

# ATP6AP1 (85.1): sc-81886



The Power to Question

## BACKGROUND

Vacuolar-type H<sup>+</sup>-ATPase (V-ATPase) is a multisubunit enzyme responsible for acidification of eukaryotic intracellular organelles. V-ATPases pump protons against an electrochemical gradient, thereby synthesizing ATP. A peripheral V<sub>1</sub> domain, which is responsible for ATP hydrolysis, and an integral V<sub>0</sub> domain, which is responsible for proton translocation, compose the V-ATPase. Nine subunits (A-H) make up the V<sub>1</sub> domain and five subunits (a, d, c, c' and c'') make up the V<sub>0</sub> domain. ATP6AP1 (ATPase, H<sup>+</sup> transporting, lysosomal accessory protein 1), also known as 16A, CF2, Ac45, XAP3, ATP6S1, VATPS1 (vacuolar ATP synthase S1 accessory protein) or ATP6IP1, is a type I transmembrane, V-ATPase accessory protein that is predominantly expressed in endocrine and neuronal cells. ATP6AP1 is responsible for targeting the V-ATPase enzyme to specialized complex vacuolar systems. Via its cytoplasmic tail, ATP6AP1 interacts with subunits of the V<sub>0</sub> domain. The disruption of this interaction in osteoclasts results in impaired bone resorption, suggesting an important role for ATP6AP1 in proper osteoclastic bone resorption.

## REFERENCES

1. Supek, F., et al. 1994. A novel accessory subunit for vacuolar H<sup>+</sup>-ATPase from chromaffin granules. *J. Biol. Chem.* 269: 24102-24106.
2. Getlawi, F., et al. 1996. Chromaffin granule membrane glycoprotein IV is identical with Ac45, a membrane-integral subunit of the granule's H<sup>+</sup>-ATPase. *Neurosci. Lett.* 219: 13-16.

## CHROMOSOMAL LOCATION

Genetic locus: ATP6AP1 (human) mapping to Xq28; Atp6ap1 (mouse) mapping to X A7.3.

## SOURCE

ATP6AP1 (85.1) is a mouse monoclonal antibody raised against recombinant ATP6AP1 of human origin.

## PRODUCT

Each vial contains 100 µg IgG<sub>1</sub> kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

## APPLICATIONS

ATP6AP1 (85.1) is recommended for detection of ATP6AP1 of mouse, rat and human origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000), immunoprecipitation [1-2 µg per 100-500 µg of total protein (1 ml of cell lysate)] and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

Suitable for use as control antibody for ATP6AP1 siRNA (h): sc-91265, ATP6AP1 siRNA (m): sc-141357, ATP6AP1 shRNA Plasmid (h): sc-91265-SH, ATP6AP1 shRNA Plasmid (m): sc-141357-SH, ATP6AP1 shRNA (h) Lentiviral Particles: sc-91265-V and ATP6AP1 shRNA (m) Lentiviral Particles: sc-141357-V.

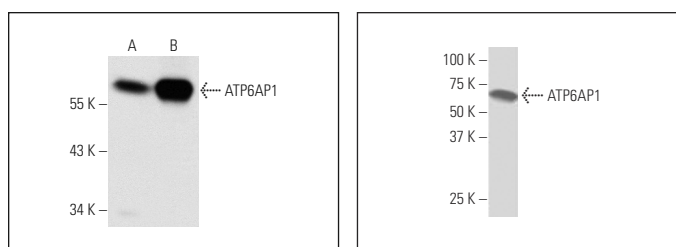
Molecular Weight of ATP6AP1: 45 kDa.

Positive Controls: HeLa whole cell lysate: sc-2200 or ATP6AP1 (h4): 293T Lysate: sc-175196.

## RECOMMENDED SUPPORT REAGENTS

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-IgGκ BP-HRP: sc-516102 or m-IgGκ BP-HRP (Cruz Marker): sc-516102-CM (dilution range: 1:1000-1:10000), Cruz Marker™ Molecular Weight Standards: sc-2035, UltraCruz® Blocking Reagent: sc-516214 and Western Blotting Luminol Reagent: sc-2048. 2) Immunoprecipitation: use Protein A/G PLUS-Agarose: sc-2003 (0.5 ml agarose/2.0 ml).

## DATA



ATP6AP1 (85.1): sc-81886. Western blot analysis of ATP6AP1 expression in non-transfected: sc-117752 (A) and human ATP6AP1 transfected: sc-175196 (B) 293T whole cell lysates.

ATP6AP1 (85.1): sc-81886. Western blot analysis of ATP6AP1 expression in HeLa whole cell lysate.

## SELECT PRODUCT CITATIONS

1. Anderson, K.S., et al. 2011. Protein microarray signature of autoantibody biomarkers for the early detection of breast cancer. *J. Proteome Res.* 10: 85-96.
2. Pareja, F., et al. 2018. Loss-of-function mutations in ATP6AP1 and ATP6AP2 in granular cell tumors. *Nat. Commun.* 9: 3533.
3. Fassl, A., et al. 2020. Increased lysosomal biomass is responsible for the resistance of triple-negative breast cancers to CDK4/6 inhibition. *Sci. Adv.* 6: eabb2210.
4. Perez-Canamas, A., et al. 2021. Fronto-temporal dementia risk gene TMEM106B has opposing effects in different lysosomal storage disorders. *Brain Commun.* 3: fcaa200.
5. Sun, X., et al. 2022. SARS-CoV-2 non-structural protein 6 triggers NLRP3-dependent pyroptosis by targeting ATP6AP1. *Cell Death Differ.* E-published.

## STORAGE

Store at 4° C, \*\*DO NOT FREEZE\*\*. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

## RESEARCH USE

For research use only, not for use in diagnostic procedures.

## PROTOCOLS

See our web site at [www.scbt.com](http://www.scbt.com) for detailed protocols and support products.