

Product datasheet for PH307908

OriGene Technologies, Inc.

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ATP5PD (NM_006356) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

Description: ATP5H MS Standard C13 and N15-labeled recombinant protein (NP_006347)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC207908

or AA Sequence: Predicted MW:

18.5 kDa

Protein Sequence: >RC207908 protein sequence

Red=Cloning site Green=Tags(s)

MAGRKLALKTIDWVAFAEIIPQNQKAIASSLKSWNETLTSRLAALPENPPAIDWAYYKANVAKAGLVDDF EKKFNALKVPVPEDKYTAQVDAEEKEDVKSCAEWVSLSKARIVEYEKEMEKMKNLIPFDQMTIEDLNEAF

PETKLDKKKYPYWPHQPIENL

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-Myc/DDK

Purity: > 80% as determined by SDS-PAGE and Coomassie blue staining

Concentration: $>0.05 \mu g/\mu L$ as determined by microplate BCA method

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

Storage: Store at -80°C. Avoid repeated freeze-thaw cycles.

Stability: Stable for 3 months from receipt of products under proper storage and handling conditions.

RefSeq: NP 006347

RefSeq Size: 628 RefSeq ORF: 483

Synonyms: APT5H; ATP5H; ATPQ

Locus ID: 10476

UniProt ID: <u>075947</u>, <u>A0PJH2</u>





Cytogenetics: 17q25.1

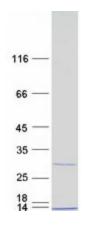
Summary: Mitochondrial ATP synthase catalyzes ATP synthesis, utilizing an electrochemical gradient of

protons across the inner membrane during oxidative phosphorylation. It is composed of two linked multi-subunit complexes: the soluble catalytic core, F1, and the membrane-spanning component, Fo, which comprises the proton channel. The F1 complex consists of 5 different subunits (alpha, beta, gamma, delta, and epsilon) assembled in a ratio of 3 alpha, 3 beta, and a single representative of the other 3. The Fo seems to have nine subunits (a, b, c, d, e, f, g, F6 and 8). This gene encodes the d subunit of the Fo complex. Alternatively spliced transcript variants encoding different isoforms have been identified for this gene. In addition, three pseudogenes are located on chromosomes 9, 12 and 15. [provided by RefSeq, Jun 2010]

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

Parkinson's disease

Product images:



Coomassie blue staining of purified ATP5PD protein (Cat# [TP307908]). The protein was produced from HEK293T cells transfected with ATP5PD cDNA clone (Cat# [RC207908]) using MegaTran 2.0 (Cat# [TT210002]).