

# Anti-ATG16L1 Antibody

Rabbit polyclonal antibody to ATG16L1

Catalog # AP60427

## Product Information

Application	WB
Primary Accession	<a href="#">Q676U5</a>
Other Accession	<a href="#">Q8CQJ2</a>
Reactivity	Human, Rat
Host	Rabbit
Clonality	Polyclonal
Calculated MW	68265

## Additional Information

Gene ID	55054
Other Names	APG16L; Autophagy-related protein 16-1; APG16-like 1
Target/Specificity	Recognizes endogenous levels of ATG16L1 protein.
Dilution	WB~~WB (1/500 - 1/1000)
Format	Liquid in 0.42% Potassium phosphate, 0.87% Sodium chloride, pH 7.3, 30% glycerol, and 0.09% (W/V) sodium azide.
Storage	Store at -20 °C.Stable for 12 months from date of receipt

## Protein Information

Name	ATG16L1 {ECO:0000303 PubMed:17200669, ECO:0000312 HGNC:HGNC:21498}
Function	Plays an essential role in both canonical and non-canonical autophagy: interacts with ATG12-ATG5 to mediate the lipidation to ATG8 family proteins (MAP1LC3A, MAP1LC3B, MAP1LC3C, GABARAPL1, GABARAPL2 and GABARAP) (PubMed: <a href="#">23376921</a> , PubMed: <a href="#">23392225</a> , PubMed: <a href="#">24553140</a> , PubMed: <a href="#">24954904</a> , PubMed: <a href="#">27273576</a> , PubMed: <a href="#">29317426</a> , PubMed: <a href="#">30778222</a> , PubMed: <a href="#">33909989</a> ). Acts as a molecular hub, coordinating autophagy pathways via distinct domains that support either canonical or non- canonical signaling (PubMed: <a href="#">29317426</a> , PubMed: <a href="#">30778222</a> ). During canonical autophagy, interacts with ATG12-ATG5 to mediate the conjugation of phosphatidylethanolamine (PE) to ATG8 proteins, to produce a membrane-bound activated form of ATG8 (PubMed: <a href="#">23376921</a> , PubMed: <a href="#">23392225</a> , PubMed: <a href="#">24553140</a> , PubMed: <a href="#">24954904</a> , PubMed: <a href="#">27273576</a> ). Thereby, controls the elongation of the nascent autophagosomal membrane (PubMed: <a href="#">23376921</a> , PubMed: <a href="#">23392225</a> ,

PubMed:[24553140](#), PubMed:[24954904](#), PubMed:[27273576](#)). As part of the ATG8 conjugation system with ATG5 and ATG12, required for recruitment of LRRK2 to stressed lysosomes and induction of LRRK2 kinase activity in response to lysosomal stress (By similarity). Also involved in non-canonical autophagy, a parallel pathway involving conjugation of ATG8 proteins to single membranes at endolysosomal compartments, probably by catalyzing conjugation of phosphatidylserine (PS) to ATG8 (PubMed:[33909989](#)). Non-canonical autophagy plays a key role in epithelial cells to limit lethal infection by influenza A (IAV) virus (By similarity). Regulates mitochondrial antiviral signaling (MAVS)-dependent type I interferon (IFN-I) production (PubMed:[22749352](#), PubMed:[25645662](#)). Negatively regulates NOD1- and NOD2-driven inflammatory cytokine response (PubMed:[24238340](#)). Instead, promotes an autophagy-dependent antibacterial pathway together with NOD1 or NOD2 (PubMed:[20637199](#)). Plays a role in regulating morphology and function of Paneth cell (PubMed:[18849966](#)).

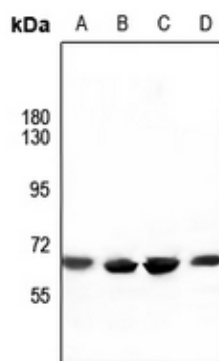
## Cellular Location

Cytoplasm. Preautophagosomal structure membrane; Peripheral membrane protein. Endosome membrane; Peripheral membrane protein. Lysosome membrane; Peripheral membrane protein. Note=Recruited to omegasomes membranes by WIPI2 (By similarity). Omegasomes are endoplasmic reticulum connected structures at the origin of preautophagosomal structures (By similarity). Localized to preautophagosomal structure (PAS) where it is involved in the membrane targeting of ATG5 (By similarity). Also localizes to discrete punctae along the ciliary axoneme (By similarity). Upon activation of non-canonical autophagy, recruited to single-membrane endolysosomal compartments (PubMed:29317426). Under starved conditions, the ATG12-ATG5-ATG16L1 complex is translocated to phagophores driven by RAB33B (PubMed:32960676). {ECO:0000250|UniProtKB:Q8C0J2, ECO:0000269|PubMed:29317426, ECO:0000269|PubMed:32960676}

## Background

KLH-conjugated synthetic peptide encompassing a sequence within the center region of human ATG16L1. The exact sequence is proprietary.

## Images



Western blot analysis of ATG16L1 expression in HepG2 (A), HCT116 (B), PC3 (C), A549 (D) whole cell lysates.

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