

Product datasheet for SC201066

OriGene Technologies, Inc.

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Acid Phosphatase 2 (ACP2) (NM_001131064) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Acid Phosphatase 2 (ACP2) (NM_001131064) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: ACP2

Synonyms: acid phosphatase 2, lysosomal; Acp-2; LAP; OTTMUSP00000015308

ACCN: NM_001131064

Insert Size: 160 bp

Insert Sequence: >SC201066 3'UTR clone of NM_001131064

The sequence shown below is from the reference sequence of NM_001131064. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AGAGTGGCCAGCCCTTCCCTGGGGTGGTGAGGGGACAGCTCTGGCCGTAGGCCTGCTGATGCCAGGCTCCTTTTCCCGCTGTTTCCCCGCTTCGCTCTACAGCTGCTGAAGTTCCCGTTTGGGCCCATGTCCCCG

TTATGAGCAGCTGCAGAACGAG

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 001131064.1





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Summary:

The protein encoded by this gene belongs to the histidine acid phosphatase family, which hydrolyze orthophosphoric monoesters to alcohol and phosphate. This protein is localized to the lysosomal membrane, and is chemically and genetically distinct from the red cell acid phosphatase. Mice lacking this gene showed multiple defects, including bone structure alterations, lysosomal storage defects, and an increased tendency towards seizures. An enzymatically-inactive allele of this gene in mice showed severe growth retardation, hair-follicle abnormalities, and an ataxia-like phenotype. Alternatively spliced transcript variants have been found for this gene. A C-terminally extended isoform is also predicted to be produced by the use of an alternative in-frame translation termination codon via a stop codon readthrough mechanism. [provided by RefSeq, Oct 2017]

Locus ID: 53

MW: 5.5