

Datasheet for ABIN3089640 Aurora Kinase B Protein (AURKB) (AA 1-344) (Strep Tag)



Overview

Quantity:	250 µg
Target:	Aurora Kinase B (AURKB)
Protein Characteristics:	AA 1-344
Origin:	Human
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	This Aurora Kinase B protein is labelled with Strep Tag.
Application:	SDS-PAGE (SDS), Western Blotting (WB), ELISA

Product Details

Brand:	AliCE®
Sequence:	MAQKENSYPW PYGRQTAPSG LSTLPQRVLR KEPVTPSALV LMSRSNVQPT AAPGQKVMEN
	SSGTPDILTR HFTIDDFEIG RPLGKGKFGN VYLAREKKSH FIVALKVLFK SQIEKEGVEH
	QLRREIEIQA HLHHPNILRL YNYFYDRRRI YLILEYAPRG ELYKELQKSC TFDEQRTATI
	MEELADALMY CHGKKVIHRD IKPENLLLGL KGELKIADFG WSVHAPSLRR KTMCGTLDYL
	PPEMIEGRMH NEKVDLWCIG VLCYELLVGN PPFESASHNE TYRRIVKVDL KFPASVPMGA
	QDLISKLLRH NPSERLPLAQ VSAHPWVRAN SRRVLPPSAL QSVA
	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:

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- 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 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 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:

Aurora Kinase B (AURKB)

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Target Details	
Alternative Name:	AURKB (AURKB Products)
Background:	Aurora kinase B (EC 2.7.11.1) (Aurora 1) (Aurora- and IPL1-like midbody-associated protein 1)
	(AIM-1) (Aurora/IPL1-related kinase 2) (ARK-2) (Aurora-related kinase 2) (STK-1)
	(Serine/threonine-protein kinase 12) (Serine/threonine-protein kinase 5) (Serine/threonine-
	protein kinase aurora-B),FUNCTION: Serine/threonine-protein kinase component of the
	chromosomal passenger complex (CPC), a complex that acts as a key regulator of mitosis
	(PubMed:11516652, PubMed:12925766, PubMed:14610074, PubMed:14722118,
	PubMed:29449677). The CPC complex has essential functions at the centromere in ensuring
	correct chromosome alignment and segregation and is required for chromatin-induced
	microtubule stabilization and spindle assembly (PubMed:11516652, PubMed:12925766,
	PubMed:14610074, PubMed:14722118, PubMed:26829474). Involved in the bipolar attachment
	of spindle microtubules to kinetochores and is a key regulator for the onset of cytokinesis
	during mitosis (PubMed:15249581). Required for central/midzone spindle assembly and
	cleavage furrow formation (PubMed:12458200, PubMed:12686604). Key component of the
	cytokinesis checkpoint, a process required to delay abscission to prevent both premature
	resolution of intercellular chromosome bridges and accumulation of DNA damage:
	phosphorylates CHMP4C, leading to retain abscission-competent VPS4 (VPS4A and/or VPS4B)
	at the midbody ring until abscission checkpoint signaling is terminated at late cytokinesis
	(PubMed:22422861, PubMed:24814515). AURKB phosphorylates the CPC complex subunits
	BIRC5/survivin, CDCA8/borealin and INCENP (PubMed:11516652, PubMed:12925766,
	PubMed:14610074). Phosphorylation of INCENP leads to increased AURKB activity
	(PubMed:11516652, PubMed:12925766, PubMed:14610074). Other known AURKB substrates
	involved in centromeric functions and mitosis are CENPA, DES/desmin, GPAF, KIF2C, NSUN2,
	RACGAP1, SEPTIN1, VIM/vimentin, HASPIN, and histone H3 (PubMed:11784863,
	PubMed:12689593, PubMed:14602875, PubMed:11856369, PubMed:16103226,
	PubMed:21658950, PubMed:11756469). A positive feedback loop involving HASPIN and AURKB
	contributes to localization of CPC to centromeres (PubMed:21658950). Phosphorylation of VIM
	controls vimentin filament segregation in cytokinetic process, whereas histone H3 is
	phosphorylated at 'Ser-10' and 'Ser-28' during mitosis (H3S10ph and H3S28ph, respectively)
	(PubMed:11784863, PubMed:11856369). AURKB is also required for kinetochore localization of
	BUB1 and SGO1 (PubMed:15020684, PubMed:17617734). Phosphorylation of p53/TP53
	negatively regulates its transcriptional activity (PubMed:20959462). Key regulator of active
	promoters in resting B- and T-lymphocytes: acts by mediating phosphorylation of H3S28ph at
	active promoters in resting B-cells, inhibiting RNF2/RING1B-mediated ubiquitination of histone
	H2A and enhancing binding and activity of the USP16 deubiquitinase at transcribed genes (By

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	similarity). Acts as an inhibitor of CGAS during mitosis: catalyzes phosphorylation of the N-
	terminus of CGAS during the G2-M transition, blocking CGAS liquid phase separation and
	activation, and thereby preventing CGAS-induced autoimmunity (PubMed:33542149).
	Phosphorylates KRT5 during anaphase and telophase (By similarity).
	{ECO:0000250 UniProtKB:070126, ECO:0000269 PubMed:11516652,
	ECO:0000269 PubMed:11756469, ECO:0000269 PubMed:11784863,
	ECO:0000269 PubMed:11856369, ECO:0000269 PubMed:12458200,
	ECO:0000269 PubMed:12686604, ECO:0000269 PubMed:12689593,
	ECO:0000269 PubMed:12925766, ECO:0000269 PubMed:14602875,
	ECO:0000269 PubMed:14610074, ECO:0000269 PubMed:14722118,
	ECO:0000269 PubMed:15020684, ECO:0000269 PubMed:15249581,
	ECO:0000269 PubMed:16103226, ECO:0000269 PubMed:17617734,
	ECO:0000269 PubMed:20959462, ECO:0000269 PubMed:21658950,
	ECO:0000269 PubMed:22422861, ECO:0000269 PubMed:24814515,
	ECO:0000269 PubMed:26829474, ECO:0000269 PubMed:29449677,
	ECO:0000269 PubMed:33542149}.
Molecular Weight:	39.3 kDa
UniProt:	Q96GD4
Pathways:	TCR Signaling, Cell Division Cycle, Maintenance of Protein Location, Hepatitis C, Toll-Like
	Receptors Cascades
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

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Application Details		
	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. Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol Might differ depending on protein.	
Handling Advice:	Avoid repeated freeze-thaw cycles.	
Storage:	-80 °C	
Storage Comment:	Store at -80°C.	
Expiry Date:	12 months	