

**ATG4B Antibody (Center)**  
**Affinity Purified Rabbit Polyclonal Antibody (Pab)**  
**Catalog # AP20544c**

**Specification**

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**ATG4B Antibody (Center) - Product Information**

Application	WB,E
Primary Accession	<a href="#">O9Y4P1</a>
Other Accession	<a href="#">O8BGE6</a> , <a href="#">O6DG88</a> , <a href="#">O6PZ02</a> , <a href="#">O6PZ03</a> , <a href="#">A0A0G2QC33</a>
Reactivity	Human
Predicted	Bovine, Chicken, Zebrafish, Mouse, Rat
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	44294
Antigen Region	250-290

**ATG4B Antibody (Center) - Additional Information**

**Gene ID** 23192

**Other Names**

Cysteine protease ATG4B, 3422-, AUT-like 1 cysteine endopeptidase, Autophagin-1, Autophagy-related cysteine endopeptidase 1, Autophagy-related protein 4 homolog B, hAPG4B, ATG4B, APG4B, AUTL1, KIAA0943

**Target/Specificity**

This ATG4B antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 250-290 amino acids from the Central region of human ATG4B.

**Dilution**

WB~~1:1000

**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**

ATG4B Antibody (Center) is for research use only and not for use in diagnostic or therapeutic procedures.

**ATG4B Antibody (Center) - Protein Information**

**Name** ATG4B {ECO:0000303|PubMed:15187094, ECO:0000312|HGNC:HGNC:20790}

**Function** Cysteine protease that plays a key role in autophagy by mediating both proteolytic activation and delipidation of ATG8 family proteins (PubMed:[15169837](#), PubMed:[15187094](#), PubMed:[17347651](#), PubMed:[19322194](#), PubMed:[21177865](#), PubMed:[22302004](#), PubMed:[26378241](#), PubMed:[27527864](#), PubMed:[28633005](#), PubMed:[28821708](#), PubMed:[29232556](#), PubMed:[30076329](#), PubMed:[30443548](#), PubMed:[30661429](#)). Required for canonical autophagy (macroautophagy), non-canonical autophagy as well as for mitophagy (PubMed:[33773106](#), PubMed:[33909989](#)). The protease activity is required for proteolytic activation of ATG8 family proteins: cleaves the C-terminal amino acid of ATG8 proteins MAP1LC3A, MAP1LC3B, MAP1LC3C, GABARAPL1, GABARAPL2 and GABARAP, to reveal a C-terminal glycine (PubMed:[15169837](#), PubMed:[15187094](#), PubMed:[17347651](#), PubMed:[19322194](#), PubMed:[20818167](#), PubMed:[21177865](#), PubMed:[22302004](#), PubMed:[27527864](#), PubMed:[28287329](#), PubMed:[28633005](#), PubMed:[29458288](#), PubMed:[30661429](#)). Exposure of the glycine at the C-terminus is essential for ATG8 proteins conjugation to phosphatidylethanolamine (PE) and insertion to membranes, which is necessary for autophagy (PubMed:[15169837](#), PubMed:[15187094](#), PubMed:[17347651](#), PubMed:[19322194](#), PubMed:[21177865](#), PubMed:[22302004](#)). Protease activity is also required to counteract formation of high-molecular weight conjugates of ATG8 proteins (ATG8ylation): acts as a deubiquitinating-like enzyme that removes ATG8 conjugated to other proteins, such as ATG3 (PubMed:[31315929](#), PubMed:[33773106](#)). In addition to the protease activity, also mediates delipidation of ATG8 family proteins (PubMed:[15187094](#), PubMed:[19322194](#), PubMed:[28633005](#), PubMed:[29458288](#), PubMed:[32686895](#), PubMed:[33909989](#)). Catalyzes delipidation of PE- conjugated forms of ATG8 proteins during macroautophagy (PubMed:[15187094](#), PubMed:[19322194](#), PubMed:[29458288](#), PubMed:[32686895](#), PubMed:[33909989](#)). Also involved in non-canonical autophagy, a parallel pathway involving conjugation of ATG8 proteins to single membranes at endolysosomal compartments, by catalyzing delipidation of ATG8 proteins conjugated to phosphatidylserine (PS) (PubMed:[33909989](#)). Compared to other members of the family (ATG4A, ATG4C or ATG4D), constitutes the major protein for proteolytic activation of ATG8 proteins, while it displays weaker delipidation activity than other ATG4 paralogs (PubMed:[29458288](#), PubMed:[30661429](#)). Involved in phagophore growth during mitophagy independently of its protease activity and of ATG8 proteins: acts by regulating ATG9A trafficking to mitochondria and promoting phagophore-endoplasmic reticulum contacts during the lipid transfer phase of mitophagy (PubMed:[33773106](#)).

#### Cellular Location

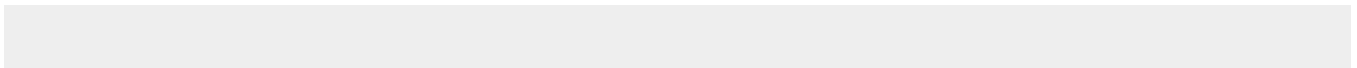
Cytoplasm. Cytoplasm, cytosol. Cytoplasmic vesicle, autophagosome. Endoplasmic reticulum. Mitochondrion. Note=Mainly localizes to the cytoplasm, including cytosol (PubMed:[29165041](#)). A samll potion localizes to mitochondria; phosphorylation at Ser-34 promotes localization to mitochondria (PubMed:[29165041](#)).

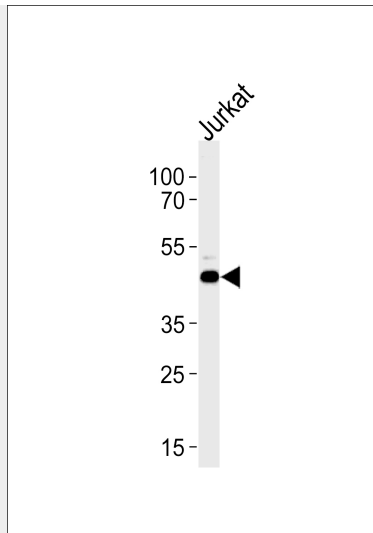
#### ATG4B Antibody (Center) - Protocols

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](#)

#### ATG4B Antibody (Center) - Images





ATG4B Antibody (Center) (Cat. #AP20544c) western blot analysis in Jurkat cell line lysates (35ug/lane). This demonstrates the ATG4B antibody detected the ATG4B protein (arrow).

#### **ATG4B Antibody (Center) - Background**

Cysteine protease required for autophagy, which cleaves the C-terminal part of either MAP1LC3, GABARAPL2 or GABARAP, allowing the liberation of form I. A subpopulation of form I is subsequently converted to a smaller form (form II). Form II, with a revealed C-terminal glycine, is considered to be the phosphatidylethanolamine (PE)-conjugated form, and has the capacity for the binding to autophagosomes.

#### **ATG4B Antibody (Center) - References**

- Marino G., et al. J. Biol. Chem. 278:3671-3678(2003).
- Kabeya Y., et al. J. Cell Sci. 117:2805-2812(2004).
- Nagase T., et al. DNA Res. 6:63-70(1999).
- Ohara O., et al. Submitted (AUG-2005) to the EMBL/GenBank/DDBJ databases.
- Ota T., et al. Nat. Genet. 36:40-45(2004).