

Datasheet for ABIN3132280

## SETDB1 Protein (AA 1-1307) (Strep Tag)



[Go to Product page](#)

### Overview

Quantity:	250 µg
Target:	SETDB1
Protein Characteristics:	AA 1-1307
Origin:	Mouse
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	This SETDB1 protein is labelled with Strep Tag.
Application:	ELISA, SDS-PAGE (SDS), Western Blotting (WB)

### Product Details

Brand:	AlIcE®
Sequence:	<p>MSSLPGCMSL AAPAAADSA EIAELQQAVV EELGISMEEL RQYIDEELEK MDCIQQRKKQ          LAELETWVLQ KESEVAYVDR LFDDASREVT NCESLVKDFY SKLGLQYHDS SSEDEASRPT          EIIEIPDEDD DVLSIDSGDA GSRTPKDQKL REAMAALRKS AQDVQKFMDA VNKKSSSQDL          HKGTLGQVSG ELSKDGD LIV SMRILGKKRT KTW HKGT LIA IQTVGLGKKY KVKFDNKGKS          LLSGNHIA YD YHPPADKLFV GSRVVAKYKD GNQVWLYAGI VAETPNVKNK LRFLIFFDDG          YASYVTQSEL YPICRPLKKT WEDIEDSSCR DFIEEYITAY PNRPMVLLKS GQLIKTEWEG          TWWKSRVEEV DGSLVRILFL DDKRCEWIYR GSTRLEPMFS MKTSSASAME KKQGGQLRTR          PNMGAVRSKG PVVQYTQDLT GTGIQFKPME PLQPIAPPAP LPIPLSPQA ADTDLESQLA          QSRKQVAKKS TSFRPGSVGS GHSSPTSSTL SENVSAGKLG INQTYRSPLA SVTSTPASAA          PPVPPVPPGP PTPPGPPAPP GPLAPPAFHG MLERAPAEPS YRAPMEKLFY LPHVCSYTCL          SRIRPMRNEQ YRGKNPLLVP LLYDFRRMTA RRRVNRKMGF HVIYKTPCGL CLRTMQEIER</p>

YLFETGCDFL FLEMFCLDPY VLVDKRFQPF KPFYYILDIT YGKEDVPLSC VNEIDTTPPP  
QVAYSKERIP GKGVFINTGP EFLVGCDCKD GCRDKSKCAC HQLTIQATAC TPGGQVNPNS  
GYQYKRLEEC LPTGVYECNK RCNCDPNMCT NRLVQHGLQV RLQLFKTQNK GWGIRCLDDI  
AKGSFVCIYA GKILTDDFAD KEGLEMGEY FANLDHIESV ENFKEGYESD VPTSSDSSGV  
DMKDQEDGNS GSEDPEESND DSSDDNFCKD EDFSTSSVWR SYATRRQTRG QKENELSEMT  
SKDSRPPDLG PPHVPISSV SVGGCNPPSS EETPKNKVAS WLSCNSVSEG GFADSDSRSS  
FKTSEGGDGR AGGGRGEAER ASTSGLSFKD EGDNKQPKKE DPENRNKMPV VTEGSQNHGH  
NPPMKSEGLR RPASKMSVLQ SQRVVTSTQS NPDDILTLSS STESEGESGT SRKPTAGHTS  
ATAVDSDDIQ TISSGSDGDD FEDKKNLSGP TKRQVAVKST RGFALKSTHG IAIKSTNMAS  
VDKGESAPVR KNTRQFYDGE ESCYIIDAKL EGNLGRYLNH SCSPNLFVQN VFVDTHDLRF  
PWVAFFASKR IRAGTELTWD YNYEVGSVEG KELLCCCGAI ECRGRLL

**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 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 -

## Product Details

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®).
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Purity:	> 70-80 % as determined by SDS PAGE, Western Blot and analytical SEC (HPLC).
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Grade:	custom-made
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## Target Details

Target:	SETDB1
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Alternative Name:	Setdb1 ( <a href="#">SETDB1 Products</a> )
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Background:	<p>Histone-lysine N-methyltransferase SETDB1 (EC 2.1.1.366) (ERG-associated protein with SET domain) (ESET) (SET domain bifurcated 1),FUNCTION: Histone methyltransferase that specifically trimethylates 'Lys-9' of histone H3 (PubMed:11791185, PubMed:22939622). H3 'Lys-9' trimethylation represents a specific tag for epigenetic transcriptional repression by recruiting HP1 (CBX1, CBX3 and/or CBX5) proteins to methylated histones. Mainly functions in euchromatin regions, thereby playing a central role in the silencing of euchromatic genes. H3 'Lys-9' trimethylation is coordinated with DNA methylation. Probably forms a complex with MBD1 and ATF7IP that represses transcription and couples DNA methylation and histone 'Lys-9' trimethylation. Its activity is dependent on MBD1 and is heritably maintained through DNA replication by being recruited by CAF-1. SETDB1 is targeted to histone H3 by TRIM28/TIF1B, a factor recruited by KRAB zinc-finger proteins. Probably forms a corepressor complex required for activated KRAS-mediated promoter hypermethylation and transcriptional silencing of tumor suppressor genes (TSGs) or other tumor-related genes in colorectal cancer (CRC) cells (By similarity). Required to maintain a transcriptionally repressive state of genes in undifferentiated embryonic stem cells (ESCs) (By similarity). In ESCs, in collaboration with TRIM28, is also required for H3K9me3 and silencing of endogenous and introduced retroviruses in a DNA-methylation independent-pathway (PubMed:20164836, PubMed:29728365). Associates at promoter regions of tumor suppressor genes (TSGs) leading to their gene silencing. The</p>
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## Target Details

SETDB1-TRIM28-ZNF274 complex may play a role in recruiting ATRX to the 3'-exons of zinc-finger coding genes with atypical chromatin signatures to establish or maintain/protect H3K9me3 at these transcriptionally active regions (By similarity).

{ECO:0000250|UniProtKB:Q15047, ECO:0000269|PubMed:11791185, ECO:0000269|PubMed:20164836, ECO:0000269|PubMed:22939622, ECO:0000269|PubMed:29728365}.

Molecular Weight: 144.5 kDa

UniProt: [O88974](#)

## 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.

Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol **Might differ depending on protein.**

Handling Advice: Avoid repeated freeze-thaw cycles.

Storage: -80 °C

Storage Comment: Store at -80°C.

## Handling

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Expiry Date: 12 months