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Meningioma 1 Protein (AA 1-1320) (Strep Tag)



Image



Overview

Quantity:	1 mg
Target:	Meningioma 1 (MN1)
Protein Characteristics:	AA 1-1320
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	This Meningioma 1 protein is labelled with Strep Tag.
Application:	ELISA, Western Blotting (WB), SDS-PAGE (SDS)

Product Details

Sequence:

LPSAASERRP PPPDFATSAL GGQPGFPFGA AGRQSTPHSG PGVNSPPSAG GGGGSSGGGG
GGGAYPPQPD FQPSQRTSAS KLGALSLGSF NKPSSKDNLF GQSCLAALST ACQNMIASLG
APNLNVTFNK KNPPEGKRKL SQNETDGAAV AGNPGSDYFP GGTAPGAPGP GGPSGTSSSG
SKASGPPNPP AQGDGTSLSP NYTLESTSGN DGKPVSGGGG RGRGRRKRDS GHVSPGTFFD
KYSAAPDSGG APGVSPGQQQ ASGAAVGGSS AGETRGAPTP HEKALTSPSW GKGAELLLGD
QPDLIGSLDG GAKSDSSSPN VGEFASDEVS TSYANEDEVS SSSDNPQALV KASRSPLVTG
SPKLPPRGVG AGEHGPKAPP PALGLGIMSN STSTPDSYGG GGGPGHPGTP GLEQVRTPTS
SSGAPPPDEI HPLEILQAQI QLQRQQFSIS EDQPLGLKGG KKGECAVGAS GAQNGDSELG
SCCSEAVKSA MSTIDLDSLM AEHSAAWYMP ADKALVDSAD DDKTLAPWEK AKPQNPNSKE
AHDLPANKAS ASQPGSHLQC LSVHCTDDVG DAKARASVPT WRSLHSDISN RFGTFVAALT

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 correct folding and modification.
- 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 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:	Meningioma 1 (MN1)
Alternative Name:	MN1 (MN1 Products)
Background:	Transcriptional activator MN1 (Probable tumor suppressor protein MN1),FUNCTION:
	Transcriptional activator which specifically regulates expression of TBX22 in the posterior
	region of the developing palate. Required during later stages of palate development for growth
	and medial fusion of the palatal shelves. Promotes maturation and normal function of calvarial
	osteoblasts, including expression of the osteoclastogenic cytokine TNFSF11/RANKL.
	Necessary for normal development of the membranous bones of the skull (By similarity). May
	play a role in tumor suppression (Probable). {ECO:0000250 UniProtKB:D3YWE6,
	ECO:0000305 PubMed:7731706}.
Molecular Weight:	136.0 kDa
UniProt:	Q10571

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. 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)



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