

Product datasheet for SC334898

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ATP6V0B (NM_001294333) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ATP6V0B (NM_001294333) Human Untagged Clone

Tag: Tag Free Symbol: ATP6V0B

Synonyms: ATP6F; HATPL; VMA16

Mammalian Cell Neomycin

Selection:

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001294333, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

ACCN: NM 001294333

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).



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Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: NM 001294333.1, NP 001281262.1

RefSeq Size: 2068 bp
RefSeq ORF: 786 bp
Locus ID: 533
Cytogenetics: 1p34.1

Protein Families: Transmembrane

Protein Pathways: Epithelial cell signaling in Helicobacter pylori infection, Lysosome, Metabolic pathways,

Oxidative phosphorylation, Vibrio cholerae infection

Gene Summary: This gene encodes a portion of the V0 domain of vacuolar ATPase (V-ATPase), a multisubunit

enzyme that mediates acidification of eukaryotic intracellular organelles. Activity of this enzyme is necessary for such varied processes as protein sorting, zymogen activation, receptor-mediated endocytosis, and synaptic vesicle proton gradient generation. Alternative

splicing results in multiple transcript variants. [provided by RefSeq, Jun 2014]

Transcript Variant: This variant (3) contains an alternate segment in the 3' region, which results in a frameshift, compared to variant 1. The encoded isoform (3) has a longer and distinct C-terminus, compared to variant 1. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on

alignments.