

## **Product datasheet for TP305447**

#### OriGene Technologies, Inc.

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## ATP6V1B2 (NM\_001693) Human Recombinant Protein

**Product data:** 

**Product Type:** Recombinant Proteins

**Description:** Recombinant protein of human ATPase, H+ transporting, lysosomal 56/58kDa, V1 subunit B2

(ATP6V1B2), 20 µg

Species: Human
Expression Host: HEK293T

Expression cDNA Clone >RC205447 protein sequence

or AA Sequence: Red=Cloning site Green=Tags(s)

MALRAMRGIVNGAAPELPVPTGGPAVGAQEQALAVSRNYLSQPRLTYKTVSGVNGPLVILDHVKFPRYAE IVHLTLPDGTKRSGQVLEVSGSKAVVQVFEGTSGIDAKKTSCEFTGDILRTPVSEDMLGRVFNGSGKPID RGPVVLAEDFLDIMGQPINPQCRIYPEEMIRTGISAIDGMNSIARGQKIPIFSAAGLPHNEIAAQICRQA GLVKKSKDVVDYSEENFAIVFAAMGVNMETARFFKSDFEENGSMDNVCLFLNLANDPTIERIITPRLALT TAEFLAYQCEKHVLVILTDMSSYAEALREVSAAREEVPGRRGFPGYMYTDLATIYERAGRVGGRNGSITQ IPILTMPNDDITHPIPDLTGYITEGQIYVDRQLHNRQIYPPINVLPSLSRLMKSAIGEGMTRKDHADVSN QLYACYAIGKDVQAMKAVVGEEALTSDDLLYLEFLQKFERNFIAQGPYENRTVFETLDIGWQLLRIFPKE

MLKRIPQSTLSEFYPRDSAKH

**TRTRPLEQKLISEEDLAANDILDYKDDDDKV** 

Tag: C-Myc/DDK
Predicted MW: 56.3 kDa

Concentration: >0.05 µg/µL as determined by microplate BCA method

**Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining

**Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol

**Preparation:** Recombinant protein was captured through anti-DDK affinity column followed by

conventional chromatography steps.

**Note:** For testing in cell culture applications, please filter before use. Note that you may experience

some loss of protein during the filtration process.

Storage: Store at -80°C.





RefSeq ORF:

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Stability: Stable for 12 months from the date of receipt of the product under proper storage and

handling conditions. Avoid repeated freeze-thaw cycles.

RefSeq: NP 001684

Locus ID: 526

**UniProt ID:** P21281 RefSeg Size: 3054 Cytogenetics: 8p21.3

1533

Synonyms: ATP6B1B2; ATP6B2; DOOD; HO57; VATB; Vma2; VPP3; ZLS2

**Summary:** This gene encodes a component of vacuolar ATPase (V-ATPase), a multisubunit enzyme that

> mediates acidification of eukaryotic intracellular organelles. V-ATPase dependent organelle acidification is necessary for such intracellular processes as protein sorting, zymogen activation, receptor-mediated endocytosis, and synaptic vesicle proton gradient generation. V-ATPase is composed of a cytosolic V1 domain and a transmembrane V0 domain. The V1 domain consists of three A, three B, and two G subunits, as well as a C, D, E, F, and H subunit. The V1 domain contains the ATP catalytic site. The protein encoded by this gene is one of two V1 domain B subunit isoforms and is the only B isoform highly expressed in osteoclasts.

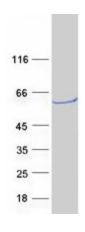
[provided by RefSeq, Jul 2008]

Druggable Genome **Protein Families:** 

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

phosphorylation, Vibrio cholerae infection

# **Product images:**



Coomassie blue staining of purified ATP6V1B2 protein (Cat# TP305447). The protein was produced from HEK293T cells transfected with ATP6V1B2 cDNA clone (Cat# [RC205447]) using

MegaTran 2.0 (Cat# [TT210002]).