

Rabbit Anti-SP1 antibody

SL0975R

Product Name:	SP1			
Chinese Name:	转录生长因子SP1抗体			
Alias:	Sp1 transcription factor isoform a; TSFP1; TSFP 1; SP1; Specificity protein 1; Transcription factor Sp1; SP1_HUMAN.			
	Specific References(2) SL0975R has been referenced in 2 publications.			
	[IF=4.15]Peng, Yanfei, et al. "Prostaglandin E2 induces Stromal cell-derived factor-1			
	expression in prostate stromal cells by activating Protein kinase A and transcription			
文献引用	factor Sp1." The International Journal of Biochemistry & Cell Biology			
	(2012). WB;Human .			
Pub	PubMed:23246486			
	[IF=2.80] Yang, J. H., et al. "Long-term persistent infection of HPV 16 E6 up-regulate			
	SP1 and hTERT by inhibiting LKB1 in lung cancer cells." PloS one 12.8 (2017):			
	e0182775. WB;Human .			
	PubMed:28813465			
Organism Species:	Rabbit			
Clonality:	Polyclonal			
React Species:	Human,Mouse,Rat,Chicken,Dog,Pig,Cow,Sheep,			
	WB=1:500-2000ELISA=1:500-1000IHC-P=1:400-800IHC-F=1:400-800Flow-			
Applications	Cyt=1ug/TestIF=1:100-500 (Paraffin sections need antigen repair)			
	not yet tested in other applications.			
	optimal dilutions/concentrations should be determined by the end user.			
Molecular weight:	81kDa			
Cellular localization:	The nucleuscytoplasmic			
Form:	Lyophilized or Liquid			
Concentration:	1mg/ml			

immunogen:	KLH conjugated synthetic peptide derived from human TSFP1:701-785/785			
Lsotype:	IgG			
Purification:	affinity purified by Protein A			
Storage Buffer:	0.01M TBS(pH7.4) with 1% BSA, 0.03% Proclin300 and 50% Glycerol.			
Storage:	Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. The lyophilized antibody is stable at room temperature for at least one month and for greater than a year when kept at -20°C. When reconstituted in sterile pH 7.4 0.01M PBS or diluent of antibody the antibody is stable for at least two weeks at 2-4 °C.			
PubMed:	PubMed			
Product Detail:	 Profound changes in patterns of gene expression can result from relatively small changes in the concentrations of sequence specific transcription factors. Synergistic interaction between factors bound to different sites within a transcriptional control region is supported by the work of Courey et al. (1989). Sp1 is a sequence specific transcription factor that recognizes GGGGCGGGG and closely related sequences, which are often referred to as GC boxes. Sp1 binds to GC box promoters elements and selectively activates mRNA synthesis from genes that contain functional recognition sites. SP1 can interact with G/C rich motifs from serotonin receptor promoter. Kadonaga et al. (1987) cloned the human Sp1 cDNA and showed that it has contiguous zinc finger motifs and requires zinc for sequence specific binding to DNA. Altername: Sp1 transcription factor isoform a; TSFP1; TSFP 1; Specificity protein 1; Transcription factor Sp1. Function: 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. 			
	Interacts with ATF7IP, ATF7IP2, BAHD1, POGZ, HCFC1, AATF and PHC2. Interacts with varicella-zoster virus IE62 protein. Interacts with HIV-1 Vpr; the interaction is inhibited by SP1 O-glycosylation. Interacts with SV40 VP2/3 proteins. Interacts with SV40 major capsid protein VP1; this interaction leads to a cooperativity between the 2 proteins in DNA binding. Interacts with HLTF; the interaction may be required for basal transcriptional activity of HLTF. Interacts (deacetylated form) with EP300; the interaction enhances gene expression. Interacts with HDAC1 and JUN. Interacts with ELF1; the interaction is inhibited by glycosylation of SP1. Interaction with NFYA; the interaction is inhibited by glycosylation of SP1. Interacts with SMARCA4/BRG1. Interacts with ATF7IP and TBP. Interacts with MEIS2 isoform 4 and PBX1 isoform PBX1a.			

Subcellular Location:

Nucleus. Cytoplasm. Nuclear location is governed by glycosylated/phosphorylated states. Insulin promotes nuclear location, while glucagon favors cytoplasmic location.

Tissue Specificity:

Up-regulated in adenocarcinomas of the stomach (at protein level).

Post-translational modifications:

Phosphorylated on multiple serine and threonine residues. Phosphorylation is coupled t	0
ubiguitination, sumovation 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	3
no effect on transcriptional activity MAPK1/MAPK3-mediated phosphorylation on Th	r-
453 and Thr-739 enhances VEGE transcription but represses EGE2-triggered PDGER-	1
alpha transcription Also implicated in the repression of RECK by ERBB?	
Hyperphosphorylated on Thr 278 and Thr 730 during mitosis by MADK8 shielding SP	1
from degradation by the ubiquitin dependent nathway. Phosphorylated in the zinc finge	1 ar
domain by calmodulin activated PKCzota, Phosphorylation on Sor 641 by PKCzota is	Л
abiliant by cannodumi-activated FKCzeta. Phosphorylation on Ser-041 by FKCzeta is	
critical for TSA-activated LHR gene expression infough release of its repressor, p10/.	_
Phosphorylation on Thr-608, Ser-670 and Thr-681 is stimulated by anglotensin II via th	le
A I I 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	
rephosphorylated by PP2A during cell-cycle interphase. Dephosphorylation on Ser-59	
eads to increased chromatin association during interphase and increases the	
ranscriptional 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 with 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 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).	

Similarity:

	Belongs to the Sp1 C2H2-type zinc-finger protein family.		
	Contains 3 C2H2-type zinc tingers.		
	SWISS:		
	P08047		
	Gene ID:		
	6667		
	Database links:		
	Entrez Gene: 6667Human		
	Entrez Gene: 395303Chicken		
	Entrez Gene: 540741Cow		
	Entrez Gene: 20683 Mouse		
	Entrez Gene: 397314Pig		
	Entrez Gene: 24790Rat		
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	Important Note:		
	This product as supplied is intended for research use only, not for use in human,		
	therapeutic or diagnostic applications.		
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	1.	25AP	
	1	80 —	
	S 1	35	
	1	00 —	
Picture:	Mag	75 —	
		63 —	
		48 —	

Sample:

A549(Human) Cell Lysate at 30 ug

Primary: Anti-SP1 (SL0975R) at 1/300 dilution

Secondary: IRDye800CW Goat Anti-Rabbit IgG at 1/20000 dilution

Predicted band size: 81 kD

Observed band size: 111 kD



Paraformaldehyde-fixed, paraffin embedded (Human stomach cancer); Antigen retrieval by boiling in sodium citrate buffer (pH6.0) for 15min; Block endogenous peroxidase by 3% hydrogen peroxide for 20 minutes; Blocking buffer (normal goat serum) at 37°C for 30min; Antibody incubation with (TSFP1) Polyclonal Antibody, Unconjugated (SL0975R) at 1:200 overnight at 4°C, followed by operating according to SP Kit(Rabbit) (sp-0023) instructionsand DAB staining.

