antibodies

Datasheet for ABIN3118594 ELOVL2 Protein (AA 1-296) (Strep Tag)





Overview

Quantity:	1 mg
Target:	ELOVL2
Protein Characteristics:	AA 1-296
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	This ELOVL2 protein is labelled with Strep Tag.
Application:	ELISA, Western Blotting (WB), SDS-PAGE (SDS)

Product Details

Sequence:	MEHLKAFDDE INAFLDNMFG PRDSRVRGWF MLDSYLPTFF LTVMYLLSIW LGNKYMKNRP
	ALSLRGILTL YNLGITLLSA YMLAELILST WEGGYNLQCQ DLTSAGEADI RVAKVLWWYY
	FSKSVEFLDT IFFVLRKKTS QITFLHVYHH ASMFNIWWCV LNWIPCGQSF FGPTLNSFIH
	ILMYSYYGLS VFPSMHKYLW WKKYLTQAQL VQFVLTITHT MSAVVKPCGF PFGCLIFQSS
	YMLTLVILFL NFYVQTYRKK PMKKDMQEPP AGKEVKNGFS KAYFTAANGV MNKKAQ
	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
	system, a different complexity of the protein could make another tag necessary. In case you have a special request, please contact us.
Characteristics:	
Characteristics:	have a special request, please contact us.

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- 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 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!

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®):
	 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)

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Product Details

Grade:

Crystallography grade

Target Details

Target:	ELOVL2
Alternative Name:	ELOVL2 (ELOVL2 Products)
Background:	Very long chain fatty acid elongase 2 (EC 2.3.1.199) (3-keto acyl-CoA synthase ELOVL2) (ELOVI
	fatty acid elongase 2) (ELOVL FA elongase 2) (Elongation of very long chain fatty acids protein
	2) (Very long chain 3-ketoacyl-CoA synthase 2) (Very long chain 3-oxoacyl-CoA synthase
	2),FUNCTION: Catalyzes the first and rate-limiting reaction of the four reactions that constitute
	the long-chain fatty acids elongation cycle. This endoplasmic reticulum-bound enzymatic
	process allows the addition of 2 carbons to the chain of long- and very long-chain fatty acids
	(VLCFAs) per cycle. Condensing enzyme that catalyzes the synthesis of polyunsaturated very
	long chain fatty acid (C20- and C22-PUFA), acting specifically toward polyunsaturated acyl-CoA
	with the higher activity toward C20:4(n-6) acyl-CoA. May participate in the production of
	polyunsaturated VLCFAs of different chain lengths that are involved in multiple biological
	processes as precursors of membrane lipids and lipid mediators. {ECO:0000255 HAMAP-
	Rule:MF_03202, ECO:0000269 PubMed:11734209, ECO:0000269 PubMed:12371743,
	EC0:0000269 PubMed:19575253, EC0:0000269 PubMed:20937905}.
Molecular Weight:	34.6 kDa
UniProt:	Q9NXB9
UniProt: Application Details	Q9NXB9
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Application Details	
	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:	Store at -80°C.
Expiry Date:	Unlimited (if stored properly)

Images

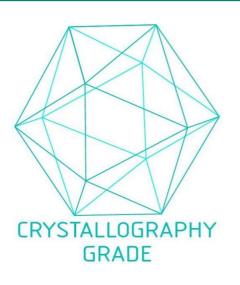


Image 1. "Crystallography Grade" protein due to multi-step, protein-specific purification process

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