

Product datasheet for SC203800

SPHK1 (NM_021972) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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3' UTR Clones
SPHK1 (NM_021972) Human 3' UTR Clone
pMirTarget (PS100062)
SPHK1
SPHK
NM_021972
299 bp
<pre>>SC203800 3'UTR clone of NM_021972 The sequence shown below is from the reference sequence of NM_021972. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CAGATGCCACCGCCAGAAGAGCCCTTATGACCCCTGGGCCGCGCTGTGCCTTAGTGTCTACTTGCAGGA CCCTTCCTCCTTCCTAGGGCTGCAGGGCCTGTCCACAGCTCCTGTGGGGGTGGAGGAGACTCCTCTGG AGAAGGGTGAGAAAGTGGAGGCTATGCTTTGGGGGGACAGGCCAGAATGAAGTCCTGGGTCAGGAGCCC AGCTGGCCCAGCTGCCTATGTAAGGCCTTCTAGTTTGTTCTGAGACCCCCACCACAGAACCAA ATCCAAATAAAGTGACATTCCA ACGCGTAAGCGGCCGCGCGCTCTCAGATTCGAAGAAATGACCGACC</pre>
Sgfl-Mlul
Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).
The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.
<u>NM 021972.4</u>



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Summary:	The protein encoded by this gene catalyzes the phosphorylation of sphingosine to form sphingosine-1-phosphate (S1P), a lipid mediator with both intra- and extracellular functions. Intracellularly, S1P regulates proliferation and survival, and extracellularly, it is a ligand for cell surface G protein-coupled receptors. This protein, and its product S1P, play a key role in TNF- alpha signaling and the NF-kappa-B activation pathway important in inflammatory, antiapoptotic, and immune processes. Phosphorylation of this protein alters its catalytic activity and promotes its translocation to the plasma membrane. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Sep 2017]
Locus ID:	8877
MW:	11.3

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