

**HDAC6 Rabbit mAb**  
Catalog # AP78985**Specification****HDAC6 Rabbit mAb - Product Information**

Application	<b>WB</b>
Primary Accession	<a href="#">Q9UBN7</a>
Reactivity	<b>Human</b>
Host	<b>Rabbit</b>
Clonality	<b>Monoclonal Antibody</b>
Calculated MW	<b>131419</b>

**HDAC6 Rabbit mAb - Additional Information****Gene ID** 10013**Other Names**  
HDAC6**Dilution**  
WB~~1/500-1/1000**Format**  
Liquid**HDAC6 Rabbit mAb - Protein Information****Name** HDAC6 {ECO:0000303|PubMed:10220385, ECO:0000312|HGNC:HGNC:14064}**Function**

Responsible for the deacetylation of lysine residues on the N-terminal part of the core histones (H2A, H2B, H3 and H4) (PubMed: [10220385](http://www.uniprot.org/citations/10220385)). Histone deacetylation gives a tag for epigenetic repression and plays an important role in transcriptional regulation, cell cycle progression and developmental events (PubMed: [10220385](http://www.uniprot.org/citations/10220385)). Histone deacetylases act via the formation of large multiprotein complexes (PubMed: [10220385](http://www.uniprot.org/citations/10220385)). In addition to histones, deacetylates other proteins, such as CTTN, tubulin and SQSTM1 (PubMed: [12024216](http://www.uniprot.org/citations/12024216), PubMed: [20308065](http://www.uniprot.org/citations/20308065), PubMed: [26246421](http://www.uniprot.org/citations/26246421), PubMed: [30538141](http://www.uniprot.org/citations/30538141), PubMed: [31857589](http://www.uniprot.org/citations/31857589)). Plays a central role in microtubule-dependent cell motility by mediating deacetylation of tubulin (PubMed: [12024216](http://www.uniprot.org/citations/12024216), PubMed: [20308065](http://www.uniprot.org/citations/20308065), PubMed: [26246421](http://www.uniprot.org/citations/26246421)). 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>). 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>). Involved in the MTA1-mediated epigenetic regulation of ESR1 expression in breast cancer (PubMed:<a href="http://www.uniprot.org/citations/24413532" target="\_blank">24413532</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 juxtannuclear 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 target them to the aggresome, facilitating their clearance by autophagy (PubMed:<a href="http://www.uniprot.org/citations/17846173" target="\_blank">17846173</a>).

#### Cellular Location

Cytoplasm. Cytoplasm, cytoskeleton. Nucleus {ECO:0000250|UniProtKB:Q9Z2V5}. 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=It is mainly cytoplasmic, where it is associated with microtubules

#### HDAC6 Rabbit mAb - Protocols

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

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

#### HDAC6 Rabbit mAb - Images



