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Datasheet for ABIN3079307

Crossover junction endonuclease EME1 (EME1) (AA 1-570) protein (Strep Tag)



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Overview

Quantity:	1 mg
Target:	Crossover junction endonuclease EME1 (EME1)
Protein Characteristics:	AA 1-570
Origin:	Human
Source:	Tobacco (Nicotiana tabacum)
Protein Type:	Recombinant
Purification tag / Conjugate:	Strep Tag
Application:	SDS-PAGE (SDS), ELISA, Western Blotting (WB)

Product Details

Sequence:

MALKKSSPSL DSGDSDSEEL PTFAFLKKEP SSTKRRQPER EEKIVVVDIS DCEASCPPAP ELFSPPVPEI AETVTQTQPV RLLSSESEDE EEFIPLAQRL TCKFLTHKQL SPEDSSSPVK SVLDHQNNEG ASCDWKKPFP KIPEVPLHDT PERSAADNKD LILDPCCQLP AYLSTCPGQS SSLAVTKTNS DILPPQKKTK PSQKVQGRGS HGCRQQRQAR QKESTLRRQE RKNAALVTRM KAQRPEECLK HIIVVLDPVL LQMEGGGQLL GALQTMECRC VIEAQAVPCS VTWRRRAGPS EDREDWVEEP TVLVLLRAEA FVSMIDNGKQ GSLDSTMKGK ETLQGFVTDI TAKTAGKALS LVIVDQEKCF SAQNPPRRGK QGANKQTKKQ QQRQPEASIG SMVSRVDAEE ALVDLQLHTE AQAQIVQSWK ELADFTCAFT KAVAEAPFKK LRDETTFSFC LESDWAGGVK VDLAGRGLAL VWRRQIQQLN RVSLEMASAV VNAYPSPQLL VQAYQQCFSD KERQNLLADI QVRRGEGVTS TSRRIGPELS RRIYLQMTTL QPHLSLDSAD

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:	Crossover junction endonuclease EME1 (EME1)
Alternative Name:	EME1 (EME1 Products)
Background:	Crossover junction endonuclease EME1 (EC 3.1.22) (MMS4 homolog) (hMMS4),FUNCTION: Interacts with MUS81 to form a DNA structure-specific endonuclease with substrate preference for branched DNA structures with a 5'-end at the branch nick. Typical substrates include 3'-flap structures, replication forks and nicked Holliday junctions. May be required in mitosis for the processing of stalled or collapsed replication forks. {ECO:0000269 PubMed:12686547, ECO:0000269 PubMed:12721304, ECO:0000269 PubMed:14617801, ECO:0000269 PubMed:17289582}.
Molecular Weight:	63.3 kDa
UniProt:	Q96AY2
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

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

CRYSTALLOGRAPHY
GRADE

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