

Recombinant Human ATP6V1F 293 Cell Lysate

Cat. No. ATP6V1F-8578HCL **Lot. No.** (See product label)

SPECIFICATION

Species	Human
Source	HEK293
Description	Antigen standard for ATPase, H ⁺ transporting, lysosomal 14kDa, V1 subunit F (ATP6V1F) 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.
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

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mixture at room temperature for 30 min). Load 5 ug lysate per lane.

GENE INFORMATION

Gene Name [ATP6V1F ATPase, H+ transporting, lysosomal 14kDa, V1 subunit F \[Homo sapiens \]](#)

Official Symbol [ATP6V1F](#)

Synonyms [ATP6V1F](#); [ATPase, H+ transporting, lysosomal 14kDa, V1 subunit F](#); [V-type proton ATPase subunit F](#); [ATP6S14](#); [VATF](#); [Vma7](#); [V-ATPase F subunit](#); [V-ATPase subunit F](#); [ATPase, vacuolar, 14 kD](#); [V-ATPase 14 kDa subunit](#); [vacuolar proton pump F subunit](#); [vacuolar proton pump subunit F](#); [vacuolar ATP synthase subunit F](#); [adenosinetriphosphatase 14k chain](#); [H\(+\)-transporting two-sector ATPase, 14kD subunit](#); [MGC117321](#); [MGC126037](#); [MGC126038](#);

Gene ID [9296](#)

mRNA Refseq [NM_001198909](#)

Protein Refseq [NP_001185838](#)

MIM [607160](#)

UniProt ID [Q16864](#)

Chromosome Location [7q32.1](#)

Pathway [Collecting duct acid secretion, organism-specific biosystem](#); [Collecting duct acid secretion, conserved biosystem](#); [Epithelial cell signaling in Helicobacter pylori infection, organism-specific biosystem](#); [Epithelial cell signaling in Helicobacter pylori](#)

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infection, conserved biosystem; Insulin receptor recycling, organism-specific biosystem; Iron uptake and transport, organism-specific biosystem; Metabolic pathways, organism-specific biosystem;

Function

ATPase activity, uncoupled; hydrogen ion transmembrane transporter activity; hydrogen ion transporting ATP synthase activity, rotational mechanism; protein binding; proton-transporting ATPase activity, rotational mechanism;

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