

ATP5G1 Antibody (Center)

Purified Rabbit Polyclonal Antibody (Pab)
Catalog # AP22100c

Specification

ATP5G1 Antibody (Center) - Product Information

Application WB,E
Primary Accession P05496

Other Accession <u>P32876</u>, <u>Q9CR84</u>, <u>A1XQS5</u>, <u>Q06645</u>, <u>P17605</u>,

P07926, Q06055, P56383, Q5RAP9, Q06646, Q06056, Q3ZC75, P48201, P56384, Q5RFL2,

071S46

Reactivity Human, Rat

Predicted Bovine, Mouse, Pig, Sheep

Host Rabbit
Clonality polyclonal
Isotype Rabbit IgG
Calculated MW 14277

ATP5G1 Antibody (Center) - Additional Information

Gene ID 516

Other Names

ATP synthase F(0) complex subunit C1, mitochondrial, ATP synthase lipid-binding protein, ATP synthase proteolipid P1, ATP synthase proton-transporting mitochondrial F(0) complex subunit C1, ATPase protein 9, ATPase subunit c, ATP5G1

Target/Specificity

This ATP5G1 antibody is generated from a rabbit immunized with a KLH conjugated synthetic peptide between 41-71 amino acids from the Central region of human ATP5G1.

Dilution

WB~~1:2000

Format

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is purified through a protein A column, followed by peptide affinity purification.

Storage

Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

Precautions

ATP5G1 Antibody (Center) is for research use only and not for use in diagnostic or therapeutic procedures.

ATP5G1 Antibody (Center) - Protein Information



Name ATP5MC1 (HGNC:841)

Function Mitochondrial membrane ATP synthase (F(1)F(0) ATP synthase or Complex V) produces ATP from ADP in the presence of a proton gradient across the membrane which is generated by electron transport complexes of the respiratory chain. F-type ATPases consist of two structural domains, F(1) - containing the extramembraneous catalytic core and F(0) - containing the membrane proton channel, linked together by a central stalk and a peripheral stalk. During catalysis, ATP synthesis in the catalytic domain of F(1) is coupled via a rotary mechanism of the central stalk subunits to proton translocation. Part of the complex F(0) domain. A homomeric c-ring of probably 10 subunits is part of the complex rotary element.

Cellular Location

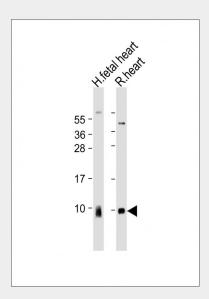
Mitochondrion membrane; Multi-pass membrane protein

ATP5G1 Antibody (Center) - Protocols

Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- <u>Immunoprecipitation</u>
- Flow Cytomety
- Cell Culture

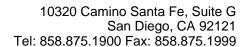
ATP5G1 Antibody (Center) - Images



All lanes: Anti-ATP5G1 Antibody (Center) at 1:2000 dilution Lane 1: human fetal heart lysate Lane 2: rat heart lysate Lysates/proteins at 20 μ g per lane. Secondary Goat Anti-Rabbit IgG, (H+L), Peroxidase conjugated at 1/10000 dilution. Predicted band size: 14 kDa Blocking/Dilution buffer: 5% NFDM/TBST.

ATP5G1 Antibody (Center) - Background

Mitochondrial membrane ATP synthase (F(1)F(0) ATP synthase or Complex V) produces ATP from





ADP in the presence of a proton gradient across the membrane which is generated by electron transport complexes of the respiratory chain. F-type ATPases consist of two structural domains, F(1) - containing the extramembraneous catalytic core and F(0) - containing the membrane proton channel, linked together by a central stalk and a peripheral stalk. During catalysis, ATP synthesis in the catalytic domain of F(1) is coupled via a rotary mechanism of the central stalk subunits to proton translocation. Part of the complex F(0) domain. A homomeric c-ring of probably 10 subunits is part of the complex rotary element.

ATP5G1 Antibody (Center) - References

Dyer M.R.,et al.Biochem. J. 293:51-64(1993). Higuti T.,et al.Biochim. Biophys. Acta 1173:87-90(1993). Wiemann S.,et al.Genome Res. 11:422-435(2001). Kalnine N.,et al.Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases. Farrell L.B.,et al.Biochem. Biophys. Res. Commun. 144:1257-1264(1987).