abcam

Product datasheet

Recombinant Human CREBBP protein ab56272

1 图像

描述

产品名称
重组人CREBBP蛋白

纯**度** > 90 % Densitometry.

表达系统 Escherichia coli

蛋白长度 Protein fragment

无动物成分 No

性质 Recombinant

种属 Human

氨基酸 1319 to 1710

技术指标

Our <u>Abpromise guarantee</u> covers the use of ab56272 in the following tested applications.

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

应用 SDS-PAGE

形式 Liquid

制备和贮存

稳定性和存储 Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles.

pH: 7.50

Constituents: 0.00174% PMSF, 0.00385% DTT, 0.79% Tris HCl, 25% Glycerol (glycerin,

glycerine), 0.87% Sodium chloride

常规信息

功能 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.

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疾病相关 Note=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.

序列相似性 Contains 1 bromo domain.

Contains 1 KIX domain.

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

结构域 The KIX domain mediates binding to HIV-1 Tat.

翻译后修饰 Methylation of the KIX domain by CARM1 blocks association with CREB. This results in the

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

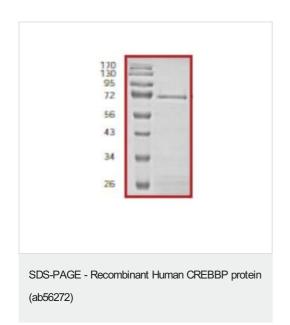
Phosphorylated upon DNA damage, probably by ATM or ATR.

Sumoylation negatively regulates transcriptional activity via the recruitment of DAAX.

细胞定位 Cytoplasm. Nucleus. Recruited to nuclear bodies by SS18L1/CREST. In the presence of ALX1

relocalizes from the cytoplasm to the nucleus.

图片



ab56272 on SDS-PAGE.

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