

# Atg7(Apg7) Antibody

Rabbit mAb Catalog # AP91480

## **Product Information**

Application	WB, IF, ICC
Primary Accession	<u>095352</u>
Reactivity	Rat, Human, Mouse
Clonality	Monoclonal
Other Names	hAGP7; Ubiquitin-activating enzyme E1-like protein; APG7L;
lsotype	Rabbit IgG
Host	Rabbit
Calculated MW	77960

### **Additional Information**

Dilution Purification Immunogen	WB 1:1000~1:5000 ICC/IF 1:50~1:200 Affinity-chromatography A synthesized peptide derived from human Atg7(Apg7)
Description	Formation of the autophagosome involves a ubiquitin-like conjugation system in which Atg12 is covalently bound to Atg5 and targeted to autophagosome vesicles. This conjugation reaction is mediated by the ubiquitin E1-like enzyme Atg7 and the E2-like enzyme Atg10.
Storage Condition and Buffer	5 , 5

#### **Protein Information**

Name	ATG7 ( <u>HGNC:16935</u> )
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 autophagosomes 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: <u>34161705</u> ). 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=Also localizes 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.

## Images



Western blot analysis of Atg7(Apg7) expression in (1) HepG2 cell lysate; (2) Mouse spleen lysate; (3) Rat kidney lysate.

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