

Rabbit Anti-phospho-HDAC2 (Ser394) antibody

SL5389R

Product Name:	phospho-HDAC2 (Ser394)
Chinese Name:	磷酸化组蛋白去乙酰化酶2抗体
Alias:	HDAC2 (Phospho-Ser394); HDAC2 (phospho S394); HDAC2 (phospho Ser394); p- HDAC2 (Ser394); histone deacetylase 2; D10Wsu179e; EC 3.5.1.98; HD2; HDAC 2; Hdac2; histone deacetylase 2; RPD3; transcriptional regulator homolog RPD3; YAF1; YY1 associated factor 1; YY1 transcription factor binding protein; Yy1bp; HDAC2_HUMAN.
	Specific References(1) SL5389R has been referenced in 1 publications.
文献引用	[IF=3.81] Jou, Yu-Jen, et al. "Quantitative phosphoproteomic analysis reveals
Pub	γ -bisabolene inducing p53-mediated apoptosis of human oral squamous cell carcinoma
:	via HDAC2 inhibition and ERK1/2 activation." Proteomics (2015).WB;Human.
	PubMed:26194454
Organism Species:	Rabbit
Clonality:	Polyclonal
React Species:	Human,Mouse,Rat,Chicken,Dog,Pig,Cow,Horse,
Applications:	ELISA=1:500-1000IHC-P=1:400-800IHC-F=1:400-800ICC=1:100-500IF=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:	55kDa
Cellular localization:	The nucleus
Form:	Lyophilized or Liquid
Concentration:	1mg/ml
immunogen:	KLH conjugated Synthesised phosphopeptide derived from human HDAC2 around the phosphorylation site of Ser394:ED(p-S)GD
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:	 This gene product belongs to the histone deacetylase family. Histone deacetylases act via the formation of large multiprotein complexes, and are responsible for the deacetylation of lysine residues at the N-terminal regions of core histones (H2A, H2B, H3 and H4). This protein forms transcriptional repressor complexes by associating with many different proteins, including YY1, a mammalian zine-finger transcription factor. Thus, it plays an important role in transcriptional regulation, cell cycle progression and developmental events. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Apr 2010]. Function: Responsible for the deacetylation of lysine residues on the N-terminal part of the core histones (H2A, H2B, H3 and H4). 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. Forms transcriptional repressor complexes by associating with MAD, SIN3, YY1 and N-COR. Interacts in the late S-phase of DNA-replication with DNMT1 in the other transcriptional repressor complex composed of DNMT1, DMAP1, PCNA, CAF1. Deacetylates TSHZ3 and regulates its transcriptional repressor activity. Component of a RCOR/GFI/KDM1A/HDAC complex. Interacts directly with GFI1 and GFI1B. Interacts with SNW1, HDAC7, PRDM6, SAP30, SETDB1 and SUV39H1. Interacts with bRV1, with COR complex composed of HDAC1, HDAC2, RBBP4 and RBBP7. The core complex associates with MTA2, MBD3, MTA1L1, CHD3 and CHD4 to form the nucleosome remodeling and histone deacetylation (NuRD) complex, or with SIN3, SAP18 and SAP30 to form the SIN3 HDAC complex. Component of a BHC complex may also contain ZMYM2, ZNF217, ZMYM3, GSE1 and GT21. Part of a complex containing the core histones H2A, H2B, H3 and H4, DEX and unphosphorylated DAXX. Part of a complex contaning ATR and CHD4. Forms a heterologous complex at least with

with APEX1; the interaction is not dependent on the acetylated status of APEX1. Part of a complex composed of TRIM28, HDAC1, HDAC2 and EHMT2. Subcellular Location: Nucleus. **Tissue Specificity:** Widely expressed; lower levels in brain and lung. **Post-translational modifications:** S-nitrosylated by GAPDH. In neurons, S-Nitrosylation at Cys-262 and Cys-274 does not affect the enzyme activity but abolishes chromatin-binding, leading to increases acetylation of histones and activate genes that are associated with neuronal development. In embryonic cortical neurons, S-Nitrosylation regulates dendritic growth and branching. Similarity: Belongs to the histone deacetylase family. HD type 1 subfamily. **SWISS:** 092769 Gene ID: 3066 Database links: Entrez Gene: 3066Human Entrez Gene: 15182Mouse Entrez Gene: 84577Rat Omim: 605164Human SwissProt: Q92769Human SwissProt: P70288Mouse Unigene: 3352Human Unigene: 19806Mouse Unigene: 1797Rat Important Note: This product as supplied is intended for research use only, not for use in human, therapeutic or diagnostic applications.

