

Rabbit Anti-Phospho-HDAC3 (Ser424) antibody

SL3174R

Product Name:	Phospho-HDAC3 (Ser424)
Chinese Name:	磷酸化组蛋白去乙酰化酶3抗体
Alias:	HDAC3 (phospho S424); HDAC3 (phospho Ser424); p-HDAC3 (Ser424); RPD3 2; RPD3-2; HD 3; HD3; HDAC 3; Histone deacetylase 3 (HD3) (RPD3-2); histone deacetylase 3; RPD 3; RPD3; SMAP 45; SMAP-45; SMAP45; HDAC3_HUMAN.
Organism Species:	Rabbit
Clonality:	Polyclonal
React Species:	Human, Mouse, Rat,
Applications:	WB=1:500-2000ELISA=1:500-1000IHC-P=1:400-800IHC-F=1:400-800IF=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:	50kDa
Cellular localization:	The nucleuscytoplasmic
Form:	Lyophilized or Liquid
Concentration:	1mg/ml
immunogen:	KLH conjugated Synthesised phosphopeptide derived from human HDAC3 around the phosphorylation site of Ser424:KE(p-S)DV
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:	Histones play a critical role in transcriptional regulation, cell cycle progression, and
	developmental events. Histone acetylation/deacetylation alters chromosome structure
	and affects transcription factor access to DNA. The protein encoded by this gene belongs

to the histone deacetylase/acuc/apha family. It has histone deacetylase activity and represses transcription when tethered to a promoter. It may participate in the regulation of transcription through its binding with the zinc-finger transcription factor YY1. This protein can also down-regulate p53 function and thus modulate cell growth and apoptosis. This gene is regarded as a potential tumor suppressor gene. [provided by RefSeq, Jul 2008].

Function:

Responsible for the deacetylation of lysine residues on the N-terminal part of the core histones (H2A, H2B, H3 and H4), and some other non-histone substrates. Histone deacetylation gives a tag for epigenetic repression and plays an important role in transcriptional regulation, cell cycle progression and developmental events. Histone deacetylases act via the formation of large multiprotein complexes. Probably participates in the regulation of transcription through its binding to the zinc-finger transcription factor YY1; increases YY1 repression activity. Required to repress transcription of the POU1F1 transcription factor. Acts as a molecular chaperone for shuttling phosphorylated NR2C1 to PML bodies for sumoylation.

Subunit:

Interacts with HDAC7 and HDAC9. Forms a heterologous complex at least with YY1. Interacts with DAXX, HDAC10 and DACH1. Found in a complex with NCOR1 and NCOR2. Component of the N-Cor repressor complex, at least composed of NCOR1, NCOR2, HDAC3, TBL1X, TBL1R, CORO2A and GPS2. Interacts with BCOR, MJD2A/JHDM3A, NRIP1, PRDM6 and SRY. Interacts with BTBD14B. Interacts with GLIS2. Interacts (via the DNA-binding domain) with NR2C1; the interaction recruits phosphorylated NR2C1 to PML bodies for sumoylation. Component of the Notch corepressor complex. Interacts with CBFA2T3 and NKAP. Interacts with APEX1; the interaction is not dependent on the acetylated status of APEX1. Interacts with and deacetylates MAPK14. Interacts with ZMYND15.

Subcellular Location: Nucleus.

Tissue Specificity: Widely expressed.

Post-translational modifications: Sumoylated in vitro.

Similarity:

Belongs to the histone deacetylase family. HD type 1 subfamily.

SWISS: 015379

Gene ID:

8841

Database links:

Entrez Gene: 8841Human

Entrez Gene: 15183Mouse

Entrez Gene: 84578Rat

Omim: 605166Human

SwissProt: O15379Human

SwissProt: 088895Mouse

SwissProt: Q6P6W3Rat

Unigene: 519632Human

Unigene: 20521Mouse

Unigene: 17284Rat

Important Note:

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