

# **HDAC6** Antibody

Rabbit mAb Catalog # AP90889

## **Specification**

# **HDAC6 Antibody - Product Information**

Application WB, IHC, ICC, IP

Primary Accession
Clonality

O9UBN7

Monoclonal

**Other Names** 

HD 6; HDAC 6; Histone deacetylase 6 (HD6); JM 21;

Isotype Rabbit IgG
Host Rabbit
Calculated MW 131419 Da

## **HDAC6 Antibody - Additional Information**

Dilution WB~~1:1000

IHC~~1:100~500

ICC~~N/A IP~~N/A

Purification Affinity-chromatography

Immunogen A synthesized peptide derived from human

HDAC6

Description Involved in the regulation of many cellular

processes, including cell migration,

immune synapse formation, viral infection, and degradation of misfolded proteins. HDAC6 binds to both poly-ubiquitinated misfolded proteins and dynein motors, facilitating the transport of misfolded proteins to the aggresome. Required for subsequent recruitment of the autophagic machinery and clearance of aggresomes from the cell. Plays a key role in the protection against the deleterious effects of pathological protein aggregation that

occurs in various diseases.

Storage Condition and Buffer Rabbit IgG in phosphate buffered saline,

pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid

freeze / thaw cycle.

# **HDAC6 Antibody - Protein Information**

Name HDAC6 {ECO:0000303|PubMed:10220385, ECO:0000312|HGNC:HGNC:14064}



#### **Function**

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Deacetylates a wide range of non-histone substrates (PubMed:<a
href="http://www.uniprot.org/citations/12024216" target=" blank">12024216</a>, PubMed:<a
href="http://www.uniprot.org/citations/18606987" target="_blank">18606987</a>, PubMed:<a
href="http://www.uniprot.org/citations/20308065" target="blank">20308065</a>, PubMed:<a
href="http://www.uniprot.org/citations/24882211" target="blank">24882211</a>, PubMed:<a
href="http://www.uniprot.org/citations/26246421" target=" blank">26246421</a>, PubMed:<a
href="http://www.uniprot.org/citations/30538141" target="blank">30538141</a>, PubMed:<a
href="http://www.uniprot.org/citations/31857589" target="blank">31857589</a>, PubMed:<a
href="http://www.uniprot.org/citations/30770470" target="_blank">30770470</a>, PubMed:<a
href="http://www.uniprot.org/citations/38534334" target="blank">38534334</a>, PubMed:<a
href="http://www.uniprot.org/citations/39567688" target=" blank">39567688</a>). Plays a
central role in microtubule- dependent cell motility by mediating deacetylation of tubulin
(PubMed:<a href="http://www.uniprot.org/citations/12024216" target=" blank">12024216</a>,
PubMed:<a href="http://www.uniprot.org/citations/20308065" target=" blank">20308065</a>,
PubMed:<a href="http://www.uniprot.org/citations/26246421" target="_blank">26246421</a>).
Required for cilia disassembly via deacetylation of alpha-tubulin (PubMed: <a
href="http://www.uniprot.org/citations/17604723" target="_blank">17604723</a>, PubMed:<a
href="http://www.uniprot.org/citations/26246421" target=" blank">26246421</a>). Alpha-tubulin
deacetylation results in destabilization of dynamic microtubules (By similarity). Promotes
deacetylation of CTTN, leading to actin polymerization, promotion of autophagosome-lysosome
fusion and completion of autophagy (PubMed: <a
href="http://www.uniprot.org/citations/30538141" target=" blank">30538141</a>). Deacetylates
SQSTM1 (PubMed:<a href="http://www.uniprot.org/citations/31857589"
target=" blank">31857589</a>). Deacetylates peroxiredoxins PRDX1 and PRDX2, decreasing
their reducing activity (PubMed: <a href="http://www.uniprot.org/citations/18606987"
target=" blank">18606987</a>). Deacetylates antiviral protein RIGI in the presence of viral
mRNAs which is required for viral RNA detection by RIGI (By similarity). Sequentially deacetylates
and polyubiquitinates DNA mismatch repair protein MSH2 which leads to MSH2 degradation,
reducing cellular sensitivity to DNA-damaging agents and decreasing cellular DNA mismatch repair
activities (PubMed:<a href="http://www.uniprot.org/citations/24882211"
target=" blank">24882211</a>). Deacetylates DNA mismatch repair protein MLH1 which
prevents recruitment of the MutL alpha complex (formed by the MLH1-PMS2 heterodimer) to the
MutS alpha complex (formed by the MSH2-MSH6 heterodimer), leading to tolerance of DNA
damage (PubMed: <a href="http://www.uniprot.org/citations/30770470"
target=" blank">30770470</a>). Deacetylates RHOT1/MIRO1 which blocks mitochondrial
transport and mediates axon growth inhibition (By similarity). Deacetylates transcription factor
SP1 which leads to increased expression of ENG, positively regulating angiogenesis (PubMed: <a
href="http://www.uniprot.org/citations/38534334" target="_blank">38534334</a>). Deacetylates
KHDRBS1/SAM68 which regulates alternative splicing by inhibiting the inclusion of CD44 alternate
exons (PubMed: <a href="http://www.uniprot.org/citations/26080397"
target=" blank">26080397</a>). Acts as a valine sensor by binding to valine through the
primate-specific SE14 repeat region (PubMed:<a
href="http://www.uniprot.org/citations/39567688" target=" blank">39567688</a>). In valine
deprivation conditions, translocates from the cytoplasm to the nucleus where it deacetylates TET2
which promotes TET2-dependent DNA demethylation, leading to DNA damage (PubMed: <a
href="http://www.uniprot.org/citations/39567688" target="_blank">39567688</a>). Promotes
odontoblast differentiation following IPO7-mediated nuclear import and subsequent repression of
RUNX2 expression (By similarity). In addition to its protein deacetylase activity, plays a key role in
the degradation of misfolded proteins: when misfolded proteins are too abundant to be degraded
by the chaperone refolding system and the ubiquitin-proteasome, mediates the transport of
misfolded proteins to a cytoplasmic juxtanuclear structure called aggresome (PubMed:<a
href="http://www.uniprot.org/citations/17846173" target=" blank">17846173</a>). Probably
acts as an adapter that recognizes polyubiquitinated misfolded proteins and targets them to the
aggresome, facilitating their clearance by autophagy (PubMed: <a
href="http://www.uniprot.org/citations/17846173" target=" blank">17846173</a>). Involved in
the MTA1-mediated epigenetic regulation of ESR1 expression in breast cancer (PubMed: <a
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href="http://www.uniprot.org/citations/24413532" target="\_blank">24413532</a>).

#### **Cellular Location**

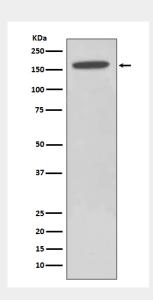
Cytoplasm. Cytoplasm, cytoskeleton. Nucleus. Perikaryon {ECO:0000250|UniProtKB:Q9Z2V5}. Cell projection, dendrite {ECO:0000250|UniProtKB:Q9Z2V5}. Cell projection, axon {ECO:0000250|UniProtKB:Q9Z2V5}. Cell projection, cilium. Cytoplasm, cytoskeleton, microtubule organizing center, centrosome. Cytoplasm, cytoskeleton, cilium basal body Note=Mainly cytoplasmic where it is associated with microtubules (PubMed:12024216). Can shuttle between the cytoplasm and the nucleus (PubMed:39567688). Retained in the cytoplasm by binding to valine via the primate-specific SE14 repeat region while valine deprivation induces nuclear localization (PubMed:39567688). Found exclusively in the cytoplasm in proliferative cells with a fraction found in the nucleus during differentiation (By similarity). May translocate to the nucleus following DNA damage (PubMed:30770470) {ECO:0000250|UniProtKB:Q9Z2V5, ECO:0000269|PubMed:12024216, ECO:0000269|PubMed:30770470, ECO:0000269|PubMed:39567688}

### **HDAC6 Antibody - Protocols**

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

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

### **HDAC6 Antibody - Images**



Western blot analysis of HDAC6 expression in HeLa cell lysate.