

# Datasheet for ABIN3092481 EXOSC6 Protein (AA 1-272) (Strep Tag)



Overview

Quantity:	250 µg
Target:	EXOSC6
Protein Characteristics:	AA 1-272
Origin:	Human
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	This EXOSC6 protein is labelled with Strep Tag.
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)

## Product Details

Brand:	AliCE®
Sequence:	MPGDHRRIRG PEESQPPQLY AADEEEAPGT RDPTRLRPVY ARAGLLSQAK GSAYLEAGGT
	KVLCAVSGPR QAEGGERGGG PAGAGGEAPA ALRGRLLCDF RRAPFAGRRR RAPPGGCEER
	ELALALQEAL EPAVRLGRYP RAQLEVSALL LEDGGSALAA ALTAAALALA DAGVEMYDLV
	VGCGLSLAPG PAPTWLLDPT RLEEERAAAG LTVALMPVLN QVAGLLGSGE GGLTESWAEA
	VRLGLEGCQR LYPVLQQSLV RAARRRGAAA QP
	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.

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- 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 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 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:	EXOSC6
Alternative Name:	EXOSC6 (EXOSC6 Products)

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Exosome complex component MTR3 (Exosome component 6) (mRNA transport regulator 3
homolog) (hMtr3) (p11),FUNCTION: Non-catalytic component of the RNA exosome complex
which has 3'->5' exoribonuclease activity and participates in a multitude of cellular RNA
processing and degradation events. In the nucleus, the RNA exosome complex is involved in
proper maturation of stable RNA species such as rRNA, snRNA and snoRNA, in the elimination
of RNA processing by-products and non-coding 'pervasive' transcripts, such as antisense RNA
species and promoter-upstream transcripts (PROMPTs), and of mRNAs with processing
defects, thereby limiting or excluding their export to the cytoplasm. The RNA exosome may be
involved in Ig class switch recombination (CSR) and/or Ig variable region somatic
hypermutation (SHM) by targeting AICDA deamination activity to transcribed dsDNA
substrates. In the cytoplasm, the RNA exosome complex is involved in general mRNA turnover
and specifically degrades inherently unstable mRNAs containing AU-rich elements (AREs)
within their 3' untranslated regions, and in RNA surveillance pathways, preventing translation of
aberrant mRNAs. It seems to be involved in degradation of histone mRNA. The catalytic
inactive RNA exosome core complex of 9 subunits (Exo-9) is proposed to play a pivotal role in
the binding and presentation of RNA for ribonucleolysis, and to serve as a scaffold for the
association with catalytic subunits and accessory proteins or complexes.
{EC0:0000269 PubMed:21255825}.
28.2 kDa
Q5RKV6
Regulation of Leukocyte Mediated Immunity, Positive Regulation of Immune Effector Process, Production of Molecular Mediator of Immune Response
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.
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Application Details	
	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 <b>Might differ depending on protein.</b>
Handling Advice:	Avoid repeated freeze-thaw cycles.
Storage:	-80 °C
Storage Comment:	Store at -80°C.
Expiry Date:	12 months