

Product datasheet for RN204104

Atp6v0e1 (NM_053578) Rat Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Atp6v0e1 (NM 053578) Rat Untagged Clone

Tag: Tag Free
Symbol: Atp6v0e1

Synonyms: Atp6k; Atp6v0e; dsr-1; Dsr-1A

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Cell Selection: Neomycin

Fully Sequenced ORF: >RN204104 representing NM_053578

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ATGGCATACCACGGCCTCACTGTGCCTTTGATCGTGATGAGCGTGTTCTGGGGCTTCGTGGGCCTCCTCGTGCCCTGGTTTATCCCCAAGGGTCCTAACCGGGGAGTTATCATCACCATGTTGGTGACCTGTTCCGTTTGCTGCTATCTCTTTTTGGCTGATTGCCGATTCTGCCGCAGCTCAATCCTCTGTTTGGACCACAGTTGAAAAAT

GAAACCATCTGGTATCTGAAGTATCACTGGCCT<mark>TGA</mark>

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM 053578

Insert Size: 246 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

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).



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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.

RefSeq: <u>NM 053578.3</u>, <u>NP 446030.3</u>

 RefSeq Size:
 835 bp

 RefSeq ORF:
 246 bp

 Locus ID:
 94170

 UniProt ID:
 Q794C0

 Cytogenetics:
 10q12

Gene Summary: Vacuolar ATPase is responsible for acidifying a variety of intracellular compartments in

eukaryotic cells.[UniProtKB/Swiss-Prot Function]