

## Recombinant Human ATP5I 293 Cell Lysate

**Cat. No.** ATP5I-8598HCL    **Lot. No.** (See product label)

### SPECIFICATION

<b>Species</b>	Human
<b>Source</b>	HEK293
<b>Description</b>	Antigen standard for ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 complex, subunit E (ATP5I), nuclear gene encoding mitochondrial protein 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.
<b>Components</b>	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).
<b>Size</b>	0.1 mg
<b>Storage Instruction</b>	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.
<b>Applications</b>	ELISA, WB, IP. WB: Mix equal volume of lysates with 2X SDS Loading Buffer. Boil

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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** [ATP5I ATP synthase, H<sup>+</sup> transporting, mitochondrial Fo complex, subunit E \[ Homo sapiens \]](#)

**Official Symbol** [ATP5I](#)

**Synonyms** [ATP5I](#); [ATP synthase, H<sup>+</sup> transporting, mitochondrial Fo complex, subunit E](#); [ATP synthase, H<sup>+</sup> transporting, mitochondrial F0 complex, subunit e](#) , [ATP synthase, H<sup>+</sup> transporting, mitochondrial F0 complex, subunit E](#); [ATP synthase subunit e, mitochondrial](#); [ATPase subunit e](#); [ATP synthase e chain, mitochondrial](#); [F1F0-ATP synthase, murine e subunit](#); [ATP synthase, H<sup>+</sup> transporting, mitochondrial F0 complex, subunit E](#); [ATP5K](#); [MGC12532](#);

**Gene ID** [521](#)

**mRNA Refseq** [NM\\_007100](#)

**Protein Refseq** [NP\\_009031](#)

**MIM** [601519](#)

**UniProt ID** [P56385](#)

**Chromosome Location** [4p16.3](#)

**Pathway** [Electron Transport Chain, organism-specific biosystem](#); [F-type ATPase, eukaryotes](#),

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organism-specific biosystem; Formation of ATP by chemiosmotic coupling, organism-specific biosystem; Metabolic pathways, organism-specific biosystem; Metabolism, organism-specific biosystem; Oxidative phosphorylation, organism-specific biosystem; Oxidative phosphorylation, organism-specific biosystem;

**Function**

contributes\_to ATPase activity; hydrogen ion transmembrane transporter activity; transmembrane transporter activity;

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