antibodies

Datasheet for ABIN3091948 Cullin 4A Protein (CUL4A) (AA 1-759) (Strep Tag)





Overview

Quantity:	1 mg
Target:	Cullin 4A (CUL4A)
Protein Characteristics:	AA 1-759
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	This Cullin 4A protein is labelled with Strep Tag.
Application:	ELISA, Western Blotting (WB), SDS-PAGE (SDS)

Product Details

Sequence:	MADEAPRKGS FSALVGRTNG LTKPAALAAA PAKPGGAGGS KKLVIKNFRD RPRLPDNYTQ
	DTWRKLHEAV RAVQSSTSIR YNLEELYQAV ENLCSHKVSP MLYKQLRQAC EDHVQAQILP
	FREDSLDSVL FLKKINTCWQ DHCRQMIMIR SIFLFLDRTY VLQNSTLPSI WDMGLELFRT
	HIISDKMVQS KTIDGILLLI ERERSGEAVD RSLLRSLLGM LSDLQVYKDS FELKFLEETN
	CLYAAEGQRL MQEREVPEYL NHVSKRLEEE GDRVITYLDH STQKPLIACV EKQLLGEHLT
	AILQKGLDHL LDENRVPDLA QMYQLFSRVR GGQQALLQHW SEYIKTFGTA IVINPEKDKD
	MVQDLLDFKD KVDHVIEVCF QKNERFVNLM KESFETFINK RPNKPAELIA KHVDSKLRAG
	NKEATDEELE RTLDKIMILF RFIHGKDVFE AFYKKDLAKR LLVGKSASVD AEKSMLSKLK
	HECGAAFTSK LEGMFKDMEL SKDIMVHFKQ HMQNQSDSGP IDLTVNILTM GYWPTYTPME
	VHLTPEMIKL QEVFKAFYLG KHSGRKLQWQ TTLGHAVLKA EFKEGKKEFQ VSLFQTLVLL
	MFNEGDGFSF EEIKMATGIE DSELRRTLQS LACGKARVLI KSPKGKEVED GDKFIFNGEF
	KHKLFRIKIN QIQMKETVEE QVSTTERVFQ DRQYQIDAAI VRIMKMRKTL GHNLLVSELY

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NQLKFPVKPG DLKKRIESLI DRDYMERDKD NPNQYHYVA

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

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	(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.
	2. 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.
^D urity:	>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	
Target:	Cullin 4A (CUL4A)
Alternative Name:	CUL4A (CUL4A Products)
Background:	Cullin-4A (CUL-4A),FUNCTION: Core component of multiple cullin-RING-based E3 ubiquitin-
	protein ligase complexes which mediate the ubiquitination of target proteins
	(PubMed:14578910, PubMed:15811626, PubMed:15548678, PubMed:15448697,
	PubMed:14739464, PubMed:16678110, PubMed:17041588, PubMed:24209620,
	PubMed:30166453, PubMed:33854232, PubMed:33854239). As a scaffold protein may
	contribute to catalysis through positioning of the substrate and the ubiquitin-conjugating
	enzyme (PubMed:14578910, PubMed:15811626, PubMed:15548678, PubMed:15448697,
	PubMed:14739464, PubMed:16678110, PubMed:17041588, PubMed:24209620). The E3
	ubiquitin-protein ligase activity of the complex is dependent on the neddylation of the cullin
	subunit and is inhibited by the association of the deneddylated cullin subunit with
	TIP120A/CAND1 (PubMed:14578910, PubMed:15811626, PubMed:15548678,
	PubMed:15448697, PubMed:14739464, PubMed:16678110, PubMed:17041588,
	PubMed:24209620). The functional specificity of the E3 ubiquitin-protein ligase complex
	depends on the variable substrate recognition component (PubMed:14578910,
	PubMed:15811626, PubMed:15548678, PubMed:15448697, PubMed:14739464,
	PubMed:16678110, PubMed:17041588, PubMed:24209620). DCX(DET1-COP1) directs
	ubiquitination of JUN (PubMed:14739464). DCX(DDB2) directs ubiquitination of XPC
	(PubMed:15811626). DCX(DDB2) ubiquitinates histones H3-H4 and is required for efficient
	histone deposition during replication-coupled (H3.1) and replication-independent (H3.3)
	nucleosome assembly, probably by facilitating the transfer of H3 from ASF1A/ASF1B to other
	chaperones involved in histone deposition (PubMed:16678110, PubMed:17041588,

PubMed:24209620). DCX(DTL) plays a role in PCNA-dependent polyubiquitination of CDT1 and
MDM2-dependent ubiquitination of p53/TP53 in response to radiation-induced DNA damage
and during DNA replication (PubMed:14578910, PubMed:15548678, PubMed:15448697).
DCX(DTL) directs autoubiquitination of DTL (PubMed:23478445). In association with DDB1 and
SKP2 probably is involved in ubiquitination of CDKN1B/p27kip (PubMed:16537899). Is involved
in ubiquitination of HOXA9 (PubMed:14609952). The DDB1-CUL4A-DTL E3 ligase complex
regulates the circadian clock function by mediating the ubiquitination and degradation of CRY1
(PubMed:26431207). A number of DCX complexes (containing either TRPC4AP or DCAF12 as
substrate-recognition component) are part of the DesCEND (destruction via C-end degrons)
pathway, which recognizes a C-degron located at the extreme C terminus of target proteins,
leading to their ubiquitination and degradation (PubMed:29779948). The DCX(AMBRA1)
complex is a master regulator of the transition from G1 to S cell phase by mediating
ubiquitination of phosphorylated cyclin-D (CCND1, CCND2 and CCND3) (PubMed:33854232,
PubMed:33854239). The DCX(AMBRA1) complex also acts as a regulator of Cul5-RING (CRL5)
E3 ubiquitin-protein ligase complexes by mediating ubiquitination and degradation of Elongin-C
(ELOC) component of CRL5 complexes (PubMed:30166453). With CUL4B, contributes to
ribosome biogenesis (PubMed:26711351). {ECO:0000269 PubMed:14578910,
ECO:0000269 PubMed:14609952, ECO:0000269 PubMed:14739464,
ECO:0000269 PubMed:15448697, ECO:0000269 PubMed:15548678,
ECO:0000269 PubMed:15811626, ECO:0000269 PubMed:16537899,
ECO:0000269 PubMed:16678110, ECO:0000269 PubMed:17041588,
ECO:0000269 PubMed:23478445, ECO:0000269 PubMed:24209620,
ECO:0000269 PubMed:26431207, ECO:0000269 PubMed:26711351,
ECO:0000269 PubMed:29779948, ECO:0000269 PubMed:30166453,
ECO:0000269 PubMed:33854232, ECO:0000269 PubMed:33854239}.

Molecular Weight:

UniProt:

Q13619

87.7 kDa

Application Details

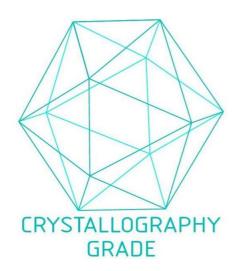
Application Notes:	In addition to the applications listed above we expect the protein to work for functional studies as well. As the protein has not been tested for functional studies yet we cannot offer a guarantee though.
Comment:	ALiCE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from Nicotiana tabacum c.v This contains all the protein expression machinery needed to produce

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	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!
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
Storage Comment:	

Images

Expiry Date:



Unlimited (if stored properly)

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