

Datasheet for ABIN3103333

Fat Storage-Inducing Transmembrane Protein 2 (FITM2) (AA 1-262) protein (Strep Tag)



()	ve	r\/i	۱۸/
\cup	V C	1 / 1	 v v

Quantity:	250 μg
Target:	Fat Storage-Inducing Transmembrane Protein 2 (FITM2)
Protein Characteristics:	AA 1-262
Origin:	Human
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	Strep Tag
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)

A I:	FLICA ODO DAGE (ODO) Washam Blatting (MD)
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)
Product Details	
Brand:	AliCE®
Sequence:	MEHLERCEWL LRGTLVRAAV RRYLPWALVA SMLAGSLLKE LSPLPESYLS NKRNVLNVYF
	VKVAWAWTFC LLLPFIALTN YHLTGKAGLV LRRLSTLLVG TAIWYICTSI FSNIEHYTGS
	CYQSPALEGV RKEHQSKQQC HQEGGFWHGF DISGHSFLLT FCALMIVEEM SVLHEVKTDR
	SHCLHTAITT LVVALGILTF IWVLMFLCTA VYFHNLSQKV FGTLFGLLSW YGTYGFWYPK
	AFSPGLPPQS CSLNLKQDSY KK
	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.

Alternative Name:

- · 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:

FITM2 (FITM2 Products)

- 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:	Fat Storage-Inducing Transmembrane Protein 2 (FITM2)

Background:

Acyl-coenzyme A diphosphatase FITM2 (EC 3.6.1.-) (Fat storage-inducing transmembrane protein 2) (Fat-inducing protein 2),FUNCTION: Fatty acyl-coenzyme A (CoA) diphosphatase that hydrolyzes fatty acyl-CoA to yield acyl-4'-phosphopantetheine and adenosine 3',5'-bisphosphate (PubMed:32915949) (By similarity). Preferentially hydrolyzes unsaturated long-chain acyl-CoA substrates such as oleoyl-CoA/(9Z)-octadecenoyl-CoA and arachidonoyl-CoA/(5Z,8Z,11Z,14Z)-eicosatetraenoyl-CoA in the endoplasmic reticulum (ER) lumen (PubMed:32915949) (By similarity). This catalytic activity is required for maintaining ER structure and for lipid droplets (LDs) biogenesis, which are lipid storage organelles involved in maintaining lipid and energy homeostasis (PubMed:18160536, PubMed:32915949) (By similarity). Directly binds to diacylglycerol (DAGs) and triacylglycerol, which is also important for LD biogenesis (By similarity). May support directional budding of nacent LDs from the ER into the cytosol by reducing DAG levels at sites of LD formation (By similarity). Plays a role in the regulation of cell morphology and cytoskeletal organization (PubMed:21834987) (By similarity). (ECO:0000255|HAMAP-Rule:MF_03230, ECO:0000269|PubMed:18160536, ECO:0000269|PubMed:21834987, ECO:0000269|PubMed:32915949}.

Molecular Weight:

29.9 kDa

UniProt:

Q8N6M3

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 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. 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