

Product datasheet for RC212265L3V

OriGene Technologies, Inc.

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S1P (MBTPS1) (NM_003791) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: S1P (MBTPS1) (NM 003791) Human Tagged ORF Clone Lentiviral Particle

Symbol: S1P

Synonyms: PCSK8; S1P; SEDKF; SKI-1

Mammalian Cell

Selection:

Puromycin

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

 Tag:
 Myc-DDK

 ACCN:
 NM_003791

 ORF Size:
 3156 bp

ORF Nucleotide

OTI Disclaimer:

3130 bp

Sequence:

The ORF insert of this clone is exactly the same as(RC212265).

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 003791.2

 RefSeq Size:
 4354 bp

 RefSeq ORF:
 3159 bp

 Locus ID:
 8720

 UniProt ID:
 Q14703

Cytogenetics: 16q23.3-q24.1 Domains: Peptidase_S8

Protein Families: Druggable Genome, Protease, Transcription Factors, Transmembrane





MW: 117.75 kDa

Gene Summary:

This gene encodes a member of the subtilisin-like proprotein convertase family, which includes proteases that process protein and peptide precursors trafficking through regulated or constitutive branches of the secretory pathway. The encoded protein undergoes an initial autocatalytic processing event in the ER to generate a heterodimer which exits the ER and sorts to the cis/medial-Golgi where a second autocatalytic event takes place and the catalytic activity is acquired. It encodes a type 1 membrane bound protease which is ubiquitously expressed and regulates cholesterol or lipid homeostasis via cleavage of substrates at non-basic residues. Mutations in this gene may be associated with lysosomal dysfunction. [provided by RefSeq, Feb 2014]