# **ATP5I Polyclonal Antibody**

Catalog Number: E-AB-18027



*Note:* Centrifuge before opening to ensure complete recovery of vial contents.

## Description

Reactivity Human, Mouse, Rat

Immunogen Synthetic peptide of human ATP5I

Host Rabbit Isotype IgG

**Purification** Antigen affinity purification

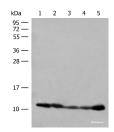
Conjugation Unconjugated

**Formulation** PBS with 0.05% NaN3 and 40% Glycerol,pH7.4

### **Applications** Recommended Dilution

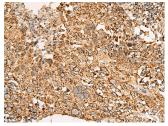
WB 1:500-1:2000 IHC 1:50-1:300 ELISA 1:5000-1:10000

#### Data

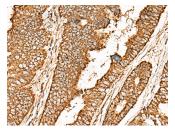


Western blot analysis of 293T cell PC-3 cell Human liver tissue lysates using ATP5I Polyclonal Antibody at dilution of 1:400

Observed Mw:Refer to figures Calculated Mw:8 kDa



Immunohistochemistry of paraffin-embedded Human cervical cancer tissue using ATP5I Polyclonal Antibody at dilution of 1:65(×200)



Immunohistochemistry of paraffin-embedded Human colorectal cancer tissue using ATP5I Polyclonal Antibody at dilution of 1:65(×200)

### **Preparation & Storage**

**Storage** Store at -20°C. Avoid freeze / thaw cycles.

## Background

#### For Research Use Only

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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 e subunit of the Fo complex. Alternative splicing results in multiple transcript variants. ATP5I (ATP Synthase, H+ Transporting, Mitochondrial Fo Complex Subunit E) is a Protein Coding gene. Among its related pathways are Respiratory electron transport, ATP synthesis by chemiosmotic coupling, and heat production by uncoupling proteins. and purine nucleotides de novo biosynthesis. GO annotations related to this gene include ATPase activity and hydrogen ion transmembrane transporter activity.

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