

## **Product datasheet for SC200045**

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## PAFAH1B3 (NM\_002573) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** PAFAH1B3 (NM\_002573) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: PAFAH1B3
Synonyms: PAFAHG

**ACCN:** NM\_002573

**Insert Size:** 73 bp

Insert Sequence: >SC200045 3'UTR clone of NM\_002573

The sequence shown below is from the reference sequence of NM\_002573. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGTGCTCCCTGCTGGAGCCCGCACCCTAAGCATCCTGCTGCCTTCCCACAACATTAAACTCTCCTTCC

**TCAG** 

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

 ${\tt CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG}$ 

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).

**Components:** 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.

**RefSeg:** NM 002573.4





## PAFAH1B3 (NM\_002573) Human 3' UTR Clone - SC200045

**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