

# ATP5EP2 Antibody (Center)

Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP21779c

### **Product Information**

**Application** WB, E **Primary Accession** Q5VTU8 Reactivity Human Host Rabbit Clonality polyclonal Isotype Rabbit IgG **Clone Names** RB53889 **Calculated MW** 5807

## **Additional Information**

Other Names ATP synthase subunit epsilon-like protein, mitochondrial, ATP5EP2

**Target/Specificity** This ATP5EP2 antibody is generated from a rabbit immunized with a KLH

conjugated synthetic peptide between 19-52 amino acids from the Central

region of human ATP5EP2.

**Dilution** WB~~1:2000 E~~Use at an assay dependent concentration.

**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** ATP5EP2 Antibody (Center) is for research use only and not for use in

diagnostic or therapeutic procedures.

## **Protein Information**

Name ATP5F1EP2 ( HGNC:34026)

**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(1) domain and of the central stalk which is part of the complex rotary element. Rotation of the central stalk against the surrounding alpha(3)beta(3) subunits leads to hydrolysis of ATP in three separate catalytic sites on the beta subunits (By similarity).

**Cellular Location** 

Mitochondrion inner membrane {ECO:0000250 | UniProtKB:P56381}

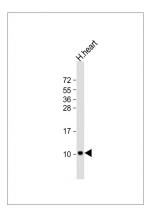
# **Background**

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(1) domain and of the central stalk which is part of the complex rotary element. Rotation of the central stalk against the surrounding alpha(3)beta(3) subunits leads to hydrolysis of ATP in three separate catalytic sites on the beta subunits (By similarity).

### References

Dunham A., et al. Nature 428:522-528(2004).

# **Images**



Anti-ATP5EP2 Antibody (Center) at 1:2000 dilution + 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: 6 kDa Blocking/Dilution buffer: 5% NFDM/TBST.

Please note: All products are 'FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES'.