

## Product datasheet for RC215618

### ATP6V0E1 (NM\_003945) Human Tagged ORF Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** ATP6V0E1 (NM\_003945) Human Tagged ORF Clone  
**Tag:** Myc-DDK  
**Symbol:** ATP6V0E1  
**Synonyms:** ATP6H; ATP6V0E; M9.2; Vma21; Vma21p  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**ORF Nucleotide Sequence:** >RC215618 representing NM\_003945  
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTGTGAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCCCGCATCGCC

ATGGCGTATCACGGCCTCACTGTGCCTCTCATTGTGATGAGCGTGTCTGGGGCTTCGTCGGCTTCTTGG  
 TGCCTTGGTTCATCCCTAAGGGTCCTAACCGGGGAGTTATCATTACCATGTTGGTGACCTGTTCAAGTTG  
 CTGCTATCTCTTTGGCTGATTGCAATTCTGGCCCACTCAACCCTCTCTTGGACCGCAATTGAAAAAT  
 GAAACCATCTGGTATCTGAAGTATCATTGGCCT

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Protein Sequence:** >RC215618 representing NM\_003945  
 Red=Cloning site Green=Tags(s)  
 MAYHGLTVPLIVMSVFWGFVGLVPWFIPKGPNGVITMLVTCSVCCYLFWLIALAQLNPLFGPQLKN  
 ETIWYLYHWP

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

**Chromatograms:** [https://cdn.origene.com/chromatograms/mk6106\\_e06.zip](https://cdn.origene.com/chromatograms/mk6106_e06.zip)

**Restriction Sites:** SgfI-MluI


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**Cloning Scheme:**


**ACCN:** NM\_003945

**ORF Size:** 243 bp

**OTI Disclaimer:** The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**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.

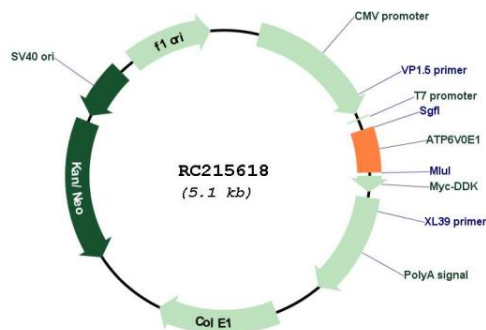
**Note:** Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.

**RefSeq:** [NM\\_003945.4](#)

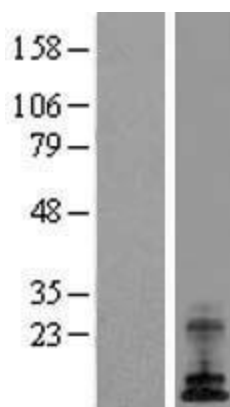
**RefSeq Size:** 894 bp

RefSeq ORF:	246 bp
Locus ID:	8992
UniProt ID:	<a href="#">O15342</a>
Cytogenetics:	5q35.1
Protein Families:	Transmembrane
Protein Pathways:	Epithelial cell signaling in Helicobacter pylori infection, Metabolic pathways, Oxidative phosphorylation, Vibrio cholerae infection
MW:	9.2 kDa
Gene Summary:	<p>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 5q35.3. [provided by RefSeq, Jul 2008]</p>

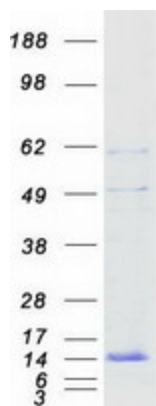
## Product images:



Circular map for RC215618



Western blot validation of overexpression lysate (Cat# [LY418332]) using anti-DDK antibody (Cat# [TA50011-100]). Left: Cell lysates from untransfected HEK293T cells; Right: Cell lysates from HEK293T cells transfected with RC215618 using transfection reagent MegaTran 2.0 (Cat# [TT210002]).



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]).