

Product datasheet for SC209576

APH1A (NM 001077628) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: APH1A (NM_001077628) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: APH1A

Synonyms: 6530402N02Rik; APH-1; APH-1A; CGI-78

ACCN: NM_001077628

Insert Size: 756 bp

Insert Sequence: >SC209576 3'UTR clone of NM_001077628

The sequence shown below is from the reference sequence of NM_001077628. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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).



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APH1A (NM_001077628) Human 3' UTR Clone - SC209576

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.

RefSeq: <u>NM 001077628.3</u>

Summary: This gene encodes a component of the gamma secretase complex that cleaves integral

membrane proteins such as Notch receptors and beta-amyloid precursor protein. The gamma secretase complex contains this gene product, or the paralogous anterior pharynx defective 1 homolog B (APH1B), along with the presenilin, nicastrin, and presenilin enhancer-2 proteins. The precise function of this seven-transmembrane-domain protein is unknown though it is suspected of facilitating the association of nicastrin and presenilin in the gamma secretase complex as well as interacting with substrates of the gamma secretase complex prior to their proteolytic processing. Polymorphisms in a promoter region of this gene have been associated with an increased risk for developing sporadic Alzheimer's disease.

Alternative splicing results in multiple protein-coding and non-protein-coding transcript

variants. [provided by RefSeq, Aug 2011]

Locus ID: 51107

MW: 28.3