

Product datasheet for TP315618

OriGene Technologies, Inc.

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ATP6V0E1 (NM_003945) Human Recombinant Protein

Product data:

Product Type: Recombinant Proteins

Description: Recombinant protein of human ATPase, H+ transporting, lysosomal 9kDa, V0 subunit e1

(ATP6V0E1), 20 µg

Species: Human
Expression Host: HEK293T

Expression cDNA Clone >RC215618 representing NM_003945

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

MAYHGLTVPLIVMSVFWGFVGFLVPWFIPKGPNRGVIITMLVTCSVCCYLFWLIAILAQLNPLFGPQLKN

ETIWYLKYHWP

TRTRPLEQKLISEEDLAANDILDYKDDDDK**V**

Tag: C-Myc/DDK

Predicted MW: 9.2 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.

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 003936

 Locus ID:
 8992

 UniProt ID:
 015342

RefSeq Size: 894



ATP6V0E1 (NM_003945) Human Recombinant Protein - TP315618

Cytogenetics: 5q35.1

RefSeq ORF: 243

Synonyms: ATP6H; ATP6V0E; M9.2; Vma21; Vma21p

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 and three B subunits, two G subunits plus the C, D, E, F, and H subunits. The V1 domain contains the ATP catalytic site. The V0 domain consists of five different subunits: a, c, c', c", and d. Additional isoforms of many of the V1 and V0 subunit proteins are encoded by multiple genes or alternatively spliced transcript variants. This encoded protein is possibly part of the V0 subunit. Since two nontranscribed pseudogenes have been found in dog, it is possible that the localization to chromosome 2 for this gene by radiation hybrid mapping is representing a pseudogene. Genomic mapping puts the

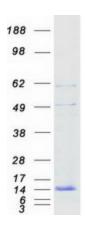
chromosomal location on 5g35.3. [provided by RefSeg, Jul 2008]

Protein Families: Transmembrane

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

phosphorylation, Vibrio cholerae infection

Product images:



Coomassie blue staining of purified ATP6V0E1 protein (Cat# TP315618). The protein was produced from HEK293T cells transfected with ATP6V0E1 cDNA clone (Cat# [RC215618]) using

MegaTran 2.0 (Cat# [TT210002]).