

ATG7 Antibody (Center)
Purified Rabbit Polyclonal Antibody (Pab)
Catalog # AP1813b

Specification

ATG7 Antibody (Center) - Product Information

Application	WB, IHC-P,E
Primary Accession	O95352
Other Accession	Q641Y5 , Q9D906 , Q5ZKY2
Reactivity	Human, Mouse
Predicted	Chicken, Rat
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Antigen Region	284-313

ATG7 Antibody (Center) - Additional Information

Gene ID 10533

Other Names

Ubiquitin-like modifier-activating enzyme ATG7, ATG12-activating enzyme E1 ATG7, Autophagy-related protein 7, APG7-like, hAGP7, Ubiquitin-activating enzyme E1-like protein, ATG7, APG7L

Target/Specificity

This ATG7 antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 284-313 amino acids from the Central region of human ATG7.

Dilution

WB~~1:1000
IHC-P~~1:50~100

Format

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is prepared by Saturated Ammonium Sulfate (SAS) precipitation followed by dialysis against PBS.

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

ATG7 Antibody (Center) is for research use only and not for use in diagnostic or therapeutic procedures.

ATG7 Antibody (Center) - Protein Information

Name ATG7 ([HGNC:16935](#))

Synonyms APG7L

Function E1-like activating enzyme involved in the 2 ubiquitin-like systems required for cytoplasm to vacuole transport (Cvt) and autophagy. Activates ATG12 for its conjugation with ATG5 as well as the ATG8 family proteins for their conjugation with phosphatidylethanolamine. Both systems are needed for the ATG8 association to Cvt vesicles and autophagosomal membranes. Required for autophagic death induced by caspase-8 inhibition. Facilitates LC3-I lipidation with phosphatidylethanolamine to form LC3-II which is found on autophagosomal membranes (PubMed:[34161705](#)). Required for mitophagy which contributes to regulate mitochondrial quantity and quality by eliminating the mitochondria to a basal level to fulfill cellular energy requirements and preventing excess ROS production. Modulates p53/TP53 activity to regulate cell cycle and survival during metabolic stress. Also plays a key role in the maintenance of axonal homeostasis, the prevention of axonal degeneration, the maintenance of hematopoietic stem cells, the formation of Paneth cell granules, as well as in adipose differentiation. Plays a role in regulating the liver clock and glucose metabolism by mediating the autophagic degradation of CRY1 (clock repressor) in a time-dependent manner (By similarity).

Cellular Location

Cytoplasm. Preautophagosomal structure. Note=Localizes also to discrete punctae along the ciliary axoneme and to the base of the ciliary axoneme

Tissue Location

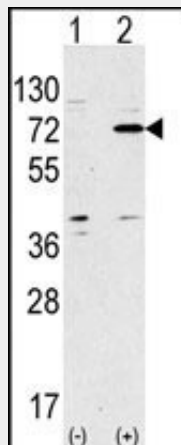
Widely expressed, especially in kidney, liver, lymph nodes and bone marrow.

ATG7 Antibody (Center) - Protocols

Provided below are standard protocols that you may find useful for product applications.

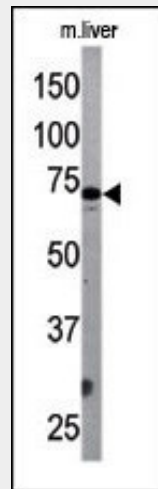
- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

ATG7 Antibody (Center) - Images

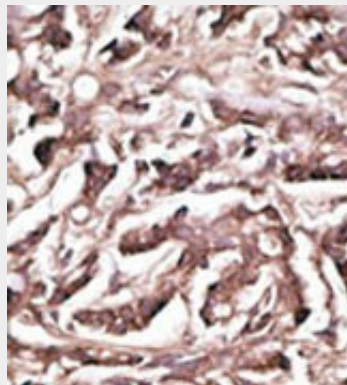


Western blot analysis of anti-ATG7 Antibody (Center) Pab (Cat. #AP1813b) in 293 cell line lysates

transiently transfected with the ATG7 gene (2ug/lane). hAPG7L-P299(arrow) was detected using the purified Pab.



The ATG7 Antibody (Center) Pab (Cat. #AP1813b) is used in Western blot to detect APG7L in mouse liver tissue lysate. APG7L (arrow) was detected using the purified Pab.



Formalin-fixed and paraffin-embedded human cancer tissue reacted with the primary antibody, which was peroxidase-conjugated to the secondary antibody, followed by DAB staining. This data demonstrates the use of this antibody for immunohistochemistry; clinical relevance has not been evaluated. BC = breast carcinoma; HC = hepatocarcinoma.

ATG7 Antibody (Center) - Background

Macroautophagy is the major inducible pathway for the general turnover of cytoplasmic constituents in eukaryotic cells, it is also responsible for the degradation of active cytoplasmic enzymes and organelles during nutrient starvation. Macroautophagy involves the formation of double-membrane bound autophagosomes which enclose the cytoplasmic constituent targeted for degradation in a membrane bound structure, which then fuse with the lysosome (or vacuole) releasing a single-membrane bound autophagic bodies which are then degraded within the lysosome (or vacuole). APG7 functions as an E1 enzyme essential for multisubstrates such as GABARAPL1 and ATG12. APG3L is an E2-like conjugating enzyme facilitating covalent binding of APG8 (MAP1LC3) to phosphatidylethanolamine (PE). APG7 (an E1-like enzyme) facilitates this reaction by forming an E1-E2 complex with APG3. Formation of the PE conjugate is essential for autophagy.

ATG7 Antibody (Center) - References

Baehrecke EH. *Nat Rev Mol Cell Biol.* 6(6):505-10. (2005) Lum JJ, et al. *Nat Rev Mol Cell Biol.* 6(6):439-48. (2005) Greenberg JT. *Dev Cell.* 8(6):799-801. (2005) Levine B. *Cell.* 120(2):159-62.

(2005) Shintani T and Klionsky DJ. Science. 306(5698):990-5. (2004) Tanida I., et al. Biochem. Biophys. Res. Commun. 292:256-262(2002) Tanida I., et al. J. Biol. Chem. 277:13739-13744(2002)

ATG7 Antibody (Center) - Citations

- [Autophagy deficiency by hepatic FIP200 deletion uncouples steatosis from liver injury in NAFLD.](#)
- [Temporal orchestration of circadian autophagy rhythm by C/EBP \$\beta\$.](#)