

Full-Length

ATP5E (Human) Recombinant Protein (P01)

Catalog # H00000514-P01 Siz

Size 25 ug, 10 ug

Applications

175 -83 -62 -47.5 -32.5 -

Specification	
Product Description	Human ATP5E full-length ORF (AAH01690, 1 a.a 51 a.a.) recombinant protein with GST-tag at N-t erminal.
Sequence	MVAYWRQAGLSYIRYSQICAKAVRDALKTEFKANAEKTSGSNVKIVKVKKE
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	31.35
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCI, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.
Storage Instruction	Store at -80°C. Aliquot to avoid repeated freezing and thawing.
Note	Best use within three months from the date of receipt of this protein.



Applications

- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — ATP5E	
Entrez GenelD	<u>514</u>
GeneBank Accession#	<u>BC001690</u>
Protein Accession#	AAH01690
Gene Name	ATP5E
Gene Alias	ATPE, MGC104243
Gene Description	ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit
Omim ID	<u>606153</u>
Gene Ontology	Hyperlink
Gene Summary	This gene encodes a subunit of mitochondrial ATP synthase. Mitochondrial ATP synthase catalyz es ATP synthesis, utilizing an electrochemical gradient of protons across the inner membrane duri ng oxidative phosphorylation. ATP synthase is composed of two linked multi-subunit complexes: t he soluble catalytic core, F1, and the membrane-spanning component, Fo, comprising the proton channel. The catalytic portion of mitochondrial ATP synthase consists of 5 different subunits (alph a, beta, gamma, delta, and epsilon) assembled with a stoichiometry of 3 alpha, 3 beta, and a sing le representative of the other 3. The proton channel consists of three main subunits (a, b, c). This g ene encodes the epsilon subunit of the catalytic core. Two pseudogenes of this gene are located on chromosomes 4 and 13. [provided by RefSeq
Other Designations	F(0)F(1)-ATPase H(+)-transporting two-sector ATPase OTTHUMP00000031404 OTTHUMP0000 0174442 OTTHUMP00000174443 mitochondrial ATP synthase epsilon chain mitochondrial ATP ase



Pathway

- Metabolic pathways
- Oxidative phosphorylation