

## Product datasheet for PH303498

### ATP6V0D1 (NM\_004691) Human Mass Spec Standard

#### Product data:

Product Type:	Mass Spec Standards
Description:	ATP6V0D1 MS Standard C13 and N15-labeled recombinant protein (NP_004682)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC203498
Predicted MW:	40.3 kDa
Protein Sequence:	>RC203498 protein sequence Red=Cloning site Green=Tags(s)  MSFFPELYFNVDNGYLEGLVRGLKAGVLSQADYLNLVQCETLEDLKLHLQSTDYGNFLANEASPLTVSVI DDRLKEKMVVEFRMRNHAYEPLASFLDFITYSYMIDNVILLITGTLHQRSIAELVPKCHPLGSFEQMEA VNIAQTPAELYNAILVDTPLAAFFQDCISEQDLDEMNIIEIRNTLYKAYLESFYKFTLLGGTTADAMCP ILEFEADDRRAFIITINSFGTELSKEDRAKLFPHCGRLYPEGLAQLARADDYEQVKNVADYYPEYKLLFEG AGSNPGDKTLEDRLFHEVVKLNKLAFLNQFHGVFYAFVKLKEQECRNIVWIAECIAQRHRAKIDNYIPI F  TRTRPLEQKLISEEDLAANDILDYKDDDDKV
Tag:	C-Myc/DDK
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Concentration:	>0.05 µg/µL as determined by microplate BCA method
Labeling Method:	Labeled with [U- <sup>13</sup> C <sub>6</sub> , <sup>15</sup> N <sub>4</sub> ]-L-Arginine and [U- <sup>13</sup> C <sub>6</sub> , <sup>15</sup> N <sub>2</sub> ]-L-Lysine
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3
Storage:	Store at -80°C. Avoid repeated freeze-thaw cycles.
Stability:	Stable for 3 months from receipt of products under proper storage and handling conditions.
RefSeq:	<a href="#">NP_004682</a>
RefSeq Size:	1688
RefSeq ORF:	1053
Synonyms:	ATP6D; ATP6DV; P39; VATX; VMA6; VPATPD
Locus ID:	9114



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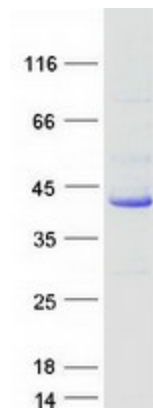
UniProt ID: [P61421](#)

Cytogenetics: 16q22.1

**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 known as the D subunit and is found ubiquitously. [provided by RefSeq, Jul 2008]

**Protein Pathways:** Epithelial cell signaling in Helicobacter pylori infection, Lysosome, Metabolic pathways, Oxidative phosphorylation, Vibrio cholerae infection

### Product images:



Coomassie blue staining of purified ATP6V0D1 protein (Cat# [TP303498]). The protein was produced from HEK293T cells transfected with ATP6V0D1 cDNA clone (Cat# [RC203498]) using MegaTran 2.0 (Cat# [TT210002]).