

## **HDAC3 Monoclonal Antibody**

Catalog No: YM1040

Reactivity: Human; Mouse; Rat; Dog; Pig; Rabbit

**Applications:** WB

Target: HDAC3

**Fields:** >>Neutrophil extracellular trap formation;>>Thyroid hormone signaling

pathway;>>Alcoholism;>>Viral carcinogenesis

Gene Name: HDAC3

**Protein Name:** Histone deacetylase 3

O88895

Human Gene Id: 8841

**Human Swiss Prot** 015379

No:

**Mouse Swiss Prot** 

No:

Rat Gene Id: 84578

Rat Swiss Prot No: Q6P6W3

**Immunogen:** Purified recombinant human HDAC3 protein fragments expressed in E.coli.

**Specificity:** HDAC3 Monoclonal Antibody detects endogenous levels of HDAC3 protein.

**Formulation :** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Source: Monoclonal, Mouse

**Dilution:** WB 1:1000 - 1:2000. Not yet tested in other applications.

Purification : Affinity purification

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**Concentration**: 1 mg/ml

**Storage Stability:** -15°C to -25°C/1 year(Do not lower than -25°C)

Molecularweight: 49kD

**Cell Pathway :** Protein\_Acetylation

**Background:** 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 zincfinger 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 :** catalytic activity:Hydrolysis of an N(6)-acetyl-lysine residue of a histone to yield a

deacetylated histone.,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. 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.,PTM:Sumoylated in

vitro., similarity: Belongs to the histone deacetylase family. Type 1

subfamily..subunit:Interacts with HDAC7 and HDAC9. Forms a heterologous

complex at least with YY1. Intera

Subcellular Location:

Nucleus . Cytoplasm . Cytoplasm, cytosol . Colocalizes with XBP1 and AKT1 in the cytoplasm (PubMed:25190803). Predominantly expressed in the nucleus in

the presence of CCAR2 (PubMed:21030595). .

**Expression:** Widely expressed.

**Sort**: 7282

**No4**: 1

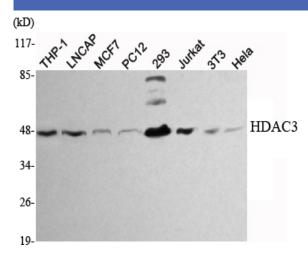
Host: Mouse

Modifications: Unmodified

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## **Products Images**



Western Blot analysis using HDAC3 Monoclonal Antibody against THP-1, LNCAP, MCF7, PC12, 293, Jurkat, 3T3, HeLa cell lysate.