



# **ATP5I** Antibody

Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP53286

#### **Product Information**

Application WB
Primary Accession P56385
Reactivity Human
Host Rabbit
Clonality Polyclonal
Calculated MW 7933

## **Additional Information**

Gene ID 521

Other Names ATP synthase subunit e, mitochondrial, ATPase subunit e, ATP5I, ATP5K

**Target/Specificity** KLH-conjugated synthetic peptide encompassing a sequence within the center

region of human ATP5I. The exact sequence is proprietary.

**Dilution** WB~~ 1:1000

Format Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.09% (W/V)

sodium azide and 50% glycerol

**Storage** Store at -20 °C.Stable for 12 months from date of receipt

#### **Protein Information**

Name ATP5ME ( HGNC:846)

**Function** Subunit e, of the mitochondrial membrane ATP synthase complex (F(1)F(0)

ATP synthase or Complex V) that 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 (PubMed:37244256). ATP synthase complex consist of a soluble F(1) head domain - the catalytic core -

and a membrane F(1) domain - the membrane proton channel

(PubMed: <u>37244256</u>). These two domains are linked by a central stalk rotating inside the F(1) region and a stationary peripheral stalk (PubMed: <u>37244256</u>). 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

(Probable). In vivo, can only synthesize ATP although its ATP hydrolase activity can be activated artificially in vitro (By similarity). Part of the complex F(0)

domain (PubMed:<u>37244256</u>).

**Cellular Location** Mitochondrion. Mitochondrion inner membrane.

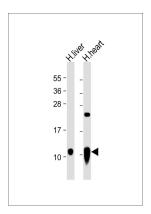
# **Background**

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. Minor subunit located with subunit a in the membrane.

## References

Fujiwara T.,et al.Submitted (NOV-1997) to the EMBL/GenBank/DDBJ databases. Kalnine N.,et al.Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases. Xu G.,et al.Proc. Natl. Acad. Sci. U.S.A. 106:19310-19315(2009). Burkard T.R.,et al.BMC Syst. Biol. 5:17-17(2011). Van Damme P.,et al.Proc. Natl. Acad. Sci. U.S.A. 109:12449-12454(2012).

# **Images**



All lanes: Anti-ATP5I Antibody at 1:1000 dilution Lane 1: human liver lysate Lane 2: human heart lysate Lysates/proteins at 20 µg per lane. Secondary Goat Anti-Rabbit IgG, (H+L),Peroxidase conjugated at 1/10000 dilution. Predicted band size: 8 kDa Blocking/Dilution buffer: 5% NFDM/TBST.

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