

Recombinant Mouse A16L1 Protein

For Research Use Only

1. Basic Product Information

Catalog No.: REP51752

Size: 10ug, 100ug

Protein Name: Autophagy-related protein 16-1 (APG16-like 1)

Synonym: Atg16l1, Apg16l

UniProt ID: Q8C0J2

UniProt Link: www.uniprot.org/uniprotkb/Q8C0J2/entry

Species Source: Mouse

Expression System: HEK293T eukaryotic cell expression system Sequence

Length: 607 AA

Molecular Weight: 68 KD

Protein Tag: C-terminal His6

2. Amino Acid Sequence

MSSGLRAADFPRWKRHIAEELRRRDRLQRQAFEEIILQYTKLLEKSDLHSVLTQKLQAEKHDMPNRH
EISPGHDGAWNDSQLQEMAQLRIKHQEELTELHKKRGELAQLVIDLNNQMQQKDKEIQMNEAKISEY
LQTISDLETNCLDLRTKLQDLEVANQTLKDEYDALQITFTALEEKLKRTTEENQELVTRWMAEKAQEA
NRLNAENEEKDSRRRQARLQKELAEAAKEPLPVEQDDIEVIVDETSDHTEETSPVRAVSRAATKRLS
QPAGLLDSITNIFGRRSVSSIPVPQDIMDTHPASGKDVRVPTTASYVFD AHDGEVNAVQFSPGSRL
LATGGMDRRVKLWEAFGDKCEF KGSLSGSNAGITSIEFDSAGAYLLAASNDFASRIWTVDDYRLRHT
LTGHSGKVLSAKFLLDNARIVSGSHDRTLKLWDLRSKVCIKTVFAGSSCNDIVCTEQCVMSGHFDKKI
RFWDIRSESVVREMELLGKITALDLPTELLSCSRD DLLKVIDLRTNAV KQTFSAPGFKCGSDWT
RVVFSPDGSYVAAGSAEGSLYVWSVLTGKVEKVL SKQHSSINAVAWAPSGLHVVSVDKGSRAVL
WAQP

3. Buffer Composition

Liquid format: PBS-based buffer containing 50% glycerol. Glycerol concentration can be customized upon customer request; please state special requirements when placing your order.

Lyophilized format: Buffer before lyophilization is PBS supplemented with 6% trehalose.

4. Storage Conditions

Upon receipt, store the product at -20 °C or -80 °C. Aliquot the product for on-demand use to preserve protein activity and prevent denaturation and inactivation caused by repeated freeze-thaw cycles.

5. Product Description

This product is recombinant Autophagy-related protein 16-1 (APG16-like 1) produced using the HEK293T eukaryotic cell expression system. It is suitable for in-vitro functional assays, protein-protein interaction studies, antibody preparation and other biomedical research applications.

Protein 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:12665549, PubMed:18849966, PubMed:19898471, PubMed:23392225, PubMed:24553140, PubMed:24954904, PubMed:33586810). Acts as a molecular hub, coordinating autophagy pathways via distinct domains that support either canonical or non-canonical signaling (PubMed:33586810). 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 (By similarity). Thereby, controls the elongation of the nascent autophagosomal membrane (By similarity). 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 (PubMed:38227290). 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 (By similarity). Non-canonical autophagy plays a key role in epithelial cells to limit lethal infection by influenza A (IAV) virus (PubMed:33586810). Regulates mitochondrial antiviral signaling (MAVS)-dependent type I interferon (IFN-I) production (By similarity). Negatively regulates NOD1- and NOD2-driven inflammatory cytokine response (PubMed:24238340). Instead, promotes an autophagy-dependent antibacterial pathway together with NOD1 or NOD2 (PubMed:19898471, PubMed:19966812, PubMed:24238340). Plays a role in regulating morphology and function of Paneth cell (By similarity).

Subcellular Localization: Cytoplasm . Preautophagosomal structure membrane ; Peripheral membrane protein . Endosome membrane ; Peripheral membrane protein . Lysosome membrane ; Peripheral membrane protein . Note=Recruited to omegasomes membranes by WIPI2 (PubMed:24954904). Omegasomes are endoplasmic reticulum connected structures at the origin of preautophagosomal structures. Localized to preautophagosomal structure (PAS) where it is involved in the membrane targeting of ATG5 (PubMed:24954904). Localizes also to discrete punctae along the ciliary axoneme (PubMed:24954904). Upon activation of non-canonical autophagy, recruited to single-membrane endolysosomal compartments (By similarity). Under starved conditions, the ATG12-ATG5-ATG16L1 complex is translocated to phagophores driven by RAB33B (By similarity).

6. Notes

- Avoid repeated freeze-thaw cycles to maintain protein stability and biological activity.
- Please specify special buffer-composition requirements at the time of order placement.
- For research use only. Not intended for clinical diagnosis, therapeutic use or commercial manufacturing.