abcam

Product datasheet

Recombinant Human SP1 protein ab82236

概述

产品名称	重组人SP1蛋白
蛋白长度	Full length protein

描述

性质	Recombinant
来源	Baculovirus

氨基酸序列

种属	Human
序列	MDEMTAVVKI EKGVGGNNGG NGNGGGAFSQ ARSSSTGSSS STGGGGQESQ PSPLALLAAT CSRIESPNEN SNNSQGPSQS GGTGELDLTA TQLSQGANGW QIISSSSGAT PTSKEQSGSS TNGSNGSESS KNRTVSGGQY VVAAAPNLQN QQVLTGLPGV MPNIQYQVIP QFQTVDGQQL QFAATGAQVQ QDGSGQIQII PGANQQIITN RGSGGNIIAA MPNLLQQAVP LQGLANNVLS GQTQYVTNVP VALNGNITLL PVNSVSAATL TPSSQAVTIS SSGSQESGSQ PVTSGTTISS ASLVSSQASS SSFFTNANSY STTTTTSNMG IMNFTTSGSS GTNSQGQTPQ RVSGLQGSDA LNIQQNQTSG GSLQAGQQKE GEQNQQTQQQ QILIQPQLVQ GGQALQALQA APLSGQTFTT QAISQETLQN LQLQAVPNSG PIIIRTPTVG PNGQVSWQTL QLQNLQVQNP QAQTITLAPM QGVSLGQTSS SNTTLTPIAS AASIPAGTVT VNAAQLSSMP GLQTINLSAL GTSGIQVHPI QGLPLAIANA PGDHGAQLGL HGAGGDGIHD DTAGGEEGEN SPDAQPQAGR RTRREACTCP YCKDSEGRGS GDPGKKKQHI CHIQGCGKVY GKTSHLRAHL RWHTGERPFM CTWSYCGKRF TRSDELQRHK RTHTGEKKFA CPECPKRFMR SDHLSKHIKT HQNKKGGPGV ALSVGTLPLD SGAGSEGSGT ATPSALITTN MVAMEAICPE GIARLANSGI NVMQVADLQS INISGNGF

分子量	82 kDa
氨基酸	1 to 785

技术指标

Our Abpromise guarantee covers the use of **ab82236** in the following tested applications.

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

应用	EMSA Western blot Functional Studies SDS-PAGE
纯度	> 90 % SDS-PAGE. ab82236 is greater than 90% homogeneous based on SDS-PAGE analysis, purified by using affinity chromatography and FPLC chromatography.
形式	Liquid
补充说明	1 unit equals 1 nanogram of purified protein. 1 unit is sufficient for a gel mobility shift assay in a 20 µl reaction; 100 units are sufficient for protein-protein interaction assays.

制备和贮存

稳定性和存储	Shipped on dry ice. Upon delivery aliquot and store at -80ºC. Avoid freeze / thaw cycles. Preservative: None Constituents: 20% Glycerol, 20mM Tris Cl, 100mM Potassium chloride, 1mM DTT, 0.2mM EDTA, pH 8.0

常规信息

功能	Transcription factor that can activate or repress transcription in response to physiological and pathological stimuli. Binds with high affinity to GC-rich motifs and regulates the expression of a large number of genes involved in a variety of processes such as cell growth, apoptosis, differentiation and immune responses. Highly regulated by post-translational modifications (phosphorylations, sumoylation, proteolytic cleavage, glycosylation and acetylation). Binds also the PDGFR-alpha G-box promoter. May have a role in modulating the cellular response to DNA damage. Implicated in chromatin remodeling. Plays a role in the recruitment of SMARCA4/BRG1 on the c-FOS promoter. Plays an essential role in the regulation of FE65 gene expression. In complex with ATF7IP, maintains telomerase activity in cancer cells by inducing TERT and TERC gene expression.
组织特异性	Up-regulated in adenocarcinomas of the stomach (at protein level).
序列相似性	Belongs to the Sp1 C2H2-type zinc-finger protein family. Contains 3 C2H2-type zinc fingers.
翻译后修饰	Phosphorylated on multiple serine and threonine residues. Phosphorylation is coupled to ubiquitination, sumoylation and proteolytic processing. Phosphorylation on Ser-59 enhances proteolytic cleavage. Phosphorylation on Ser-7 enhances ubiquitination and protein degradation. Hyperphosphorylation on Ser-101 in response to DNA damage has no effect on transcriptional activity. MAPK1/MAPK3-mediated phosphorylation on Thr-453 and Thr-739 enhances VEGF

	transcription but, represses FGF2-triggered PDGFR-alpha transcription. Also implicated in the repression of RECK by ERBB2. Hyperphosphorylated on Thr-278 and Thr-739 during mitosis by MAPK8 shielding SP1 from degradation by the ubiquitin-dependent pathway. Phosphorylated in the zinc-finger domain by calmodulin-activated PKCzeta. Phosphorylation on Ser-641 by PKCzeta is critical for TSA-activated LHR gene expression through release of its repressor, p107. Phosphorylation on Thr-668, Ser-670 and Thr-681 is stimulated by angiotensin II via the AT1 receptor inducing increased binding to the PDGF-D promoter. This phosphorylation is increased in injured artey wall. Ser-59 and Thr-681 can both be dephosphorylated by PP2A during cell-cycle interphase. Dephosphorylation on Ser-59 leads to increased chromatin association during interphase and increases the transcriptional activity. On insulin stimulation, sequentially glycosylated and phosphorylated on several C-terminal serine and threonine residues. Acetylated. Acetylation/deacetylation events affect transcriptional activity. Deacetylation leads to an increase in the expression the 12(s)-lipooxygenase gene though recruitment of p300 to the promoter. Ubiquitinated. Ubiquitination occurs on the C-terminal proteolytically-cleaved peptide and is triggered by phosphorylation. Sumoylated by SUMO1. Sumoylation modulates proteolytic cleavage of the N-terminal repressor domain. Sumoylation levels are attenuated during tumorigenesis. Phosphorylation mediates SP1 desumoylation. Proteolytic cleavage in the N-terminal repressor domain is prevented by sumoylation. The C-terminal cleaved product is susceptible to degradation. O-glycosylated; contains at least 8 N-acetylglucosamine side chains. Levels are controlled by insulin and the SP1 phosphorylation states. Insulin-mediated O-glycosylation locates SP1 to the nucleus, where it is sequentially deglycosylated and phosphorylated. O-glycosylation affects transcriptional activity through disrupting the interaction with a number of transcription factors including ELF1 and NFYA. Also inhibits interaction with the HIV1 promoter. Inhibited by peroxisomome proliferator receptor gamma (PPARgamma).
细胞定位	Nucleus. Cytoplasm. Nuclear location is governed by glycosylated/phosphorylated states. Insulin promotes nuclear location, while glucagon favors cytoplasmic location.

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