

**HDAC6 Antibody (N-term) Blocking Peptide**  
**Synthetic peptide**  
**Catalog # BP1106b****Specification**

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**HDAC6 Antibody (N-term) Blocking Peptide - Product Information**Primary Accession  
Other Accession[O9UBN7](#)  
[NP\\_006035](#)**HDAC6 Antibody (N-term) Blocking Peptide - Additional Information****Gene ID** 10013**Other Names**

Histone deacetylase 6, HD6, HDAC6, KIAA0901

**Target/Specificity**

The synthetic peptide sequence used to generate the antibody [AP1106b](/product/products/AP1106b) was selected from the N-term region of human HDAC6. A 10 to 100 fold molar excess to antibody is recommended. Precise conditions should be optimized for a particular assay.

**Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

**HDAC6 Antibody (N-term) Blocking Peptide - 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)

target="\_blank">26246421</a>, PubMed:<a href="http://www.uniprot.org/citations/31857589" target="\_blank">31857589</a>, PubMed:<a href="http://www.uniprot.org/citations/30538141" target="\_blank">30538141</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>). 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 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 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 Antibody (N-term) Blocking Peptide - Protocols

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

- [Blocking Peptides](#)

#### HDAC6 Antibody (N-term) Blocking Peptide - Images

#### HDAC6 Antibody (N-term) Blocking Peptide - Background

HDAC6 (histone deacetylase 6) is 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. HDAC6 plays a central role in microtubule-dependent cell motility via deacetylation of tubulin, and has been shown to interact with HDAC11, SIRT2, and F-actin. HDAC6 is ubiquitinated, but its polyubiquitination however does not lead to degradation. HDAC is also a potential target of sumoylation.

#### HDAC6 Antibody (N-term) Blocking Peptide - References

North, B.J., et al., Mol. Cell 11(2):437-444 (2003).Hubbert, C., et al., Nature 417(6887):455-458 (2002).Gao, L., et al., J. Biol. Chem. 277(28):25748-25755 (2002).Hook, S.S., et al., Proc. Natl. Acad. Sci. U.S.A. 99(21):13425-13430 (2002).Kirsh, O., et al., EMBO J. 21(11):2682-2691 (2002).