

Recombinant Human ATP6V0E1 293 Cell Lysate

Cat. No. ATP6V0E1-8586HCL **Lot. No.** (See product label)

SPECIFICATION

Product Overview

Species	Human
Source	HEK293
Description	Antigen standard for ATPase, H ⁺ transporting, lysosomal 9kDa, V0 subunit e1 (ATP6V0E1) is a lysate prepared from HEK293T cells transiently transfected with a TrueORF gene-carrying pCMV plasmid and then lysed in RIPA Buffer. Protein concentration was determined using a colorimetric assay. The antigen control carries a C-terminal Myc/DDK tag for detection.
Components	This product includes 3 vials: 1 vial of gene-specific cell lysate, 1 vial of control vector cell lysate, and 1 vial of loading buffer. Each lysate vial contains 0.1 mg lysate in 0.1 ml (1 mg/ml) of RIPA Buffer (50 mM Tris-HCl pH7.5, 250 mM NaCl, 5 mM EDTA, 50 mM NaF, 1% NP40). The loading buffer vial contains 0.5 ml 2X SDS Loading Buffer (125 mM Tris-Cl, pH6.8, 10% glycerol, 4% SDS, 0.002% Bromophenol blue, 5% beta-mercaptoethanol).
Size	0.1 mg
Storage Instruction	Store at -80°C. Minimize freeze-thaw cycles. After addition of 2X SDS Loading Buffer, the lysates can be stored at -20°C. Product is guaranteed 6 months from the date of shipment.

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Applications ELISA, WB, IP. WB: Mix equal volume of lysates with 2X SDS Loading Buffer. Boil the mixture for 10 min before loading (for membrane protein lysates, incubate the mixture at room temperature for 30 min). Load 5 ug lysate per lane.

GENE INFORMATION

Gene Name	ATP6V0E1?ATPase, H ⁺ transporting, lysosomal 9kDa, V0 subunit e1 [Homo sapiens?(human)]
Official Symbol	ATP6V0E1
Synonyms	ATP6V0E1; ATPase, H ⁺ transporting, lysosomal 9kDa, V0 subunit e1; M9.2; ATP6H; Vma21; Vma21p; ATP6V0E; V-type proton ATPase subunit e 1; V-ATPase H subunit; V-ATPase subunit e 1; V-ATPase M9.2 subunit; vacuolar proton pump H subunit; vacuolar ATP synthase subunit H; vacuolar proton pump subunit e 1; vacuolar proton-ATPase subunit M9.2; V-ATPase 9.2 kDa membrane accessory protein; H(+)-transporting two-sector ATPase, subunit H; NP_003936.1; EC 3.6.3.14
Gene ID	8992
mRNA Refseq	NM_003945
Protein Refseq	NP_003936
MIM	603931
UniProt ID	O15342
Chromosome Location	5q35.1

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Pathway	Collecting duct acid secretion, organism-specific biosystem; Collecting duct acid secretion, conserved biosystem; Epithelial cell signaling in Helicobacter pylori infection, conserved biosystem; Oxidative phosphorylation, organism-specific biosystem
Function	ATPase activity, coupled to transmembrane movement of ions; proton-transporting ATPase activity, rotational mechanism; transporter activity

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