



Anti-ATP1B3 monoclonal antibody, clone 6H21 (CABT-21818MH)

This product is for research use only and is not intended for diagnostic use.

PRODUCT INFORMATION

Antigen Description

The protein encoded by this gene belongs to the family of Na+/K+ and H+/K+ ATPases beta chain proteins, and to the subfamily of Na+/K+ -ATPases. Na+/K+ -ATPase is an integral membrane protein responsible for establishing and maintaining the electrochemical gradients of Na and K ions across the plasma membrane. These gradients are essential for osmoregulation, for sodium-coupled transport of a variety of organic and inorganic molecules, and for electrical excitability of nerve and muscle. This enzyme is composed of two subunits, a large catalytic subunit (alpha) and a smaller glycoprotein subunit (beta). The beta subunit regulates, through assembly of alpha/beta heterodimers, the number of sodium pumps transported to the plasma membrane. The glycoprotein subunit of Na+/K+ -ATPase is encoded by multiple genes. This gene encodes a beta 3 subunit. A pseudogene exists for this gene, and it is located on chromosome 2.

Mouse monoclonal antibody raised against native ATP1B3.

Immunogen	Native purified from human cancer cel line.
Isotype	IgG2a
Source/Host	Mouse
Species Reactivity	Human
Clone	6H21
Conjugate	Unconjugated
Applications	IHC,IP,Flow Cyt
Format	Liquid
Buffer	In PBS (0.1% proclin, 2.0% Block Ace)

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GENE INFORMATION

ATPase, Na+/K+ transporting, beta 3 polypeptide; Na, K-ATPase beta-3 polypeptide; CD298; sodium/potassium-dependent ATPase beta-3 subunit; FLJ29027; sodium/potassium-transporting ATPase beta-3 chain; Sodium/potassium-dependent ATPase subunit beta-3; sodium/potassium-transporting ATPase subunit beta-3; ATPB-3; CD298 antigen; OTTHUMP00000214031 Entrez Gene ID 483 Protein Refseq NP_001670 UniProt ID P54709 Chromosome Location 3q23 Pathway Aldosterone-regulated sodium reabsorption, organism-specific biosystem; Aldosterone-regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, conserved biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	Gene Name	ATP1B3 ATPase, Na+/K+ transporting, beta 3 polypeptide [Homo sapiens]
sodium/potassium-dependent ATPase beta-3 subunit; FLJ29027; sodium/potassium-transporting ATPase beta-3 chain; Sodium/potassium-dependent ATPase subunit beta-3; sodium/potassium-transporting ATPase subunit beta-3; ATPB-3; CD298 antigen; OTTHUMP00000214031 Entrez Gene ID 483 Protein Refseq NP 001670 UniProt ID P54709 Chromosome Location 3q23 Pathway Aldosterone-regulated sodium reabsorption, organism-specific biosystem; Aldosterone-regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, conserved biosystem; Bile secretion, conserved biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	Official Symbol	ATP1B3
Protein Refseq NP_001670 UniProt ID P54709 Chromosome Location 3q23 Pathway Aldosterone-regulated sodium reabsorption, organism-specific biosystem; Aldosterone-regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, organism-specific biosystem; Bile secretion, conserved biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	Synonyms	sodium/potassium-dependent ATPase beta-3 subunit; FLJ29027; sodium/potassium-transporting ATPase beta-3 chain; Sodium/potassium-dependent ATPase subunit beta-3; sodium/potassium-transporting ATPase subunit beta-3; ATPB-3; CD298 antigen;
UniProt ID P54709 Chromosome Location 3q23 Aldosterone-regulated sodium reabsorption, organism-specific biosystem; Aldosterone-regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, organism-specific biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	Entrez Gene ID	483
Chromosome Location 3q23 Pathway Aldosterone-regulated sodium reabsorption, organism-specific biosystem; Aldosterone-regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, organism-specific biosystem; Bile secretion, conserved biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	Protein Refseq	<u>NP 001670</u>
Pathway Aldosterone-regulated sodium reabsorption, organism-specific biosystem; Aldosterone- regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, organism-specific biosystem; Bile secretion, conserved biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	UniProt ID	<u>P54709</u>
regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, organism-specific biosystem; Bile secretion, conserved biosystem; Calcium Regulation in the Cardiac Cell, organism-specific biosystem	Chromosome Location	3q23
Function protein binding; sodium:potassium-exchanging ATPase activity	Pathway	regulated sodium reabsorption, conserved biosystem; Basigin interactions, organism-specific biosystem; Bile secretion, organism-specific biosystem; Bile secretion, conserved biosystem;
	Function	protein binding; sodium:potassium-exchanging ATPase activity