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ARL8B Protein (AA 1-186) (Strep Tag)



Image



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Quantity:	1 mg
Target:	ARL8B
Protein Characteristics:	AA 1-186
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	This ARL8B protein is labelled with Strep Tag.
Application:	ELISA, Western Blotting (WB), SDS-PAGE (SDS)

Product Details

Sequence:

MLALISRLLD WFRSLFWKEE MELTLVGLQY SGKTTFVNVI ASGQFSEDMI PTVGFNMRKV TKGNVTIKIW DIGGQPRFRS MWERYCRGVN AIVYMIDAAD REKIEASRNE LHNLLDKPQL QGIPVLVLGN KRDLPNALDE KQLIEKMNLS AIQDREICCY SISCKEKDNI DITLQWLIQH SKSRRS

Sequence without tag. The proposed Strep-Tag is based on experience s with the expression system, a different complexity of the protein could make another tag necessary. In case you have a special request, please contact us.

Characteristics:

Key Benefits:

- Made in Germany from design to production by highly experienced protein experts.
- Protein expressed with ALiCE® and purified by multi-step, protein-specific process to ensure correct folding and modification.
- These proteins are normally active (enzymatically functional) as our customers have reported (not tested by us and not guaranteed).

• State-of-the-art algorithm used for plasmid design (Gene synthesis).

This protein is a **made-to-order protein** and will be made for the first time for your order. Our experts in the lab will ensure that you receive a correctly folded protein.

The big advantage of ordering our **made-to-order proteins** in comparison to ordering custom made proteins from other companies is that there is no financial obligation in case the protein cannot be expressed or purified.

Expression System:

- ALiCE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from Nicotiana tabacum c.v.. This contains all the protein expression machinery needed to produce even the most difficult-to-express proteins, including those that require posttranslational modifications.
- During lysate production, the cell wall and other cellular components that are not required for
 protein production are removed, leaving only the protein production machinery and the
 mitochondria to drive the reaction. During our lysate completion steps, the additional
 components needed for protein production (amino acids, cofactors, etc.) are added to
 produce something that functions like a cell, but without the constraints of a living system all that's needed is the DNA that codes for the desired protein!

Concentration:

- The concentration of our recombinant proteins is measured using the absorbance at 280nm.
- The protein's absorbance will be measured in several dilutions and is measured against its specific reference buffer.
- We use the Expasy's ProtParam tool to determine the absorption coefficient of each protein.

Purification:	Two step purification of proteins expressed in Almost Living Cell-Free Expression System		
	(ALiCE®):		
	1. In a first purification step, the protein is purified from the cleared cell lysate using StrepTag capture material. Eluate fractions are analyzed by SDS-PAGE.		
	Protein containing fractions of the best purification are subjected to second purification step through size exclusion chromatography. Eluate fractions are analyzed by SDS-PAGE and Western blot.		
Purity:	>80 % as determined by SDS PAGE, Size Exclusion Chromatography and Western Blot.		
Endotoxin Level:	Low Endotoxin less than 1 EU/mg (< 0.1 ng/mg)		
Grade:	Crystallography grade		

Target Details

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Target:	ARL8B
Alternative Name:	ARL8B (ARL8B Products)
Background:	ADP-ribosylation factor-like protein 8B (EC 3.6.5.2) (ADP-ribosylation factor-like protein 10C)
	(Novel small G protein indispensable for equal chromosome segregation 1),FUNCTION: Small
	GTPase which cycles between active GTP-bound and inactive GDP-bound states
	(PubMed:15331635, PubMed:16537643). In its active state, binds to a variety of effector
	proteins playing a key role in the regulation of lysosomal positioning which is important for
	nutrient sensing, natural killer cell-mediated cytotoxicity and antigen presentation. Along with its
	effectors, orchestrates lysosomal transport and fusion (PubMed:16650381, PubMed:16537643,
	PubMed:28325809, PubMed:25898167, PubMed:27808481). Localizes specifically to lysosomal
	membranes and mediates anterograde lysosomal motility by recruiting PLEKHM2, which in turn
	recruits the motor protein kinesin-1 on lysosomes. Required for lysosomal and cytolytic granule
	exocytosis (PubMed:22172677, PubMed:29592961, PubMed:24088571). Critical factor involved
	in NK cell-mediated cytotoxicity. Drives the polarization of cytolytic granules and microtubule-
	organizing centers (MTOCs) toward the immune synapse between effector NK lymphocytes
	and target cells (PubMed:24088571). In neurons, mediates the anterograde axonal long-range
	transport of presynaptic lysosome-related vesicles required for presynaptic biogenesis and
	synaptic function (By similarity). Also acts as a regulator of endosome to lysosome trafficking
	pathways of special significance for host defense (PubMed:21802320). Regulates cargo
	trafficking to lysosomes by binding to PLEKHM1 and recruiting the HOPS subunit VPS41,
	resulting in functional assembly of the HOPS complex on lysosomal membranes
	(PubMed:16537643, PubMed:25908847). Plays an important role in cargo delivery to
	lysosomes for antigen presentation and microbial killing. Directs the intersection of CD1d with
	lipid antigens in lysosomes, and plays a role in intersecting phagosomes with lysosomes to
	generate phagolysosomes that kill microbes (PubMed:25908847, PubMed:21802320). Involved
	in the process of MHC II presentation. Regulates the delivery of antigens to lysosomes and the
	formation of MHC II-peptide complexes through the recruitment of the HOPS complex to
	lysosomes allowing the fusion of late endosomes to lysosomes (By similarity). May play a role
	in chromosome segregation (PubMed:15331635). {ECO:0000250 UniProtKB:Q9CQW2,
	ECO:0000269 PubMed:15331635, ECO:0000269 PubMed:16537643,
	ECO:0000269 PubMed:16650381, ECO:0000269 PubMed:21802320,
	ECO:0000269 PubMed:22172677, ECO:0000269 PubMed:24088571,
	ECO:0000269 PubMed:25898167, ECO:0000269 PubMed:25908847,
	ECO:0000269 PubMed:27808481, ECO:0000269 PubMed:28325809,

ECO:0000269|PubMed:29592961}., FUNCTION: (Microbial infection) During Mycobacterium

tuberculosis (Mtb) infection, is required for plasma membrane repair by controlling the exocytosis of lysosomes in macrophages. ARL8B secretion pathway is crucial to control the type of cell death of the M. tuberculosis-infected macrophages, distinguishing avirulent from virulent Mtb induced necrotic cell death. {ECO:0000269|PubMed:29592961}., FUNCTION: (Microbial infection) During infection, coronaviruses such as SARS-CoV-2 and the chaperone HSPA5/GRP78 are probably co-released through ARL8B-dependent lysosomal exocytic pathway for unconventional egress. {ECO:0000269|PubMed:33157038}.

Molecular Weight:

21.5 kDa

UniProt:

Q9NVJ2

Application Details

Application Notes:

In addition to the applications listed above we expect the protein to work for functional studies as well. As the protein has not been tested for functional studies yet we cannot offer a guarantee though.

Comment:

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During lysate production, the cell wall and other cellular components that are not required for protein production are removed, leaving only the protein production machinery and the mitochondria to drive the reaction. During our lysate completion steps, the additional components needed for protein production (amino acids, cofactors, etc.) are added to produce something that functions like a cell, but without the constraints of a living system - all that's needed is the DNA that codes for the desired protein!

Restrictions:

For Research Use only

Handling

Format:	Liquid
Buffer:	The buffer composition is at the discretion of the manufacturer. If you have a special request, please contact us.
Handling Advice:	Avoid repeated freeze-thaw cycles.
Storage:	-80 °C

Handling

Storage Comment:	Store at -80°C.
Expiry Date:	Unlimited (if stored properly)

Images



Image 1. "Crystallography Grade" protein due to multi-step, protein-specific purification process