

Datasheet for ABIN3130839

METTL8 Protein (AA 1-281) (Strep Tag)



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Quantity:	250 μg
Target:	METTL8
Protein Characteristics:	AA 1-281
Origin:	Mouse
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	This METTL8 protein is labelled with Strep Tag.
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)

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Product Details	
Brand:	AliCE®
Sequence:	MNVIWRSCIC RLRQGKVPHR CQSGVHPVAP LGSRILTDPA KVFEHNMWDH MQWSKEEEDA
	ARKKVEENSA TRVAPEEQVK FESDANKYWD IFYQTHKNKF FKNRNWLLRE FPEILPVNQN
	TKEKVGESSW DQVGSSISRT QGTETHCQES FVSPEPGSRG RSAPDPDLEE YSKGPGKTEP
	FPGSNATFRI LEVGCGAGNS VFPILNTLQN IPGSFLYCCD FASEAVELVK SHESYSEAQC
	SAFIHDVCDD GLAYPFPDGI LDVVLLVFVL SSIHPDRALF I
	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 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
Torget Details	

Target Details

Target:	METTL8
Alternative Name:	Mettl8 (METTL8 Products)

Background:

TRNA N(3)-methylcytidine methyltransferase METTL8, mitochondrial (EC 2.1.1.-) (Methyltransferase-like protein 8) (Tension-induced/inhibited protein) (mRNA N(3)methylcytidine methyltransferase METTL8) (EC 2.1.1.-),FUNCTION: Mitochondrial S-adenosyl-Lmethionine-dependent methyltransferase that mediates N(3)-methylcytidine modification of residue 32 of the tRNA anticodon loop of mitochondrial tRNA(Ser)(UCN) and tRNA(Thr) (By similarity). N(3)-methylcytidine methylation modification regulates mitochondrial translation efficiency and is required for activity of the respiratory chain (By similarity). N(3)-methylcytidine methylation of mitochondrial tRNA(Ser)(UCN) requires the formation of N(6)dimethylallyladenosine(37) (i6A37) by TRIT1 as prerequisite (By similarity). May also mediate N(3)-methylcytidine modification of mRNAs (PubMed:28655767). The existence of N(3)methylcytidine modification on mRNAs is however unclear, and additional evidences are required to confirm the role of the N(3)-methylcytidine-specific mRNA methyltransferase activity of METTL8 in vivo (By similarity). {ECO:0000250|UniProtKB:Q9H825, ECO:0000269|PubMed:28655767}., FUNCTION: [Isoform 5]: Overexpression in lung progenitor cells stimulates smooth muscle-specific gene expression and suppresses adipogenic gene expression. {ECO:0000269|PubMed:15992539}., FUNCTION: [Isoform 4]: Stimulates adipogenesis. {ECO:0000269|PubMed:18710950}., FUNCTION: [Isoform 7]: Stimulates adipogenesis. {ECO:0000269|PubMed:18710950}.

Molecular Weight:

31.7 kDa

UniProt:

A2AUU0

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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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