting caveolae formation and resulting in Limb-Girdle muscular dystrophy type-1C (LGMD-1C), hyperCKemia or rippling muscle disease (RMD). Alternative splicing has been identified for this locus, with inclusion or exclusion of a differentially spliced intron. In addition, transcripts utilize multiple polyA sites and contain two potential translation initiation sites. [provided by RefSeq, Jul 2008]