

Product datasheet for SC200049

PAFAH1B3 (NM 001145939) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: PAFAH1B3 (NM 001145939) Human 3' UTR Clone

Symbol: PAFAH1B3 **PAFAHG** Synonyms:

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

ACCN: NM 001145939

Insert Size: 73 bp

Insert Sequence: >SC200049 3'UTR clone of NM_001145939

The sequence shown below is from the reference sequence of NM_001145939. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGTGCTCCCCTGCTGGAGCCCGCACCCTAAGCATCCTGCCTCCCACAACATTAAACTCTCCTTCC

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: 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 Components:

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: NM 001145939.2



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Summary: This gene encodes an acetylhydrolase that catalyzes the removal of an acetyl group from the

glycerol backbone of platelet-activating factor. The encoded enzyme is a subunit of the platelet-activating factor acetylhydrolase isoform 1B complex, which consists of the catalytic beta and gamma subunits and the regulatory alpha subunit. This complex functions in brain development. A translocation between this gene on chromosome 19 and the CDC-like kinase 2 gene on chromosome 1 has been observed, and was associated with cognitive disability, ataxia, and atrophy of the brain. Alternatively spliced transcript variants have been described.

[provided by RefSeq, Mar 2009]

Locus ID: 5050 **MW:** 2.3