

Product datasheet for PH310456

ATP5F1E (NM_006886) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards **Description:** ATP5E MS Standard C13 and N15-labeled recombinant protein (NP_008817) Species: Human **Expression Host: HEK293** RC210456 Expression cDNA Clone or AA Sequence: Predicted MW: 5.8 kDa >RC210456 protein sequence Protein Sequence: Red=Cloning site Green=Tags(s) MVAYWRQAGLSYIRYSQICAKAVRDALKTEFKANAEKTSGSNVKIVKVKKE TRTRPLEQKLISEEDLAANDILDYKDDDDKV Tag: C-Myc/DDK > 80% as determined by SDS-PAGE and Coomassie blue staining **Purity:** >0.05 µg/µL as determined by microplate BCA method **Concentration:** Labeling Method: Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine **Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3 Store at -80°C. Avoid repeated freeze-thaw cycles. Storage: Stable for 3 months from receipt of products under proper storage and handling conditions. Stability: NP 008817 RefSeq: **RefSeq Size:** 449 **RefSeq ORF:** 153 Synonyms: ATP5E; ATPE; MC5DN3 Locus ID: 514 **UniProt ID:** P56381 Cytogenetics: 20q13.32



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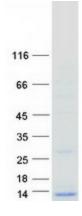
OriGene Technologies, Inc.

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Summary:	This gene encodes a subunit of mitochondrial ATP synthase. Mitochondrial ATP synthase catalyzes ATP synthesis, utilizing an electrochemical gradient of protons across the inner membrane during oxidative phosphorylation. ATP synthase is composed of two linked multi-subunit complexes: the 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 (alpha, beta, gamma, delta, and epsilon) assembled with a stoichiometry of 3 alpha, 3 beta, and a single representative of the other 3. The proton channel consists of three main subunits (a, b, c). This gene encodes the epsilon subunit of the catalytic core. Two pseudogenes of this gene are located on chromosomes 4 and 13. Read-through transcripts that include exons from this gene are expressed from the upstream gene SLMO2.[provided by RefSeq, Mar 2011]
Protein Pathway	Ve: Alzheimer's disease Huntington's disease Metabolic pathways. Ovidative phosphonylation

Protein Pathways:Alzheimer's disease, Huntington's disease, Metabolic pathways, Oxidative phosphorylation,
Parkinson's disease

Product images:



Coomassie blue staining of purified ATP5F1E protein (Cat# [TP310456]). The protein was produced from HEK293T cells transfected with ATP5F1E cDNA clone (Cat# [RC210456]) using MegaTran 2.0 (Cat# [TT210002]).

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