abcam

Product datasheet

Recombinant human CREBBP protein ab167964

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Description

Product name Recombinant human CREBBP protein

Biological activity

The specific activity of ab167964 was determined to be 32 nmol/min/mg as per activity assay

protocol.

Purity > 90 % SDS-PAGE.

Assessed by densitometry. Affinity purified.

Expression system Baculovirus infected Sf9 cells

Accession Q92793

Protein length Protein fragment

Animal free No

Nature Recombinant

Species Human

Sequence RK ENKFSAKRLQ TTRLGNHLED RVNKFLRRQN

HPEAGEVFVR VVASSDKTVE VKPGMKSRFV
DSGEMSESFP YRTKALFAFE EIDGVDVCFF
GMHVQEYGSD CPPPNTRRVY ISYLDSIHFF

RPRCLRTAVY HEILIGYLEY VKKLGYVTGH IWACPPSEGD DYIFHCHPPD QKIPKPKRLQ EWYKKMLDKA FAERIIHDYK

DIFKQATEDR LTSAKELPYF EGDFWPNVLE ESIKELEQEE EERKKEESTA ASETTEGSQG DSKNAKKKNN KKTNKNKSSI SRANKKKPSM

PNVSNDLSQK LYATMEKHKE VFFVIHLHAG PVINTLPPIV

DPDPLLSCDL MDGRDAFLTL ARDKHWEFSS LRRSKWSTLC MLVELHTQGQ DRFVYTCNEC

Predicted molecular weight 72 kDa including tags

Amino acids 1319 to 1710

Tags proprietary tag N-Terminus

Specifications

Our Abpromise guarantee covers the use of ab167964 in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

Applications Western blot

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Functional Studies

SDS-PAGE

Form Liquid

Preparation and Storage

Stability and Storage Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles.

pH: 7.50

 $Constituents: 0.31\% \ Glutathione, 0.002\% \ PMSF, 0.004\% \ DTT, 0.79\% \ Tris \ HCI, 0.003\% \ EDTA, 0.004\% \ DTT, 0.009\% \ Tris \ HCI, 0.003\% \ EDTA, 0.004\% \ DTT, 0.009\% \ Tris \ HCI, 0.000\% \ EDTA, 0.000\% \ DTT, 0.000\% \ DT$

25% Glycerol (glycerin, glycerine), 0.29% Sodium chloride

This product is an active protein and may elicit a biological response in vivo, handle with caution.

General Info

Function Acetylates histones, giving a specific tag for transcriptional activation. Also acetylates non-histone

proteins, like NCOA3 coactivator. Binds specifically to phosphorylated CREB and enhances its transcriptional activity toward cAMP-responsive genes. Acts as a coactivator of ALX1 in the

presence of EP300.

Involvement in diseaseNote=Chromosomal aberrations involving CREBBP may be a cause of acute myeloid leukemias.

Translocation t(8;16)(p11;p13) with MYST3/MOZ; translocation t(11;16)(q23;p13.3) with MLL/HRX; translocation t(10;16)(q22;p13) with MYST4/MORF. MYST3-CREBBP may induce

leukemia by inhibiting RUNX1-mediated transcription.

Defects in CREBBP are a cause of Rubinstein-Taybi syndrome type 1 (RSTS1) [MIM:180849]. RSTS1 is an autosomal dominant disorder characterized by craniofacial abnormalities, broad thumbs, broad big toes, mental retardation and a propensity for development of malignancies.

Sequence similarities Contains 1 bromo domain.

Contains 1 KIX domain.

Contains 2 TAZ-type zinc fingers. Contains 1 ZZ-type zinc finger.

Domain The KIX domain mediates binding to HIV-1 Tat.

Post-translational Methylation of the KIX domain by CARM1 blocks association with CREB. This results in the

modifications blockade of CREB signaling, and in activation of apoptotic response.

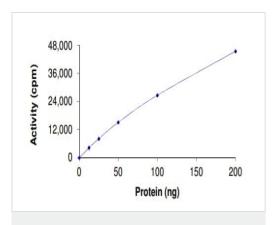
Phosphorylated upon DNA damage, probably by ATM or ATR.

Sumoylation negatively regulates transcriptional activity via the recruitment of DAAX.

Cellular localization Cytoplasm. Nucleus. Recruited to nuclear bodies by SS18L1/CREST. In the presence of ALX1

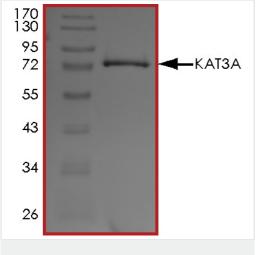
relocalizes from the cytoplasm to the nucleus.

Images



The specific activity of CREBBP (ab167964) was determined to be 33 nmol/min/mg as per activity assay protocol

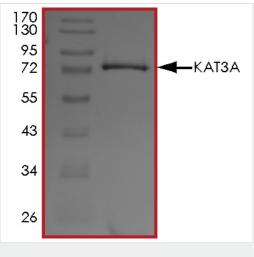




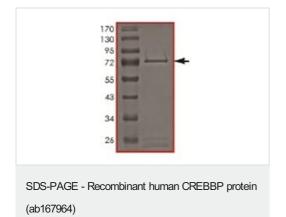
SDS-PAGE - Recombinant human CREBBP protein (ab167964)

SDS PAGE analysis of ab167964

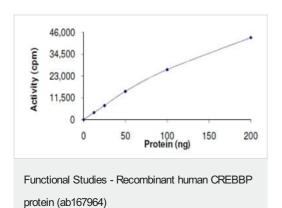




SDS-PAGE - Recombinant human CREBBP protein (ab167964)



SDS Page analysis of ab167964



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