

# ATG4A Rabbit mAb

Catalog # AP75120

## Product Information

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<b>Application</b>	WB, IHC-P
<b>Primary Accession</b>	<a href="#">Q8WYN0</a>
<b>Reactivity</b>	Rat, Human, Mouse
<b>Host</b>	Rabbit
<b>Clonality</b>	Monoclonal Antibody
<b>Isotype</b>	IgG
<b>Conjugate</b>	Unconjugated
<b>Purification</b>	Affinity Purified
<b>Calculated MW</b>	45378

## Additional Information

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<b>Gene ID</b>	115201
<b>Other Names</b>	ATG4A
<b>Dilution</b>	WB~~1/500-1/1000 IHC-P~~N/A
<b>Format</b>	Liquid in 50mM Tris-Glycine(pH 7.4), 0.15M NaCl, 40%Glycerol, 0.01% sodium azide and 0.05% BSA.
<b>Storage</b>	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.

## Protein Information

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<b>Name</b>	ATG4A {ECO:0000303   Ref.20, ECO:0000312   HGNC:HGNC:16489}
<b>Function</b>	Cysteine protease that plays a key role in autophagy by mediating both proteolytic activation and delipidation of ATG8 family proteins (PubMed: <a href="#">12473658</a> , PubMed: <a href="#">15169837</a> , PubMed: <a href="#">17347651</a> , PubMed: <a href="#">21177865</a> , PubMed: <a href="#">21245471</a> , PubMed: <a href="#">22302004</a> , PubMed: <a href="#">32732290</a> ). The protease activity is required for proteolytic activation of ATG8 family proteins: cleaves the C-terminal amino acid of ATG8 proteins to reveal a C-terminal glycine (PubMed: <a href="#">12473658</a> , PubMed: <a href="#">15169837</a> , PubMed: <a href="#">17347651</a> , PubMed: <a href="#">21177865</a> , PubMed: <a href="#">21245471</a> , PubMed: <a href="#">22302004</a> ). 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: <a href="#">12473658</a> , PubMed: <a href="#">15169837</a> , PubMed: <a href="#">17347651</a> , PubMed: <a href="#">21177865</a> , PubMed: <a href="#">21245471</a> , PubMed: <a href="#">22302004</a> ). Preferred substrate is GABARAPL2 followed by MAP1LC3A and GABARAP (PubMed: <a href="#">12473658</a> , PubMed: <a href="#">15169837</a> , PubMed: <a href="#">17347651</a> , PubMed: <a href="#">21177865</a> , PubMed: <a href="#">21245471</a> ,

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:[29458288](#), PubMed:[33909989](#)). Catalyzes delipidation of PE-conjugated forms of ATG8 proteins during macroautophagy (PubMed:[29458288](#), PubMed:[33909989](#)). Compared to ATG4B, the major protein for proteolytic activation of ATG8 proteins, shows weaker ability to cleave the C-terminal amino acid of ATG8 proteins, while it displays stronger delipidation activity (PubMed:[29458288](#)). 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 {ECO:0000250|UniProtKB:Q8BGE6}.

**Background**

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