Cat. No. PME35253



PRODUCT INFORMATION

ATP6V1F **Target**

Synonyms ATP6S14; VATF; Vma7

Recombinant protein of human ATPase, H+ transporting, lysosomal 14kDa, V1 subunit F **Description**

(ATP6V1F)

Delivery 2-3 weeks **Uniprot ID** Q16864 **Expression Host** HEK293T Tag C-Myc/DDK

Molecular

N/A Characterization

Molecular Weight 13.2 kDa

> 80% as determined by SDS-PAGE and **Purity**

Coomassie blue staining

Formulation & Reconstitution

Background

25 mM Tris.HCl, pH 7.3, 100 mM glycine, 10%

glycerol

Storage & Shipping

Store at -80°C.

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 contains the ATT catalytic site. The voldomain 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 the V1 domain F subunit protein. [provided by RefSeq, Jul 2008]

Usage Research use only

Conjugate Unconjugated



