

Datasheet for ABIN3075947

## ZNF341 Protein (AA 1-854) (Strep Tag)



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### 1 Image

#### Overview

Quantity:	1 mg
Target:	ZNF341
Protein Characteristics:	AA 1-854
Origin:	Human
Source:	Tobacco ( <i>Nicotiana tabacum</i> )
Protein Type:	Recombinant
Purification tag / Conjugate:	This ZNF341 protein is labelled with Strep Tag.
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)

#### Product Details

Sequence: MAQAIFEALE GMDNQTVLAV QSLLDGQGAV PDPTGQSVNA PPAIQPLDDE DVFLCGKCKK  
QFNSLPAFMT HKREQCQGNA PALATVSLAT NSIYPPSAAP TAVQQAPTPA NRQISTYITV  
PPSPLIQTLV QGNILVSDDV LMSAMSAFTS LDQPMPQGPP PVQSSLNMHS VPSYLTQPPP  
PPPPPPPLPP PPPPQPPPP PQLGPPGRP NPGGNGVVEV YSAAAPLAGS GTVEIQALGM  
QPYPPLEVPN QCVEPPVYPT PTVYSPGKQG FKPKGPNPAA PMTSATGGTV ATFDSPATLK  
TRRAKGARGL PEAAGKPKAQ KLKCSYCDKS FTKNFDLQQH IRSHTGEKPF QCIACGRAFA  
QKSNVKKHMQ THKVVPPGHS GGTVSRNSVT VQVMALNPSR QEDEESTGLG QPLPGAPQPQ  
ALSTAGEEEG DKPESKQVVL IDSSYLCQFC PSKFSTYFQL KSHMTQHKNE QVYKCVKSC  
AQTFPKLDTF LEHIKSHQEE LSYRCHLCGK DFPSLYDLGV HQYSHSLLPQ HSPKKDNAVY  
KCVKCVNKYS TPEALEHHLQ TATHNFPCPH CQKVFPCEY LRRHLPTHGS GGRFKCQVCK  
KFFRREHYLK LHAHIHSGEK PYKCSVCESA FNRKDKLKRH MLIHEPFKKY KCPFSTHTGC  
SKEFNRPDKL KAHILSHSGM KLHKCALCSK SFSRRAHLAE HQRAHTGNYK FRCAGCAKGF

SRHKYLDKDR CRLGPQKDKD LQTRRPPQRR AAPRSCGSGG RKVLTPLPDP LGLEELKDTG  
AGLVPEAVPG KPPFAEPDAV LSIVVGGAVG AETELVVP GH AEGLGSNLAL AELQAGAEGP  
CAMLAVPVYI QASE

**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.**

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### 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 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.

## Product Details

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Purification:	Two step purification of proteins expressed in Almost Living Cell-Free Expression System (ALiCE®): <ol style="list-style-type: none"><li>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.</li><li>2. 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.</li></ol>
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

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Target:	ZNF341
Alternative Name:	ZNF341 ( <a href="#">ZNF341 Products</a> )
Background:	Zinc finger protein 341,FUNCTION: Transcriptional activator of STAT3 involved in the regulation of immune homeostasis. Also able to activate STAT1 transcription. {ECO:0000269 PubMed:29907690, ECO:0000269 PubMed:29907691}.
Molecular Weight:	92.7 kDa
UniProt:	<a href="#">Q9BYN7</a>

## Application Details

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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 <i>Nicotiana tabacum</i> 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

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Restrictions: For Research Use only

## Handling

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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

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**Image 1.** „Crystallography Grade“ protein due to multi-step, protein-specific purification process