

Datasheet for ABIN3130666 NLR Family, Pyrin Domain Containing 1B (NLRP1B) (AA 1-1177) protein (Strep Tag)



Overview

Quantity:	250 µg
Target:	NLR Family, Pyrin Domain Containing 1B (NLRP1B)
Protein Characteristics:	AA 1-1177
Origin:	Mouse
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	Strep Tag
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)

Product Details

Brand:	AliCE®
Sequence:	MEESQYKQEH NKKVAQDEGQ EDKDTIFETI EAIEAKLMEL KTNPESTFNY GIFPEVYMNQ
	GEEILYPAWS LKEENLFQTF KSLRLFQKLC PRGSGNLVKK SWYPCVPEEG GHIINIQDLF
	GPNIGTQKEP QLVIIEGAAG IGKSTLARQV KRAWMEGELY RDHFQHVFFF SCRELAQCKK
	LSLAELITQG QDVPTAPINQ ILSHPEKLLF ILDGIDEPAW VLADQNPELC LYWSQTQPVH
	TLLGSLLGKS ILPEASFLLT TRTTALQKFI PSLPQSCQVE VLGFSDFEQE IYIYKYFAKQ
	IFGIKALMMV ESNPVLLTLC EVPWVCWLVC NCLKKQMEQG GDVSLTSQTT TAICLKYISL
	TIPVHHMRTQ LRALCSLAAE GIWKRRTLFS ESDLCKQGLD EDAVAIFLKT GVLQKQASSL
	SYSFAHLCLQ EFFASMSCIL EDSEERHGDM EMDRIVETLV ERYGRQNLFE APTVRFLFGL
	LSKEGLKEME KLFSCSLPGK TKLKLLWHIL GKSQPHQPPC LGLLHCLYEN QDMKLLTHVM
	HDLQGTIVPD TDDITHTVLQ TNVKHLVVRT DMELMVVTFC IQFCSHMRSL QLNMEGQQGY
	ALTAPRMVLY RWTPITNASW KILFYNLKFN SNLEGLDLSG NPLSYSAVQY LCDAMIYPGC

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	QLKTLWLVEC GLTPTYCSLL ASVLSACSSL RELDLQLNDL CDDGVRMLCE GLRNRACNLR
	ILRLDLYSLS AQVITELRTL EENNLKLHIS SIWMPQMMVP TENMDEEDIL TSFKQQRQQS
	GANPMEILGT EEDFWGPIGP VATEVVYRER NLYRVQLPMA GSYHCPSTRL HFVVTRAVTI
	EIEFCAWSQF LDKTPLQQSH MVVGPLFDIK AEQGAVTAVY LPHFVSLKDT KASTFDFKVA
	HFQEHGMVLE TPDRVKPGYT VLKNPSFSPM GVVLRIIPAA RHFIPITSIT LIYYRVNQEE
	VTLHLYLVPN DCTIQKAIDD EEMKFQFVRI NKPPPVDNLF IGSRYIVSGS ENLEITPKEL
	ELCYRSSKEF QLFSEIYVGN MGSEIKLQIK NKKHMKLIWE ALLKPGDLRP ALPRIAQALK
	DAPSLLHFMD QHREQLVARV TSVDPLLDKL HGLVLNEESY EAVRAENTNQ DKMRKLFNLS
	RSWSRACKDL FYQALKETHP HLVMDLLEKS GGVSLGS
	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 in one-step affinity chromatography
	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 try to ensure that you receive soluble 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 post- translational 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!

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Concentration:

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

Purification:	One-step Strep-tag purification of proteins expressed in Almost Living Cell-Free Expression System (AliCE®).
Purity:	> 70-80 % as determined by SDS PAGE, Western Blot and analytical SEC (HPLC).
Grade:	custom-made

Target Details

Target:	NLR Family, Pyrin Domain Containing 1B (NLRP1B)
Alternative Name:	NIrp1b (NLRP1B Products)
Background:	NACHT, LRR and PYD domains-containing protein 1b allele 2 (EC 3.4) [Cleaved into: NACHT,
	LRR and PYD domains-containing protein 1b, C-terminus (NIrp1b1-CT), NACHT, LRR and PYD
	domains-containing protein 1b, N-terminus (NIrp1b1-NT)],FUNCTION: Acts as the sensor
	component of the NIrp1b inflammasome, which mediates inflammasome activation in
	response to various pathogen-associated signals, leading to subsequent pyroptosis (By
	similarity). Inflammasomes are supramolecular complexes that assemble in the cytosol in
	response to pathogens and other damage-associated signals and play critical roles in innate
	immunity and inflammation (By similarity). Acts as a recognition receptor (PRR): recognizes
	specific pathogens and other damage-associated signals: in response to pathogen-associated
	signals, the N-terminal part of NIrp1b is degraded by the proteasome, releasing the cleaved C-
	terminal part of the protein (NACHT, LRR and PYD domains-containing protein 1b, C-terminus),
	which polymerizes to initiate the formation of the inflammasome complex: the inflammasome
	directly recruits pro-caspase-1 (proCASP1) independently of PYCARD/ASC and promotes
	caspase-1 (CASP1) activation, which subsequently cleaves and activates inflammatory
	cytokines IL1B and IL18 and gasdermin-D (GSDMD), leading to pyroptosis (By similarity). In the
	absence of GSDMD expression, the NIrp1b inflammasome is able to recruit and activate
	CASP8, leading to activation of gasdermin-E (GSDME) (By similarity). Activation of NIrp1b
	inflammasome is also required for HMGB1 secretion, the active cytokines and HMGB1
	stimulate inflammatory responses (By similarity). Contrary to NIrp1b allele 1, allele 2 is not
	activated by Bacillus anthracis lethal toxin (PubMed:16429160, PubMed:21170303,
	PubMed:24492532). {ECO:0000250 UniProtKB:Q2LKW6, ECO:0000269 PubMed:16429160,

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ECO:0000269 PubMed:21170303, ECO:0000269 PubMed:24492532}., FUNCTION: [NACHT, LRR
and PYD domains-containing protein 1b allele 2]: Constitutes the precursor of the NIrp1b
inflammasome, which mediates autoproteolytic processing within the FIIND domain to
generate the N-terminal and C-terminal parts, which are associated non-covalently in absence
of pathogens and other damage-associated signals. {ECO:0000250 UniProtKB:Q2LKW6}.,
FUNCTION: [NACHT, LRR and PYD domains-containing protein 1b, N-terminus]: Regulatory part
that prevents formation of the NIrp1b inflammasome: in absence of pathogens and other
damage-associated signals, interacts with the C-terminal part of NIrp1b (NACHT, LRR and PYD
domains-containing protein 1b, C-terminus), preventing activation of the NIrp1b inflammasome.
In response to pathogen-associated signals, this part is ubiquitinated by the N-end rule pathway
and degraded by the proteasome, releasing the cleaved C-terminal part of the protein, which
polymerizes and forms the NIrp1b inflammasome. {ECO:0000250 UniProtKB:Q2LKW6}.,
FUNCTION: [NACHT, LRR and PYD domains-containing protein 1b, C-terminus]: Constitutes the
active part of the NIrp1b inflammasome. In absence of pathogens and other damage-
associated signals, interacts with the N-terminal part of NIrp1b (NACHT, LRR and PYD
domains-containing protein 1b, N-terminus), preventing activation of the NIrp1b inflammasome.
In response to pathogen-associated signals, the N-terminal part of NIrp1b is degraded by the
proteasome, releasing this form, which polymerizes to form the NIrp1b inflammasome
complex: the NIrp1b inflammasome complex then directly recruits pro-caspase-1 (proCASP1)
and promotes caspase-1 (CASP1) activation, leading to gasdermin-D (GSDMD) cleavage and
subsequent pyroptosis. {ECO:0000250 UniProtKB:Q2LKW6}.

Molecular Weight: 134.2 kDa

UniProt:

A1Z198

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:	 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 post-translational 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

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Application Details	
	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
Format: Buffer:	Liquid The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein.
	The buffer composition is at the discretion of the manufacturer.
Buffer:	The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein.
Buffer: Handling Advice:	The buffer composition is at the discretion of the manufacturer. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein. Avoid repeated freeze-thaw cycles.