

Product datasheet for SC211664

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Acid Phosphatase (ACP1) (NM_007099) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Acid Phosphatase (ACP1) (NM_007099) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: ACP1

Synonyms: HAAP; LMW-PTP; LMWPTP

ACCN: NM_007099

Insert Size: 1009 bp

Insert Sequence: >SC211664 3'UTR clone of NM_007099

The sequence shown below is from the reference sequence of NM_007099. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAAATAAACTGGGGAGTTATAAAAATACAACTAGAGATATAAA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul





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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.

RefSeq: <u>NM 007099.4</u>

Summary: The product of this gene belongs to the phosphotyrosine protein phosphatase family of

proteins. It functions as an acid phosphatase and a protein tyrosine phosphatase by

hydrolyzing protein tyrosine phosphate to protein tyrosine and orthophosphate. This enzyme also hydrolyzes orthophosphoric monoesters to alcohol and orthophosphate. This gene is genetically polymorphic, and three common alleles segregating at the corresponding locus give rise to six phenotypes. Each allele appears to encode at least two electrophoretically different isozymes, Bf and Bs, which are produced in allele-specific ratios. Multiple

alternatively spliced transcript variants encoding distinct isoforms have been identified for

this gene. [provided by RefSeq, Aug 2008]

Locus ID: 52

MW: 38.3