

# **Anti-HDAC2 Antibody**

**Catalog # ABO10674** 

### **Specification**

## **Anti-HDAC2 Antibody - Product Information**

Application
Primary Accession
Host
Reactivity
Clonality
Format
WB, IHC
092769
Rabbit
Human, Rat
Polyclonal
Lyophilized

**Description** 

Rabbit IgG polyclonal antibody for Histone deacetylase 2(HDAC2) detection. Tested with WB, IHC-P in Human;Rat.

#### Reconstitution

Add 0.2ml of distilled water will yield a concentration of 500ug/ml.

# **Anti-HDAC2 Antibody - Additional Information**

**Gene ID 3066** 

#### **Other Names**

Histone deacetylase 2, HD2, 3.5.1.98, HDAC2

# **Calculated MW**

55364 MW KDa

#### **Application Details**

Immunohistochemistry(Paraffin-embedded Section), 0.5-1  $\mu$ g/ml, Human, Rat, Mouse, By Heat<br/>br>Western blot, 0.1-0.5  $\mu$ g/ml, Human, Mouse, Rat<br/>br>

#### **Subcellular Localization**

Nucleus . Cytoplasm .

# **Tissue Specificity**

Widely expressed; lower levels in brain and lung.

### **Protein Name**

Histone deacetylase 2(HD2)

#### Contents

Each vial contains 5mg BSA, 0.9mg NaCl, 0.2mg Na2HPO4, 0.05mg Thimerosal, 0.05mg NaN3.

## **Immunogen**

A synthetic peptide corresponding to a sequence at the C-terminus of human HDAC2(462-478aa KEEDKSKDNSGEKTDTK), identical to the related rat sequence, and different from the related mouse sequence by one amino acid.





Purification Immunogen affinity purified.

**Cross Reactivity**No cross reactivity with other proteins

Storage

At -20°C for one year. After r°Constitution, at 4°C for one month. It°Can also be aliquotted and stored frozen at -20°C for a longer time. Avoid repeated freezing and thawing.

#### **Sequence Similarities**

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

# **Anti-HDAC2 Antibody - Protein Information**

Name HDAC2 {ECO:0000303|PubMed:10545197, ECO:0000312|HGNC:HGNC:4853}

#### **Function**

Histone deacetylase that catalyzes the deacetylation of lysine residues on the N-terminal part of the core histones (H2A, H2B, H3 and H4) (PubMed:<a href="http://www.uniprot.org/citations/28497810" target="\_blank">28497810</a>). Histone deacetylation gives a tag for epigenetic repression and plays an important role in transcriptional

deacetylation gives a tag for epigenetic repression and plays an important role in transcriptional regulation, cell cycle progression and developmental events (By similarity). Histone deacetylases act via the formation of large multiprotein complexes (By similarity). Forms transcriptional repressor complexes by associating with MAD, SIN3, YY1 and N-COR (PubMed: <a href="http://www.uniprot.org/citations/12724404" target=" blank">12724404</a>). Component of a RCOR/GFI/KDM1A/HDAC complex that suppresses, via histone deacetylase (HDAC) recruitment, a number of genes implicated in multilineage blood cell development (By similarity). Acts as a component of the histone deacetylase NuRD complex which participates in the remodeling of chromatin (PubMed:<a href="http://www.uniprot.org/citations/16428440" target=" blank">16428440</a>, PubMed:<a href="http://www.uniprot.org/citations/28977666" target="blank">28977666</a>). Component of the SIN3B complex that represses transcription and counteracts the histone acetyltransferase activity of EP300 through the recognition H3K27ac marks by PHF12 and the activity of the histone deacetylase HDAC2 (PubMed:<a href="http://www.uniprot.org/citations/37137925" target="\_blank">37137925</a>). Also deacetylates non-histone targets: deacetylates TSHZ3, thereby regulating its transcriptional repressor activity (PubMed: <a href="http://www.uniprot.org/citations/19343227" target=" blank">19343227</a>). May be involved in the transcriptional repression of circadian target genes, such as PER1, mediated by CRY1 through histone deacetylation (By similarity). Involved in MTA1-mediated transcriptional corepression of TFF1 and CDKN1A (PubMed:<a

Involved in MTA1-mediated transcriptional corepression of TFF1 and CDKN1A (PubMed:<a href="http://www.uniprot.org/citations/21965678" target="\_blank">21965678</a>). In addition to protein deacetylase activity, also acts as a protein-lysine deacylase by recognizing other acyl groups: catalyzes removal of (2E)-butenoyl (crotonyl) and 2- hydroxyisobutyryl) acyl groups from lysine residues, leading to protein decrotonylation and de-2- hydroxyisobutyrylation, respectively (PubMed:<a

 $href="http://www.uniprot.org/citations/28497810" target="\_blank">28497810</a>, PubMed:<a href="http://www.uniprot.org/citations/29192674" target="\_blank">29192674</a>).$ 

**Cellular Location** Nucleus. Cytoplasm

### **Tissue Location**

Widely expressed; lower levels in brain and lung.

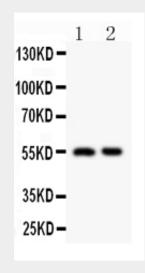


# **Anti-HDAC2 Antibody - Protocols**

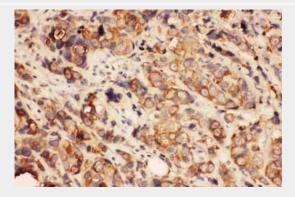
Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- <u>Immunohistochemistry</u>
- Immunofluorescence
- Immunoprecipitation
- Flow Cytomety
- Cell Culture

## **Anti-HDAC2 Antibody - Images**



Anti-HDAC2 antibody, ABO10674, Western blottingAll lanes: Anti HDAC2 (ABO10674) at 0.5ug/mlLane 1: U87 Whole Cell Lysate at 40ugLane 2: A549 Whole Cell Lysate at 40ugPredicted bind size: 55KDObserved bind size: 55KD



Anti-HDAC2 antibody, ABO10674, IHC(P)IHC(P): Human Mammary Cancer Tissue

# Anti-HDAC2 Antibody - Background

Histone deacetylase 2 is an enzyme that in humans is encoded by the HDAC2 gene. 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 on the N-terminal region of the core histones(H2A, H2B, H3 and H4). This protein also forms transcriptional





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repressor complexes by associating with many different proteins, including YY1, a mammalian zinc-finger transcription factor. Thus it plays an important role in transcriptional regulation, cell cycle progression and developmental events. Betz et al.(1998) performed PCR using HDAC2-specific primers to screen a somatic cell hybrid mapping panel. They mapped the HDAC2 gene to human chromosome 6q21, a region of the genome altered in some cancers, including retinoblastoma.