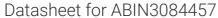
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NME3 Protein (AA 1-169) (Strep Tag)



Image



Go to Product page

Overview

Quantity:	1 mg
Target:	NME3
Protein Characteristics:	AA 1-169
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	This NME3 protein is labelled with Strep Tag.
Application:	SDS-PAGE (SDS), Western Blotting (WB), ELISA

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Product Details	
Sequence:	MICLVLTIFA NLFPAACTGA HERTFLAVKP DGVQRRLVGE IVRRFERKGF KLVALKLVQA
	SEELLREHYA ELRERPFYGR LVKYMASGPV VAMVWQGLDV VRTSRALIGA TNPADAPPGT
	IRGDFCIEVG KNLIHGSDSV ESARREIALW FRADELLCWE DSAGHWLYE
	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

· These proteins are normally active (enzymatically functional) as our customers have

reported (not tested by us and not guaranteed).

correct folding and modification.

• 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

Target:	NME3
Alternative Name:	NME3 (NME3 Products)
Background:	Nucleoside diphosphate kinase 3 (NDK 3) (NDP kinase 3) (EC 2.7.4.6) (DR-nm23) (Nucleoside
	diphosphate kinase C) (NDPKC) (nm23-H3),FUNCTION: Catalyzes the phosphorylation of
	ribonucleosides and deoxyribonucleoside diphosphates, other than ATP, into the corresponding
	triphosphates with ATP as the major phosphate donor (PubMed:11277919,
	PubMed:30587587). The ATP gamma phosphate is transferred to the nucleoside diphosphate
	beta phosphate via a ping-pong mechanism, using a phosphorylated active-site intermediate.
	Through the catalyzed exchange of gamma-phosphate between di- and triphosphonucleosides
	participates in regulation of intracellular nucleotide homeostasis (PubMed:11277919,
	PubMed:30587587). Inhibits granulocyte differentiation (PubMed:7638209). May be required for
	ciliary function during renal development (By similarity). {ECO:0000250 UniProtKB:Q9PTF3,
	ECO:0000269 PubMed:11277919, ECO:0000269 PubMed:30587587,
	ECO:0000269 PubMed:7638209}., FUNCTION: Independently of its kinase activity, facilitates
	mitochondrial tethering prior to membrane fusion through its direct membrane-binding and
	hexamerization (PubMed:30587587, PubMed:37584589). Implicated in repair of both single-
	and double-stranded breaks in DNA through its association with the ribonucleotide reductase
	complex (RNR complex) via its interaction with the histone acetyltransferase KAT5, this
	interaction enables recruitment of NME3 at DNA damage sites where it plays a role in the repair
	of DNA, independently of its kinase activity (PubMed:37584589).
	{ECO:0000269 PubMed:30587587, ECO:0000269 PubMed:37584589}.
Molecular Weight:	19.0 kDa
UniProt:	Q13232
Pathways:	Nucleotide Phosphorylation, Ribonucleoside Biosynthetic Process
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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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:	Store at -80°C.
Expiry Date:	Unlimited (if stored properly)

Images



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