

Product datasheet for RC224268L3V

OriGene Technologies, Inc.

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Phospholipase C beta 3 (PLCB3) (NM_000932) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Phospholipase C beta 3 (PLCB3) (NM_000932) Human Tagged ORF Clone Lentiviral Particle

Symbol: PLCB3
Synonyms: SMDCD

Mammalian Cell

Puromycin

Selection:

Vector:

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

 Tag:
 Myc-DDK

 ACCN:
 NM_000932

 ORF Size:
 3702 bp

ORF Nucleotide

OTI Disclaimer:

3702 bp

Sequence:

The ORF insert of this clone is exactly the same as(RC224268).

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 000932.1, NP 000923.1

 RefSeq Size:
 5728 bp

 RefSeq ORF:
 3705 bp

 Locus ID:
 5331

 UniProt ID:
 Q01970

 Cytogenetics:
 11q13.1

Protein Families: Druggable Genome





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Protein Pathways: Alzheimer's disease, Calcium signaling pathway, Chemokine signaling pathway, Gap junction,

GnRH signaling pathway, Huntington's disease, Inositol phosphate metabolism, Long-term

depression, Long-term potentiation, Melanogenesis, Metabolic pathways,

Phosphatidylinositol signaling system, Vascular smooth muscle contraction, Wnt signaling

pathway

MW: 138.8 kDa

Gene Summary: This gene encodes a member of the phosphoinositide phospholipase C beta enzyme family

that catalyze the production of the secondary messengers diacylglycerol and inositol 1,4,5-triphosphate from phosphatidylinositol in G-protein-linked receptor-mediated signal

transduction. Alternative splicing results in multiple transcript variants.[provided by RefSeq,

May 2010]