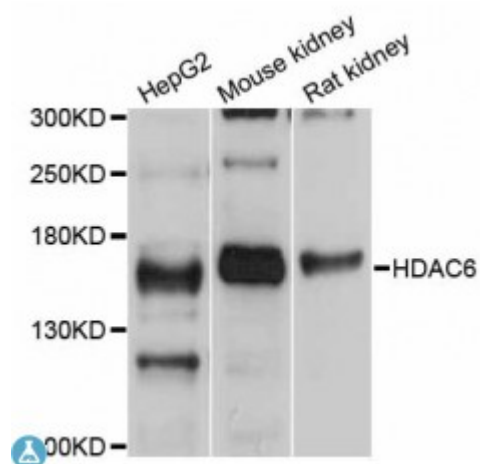


## Anti-HDAC6 Antibody



### Description

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 class II of the histone deacetylase/acuc/apha family. It contains an internal duplication of two catalytic domains which appear to function independently of each other. This protein possesses histone deacetylase activity and represses transcription.

<b>Model</b>	STJ113748
<b>Host</b>	Rabbit
<b>Reactivity</b>	Human, Mouse, Rat
<b>Applications</b>	WB
<b>Immunogen</b>	Recombinant fusion protein containing a sequence corresponding to amino acids 836-1104 of human HDAC6 (NP_006035.2).
<b>Gene ID</b>	<a href="#">10013</a>
<b>Gene Symbol</b>	<a href="#">HDAC6</a>
<b>Dilution range</b>	WB 1:500 - 1:2000
<b>Purification</b>	Affinity purification
<b>Note</b>	For Research Use Only (RUO).
<b>Protein Name</b>	Histone deacetylase 6 HD6
<b>Molecular Weight</b>	131.419 kDa

<b>Clonality</b>	Polyclonal
<b>Conjugation</b>	Unconjugated
<b>Isotype</b>	IgG
<b>Formulation</b>	PBS with 0.02% sodium azide, 50% glycerol, pH7.3.
<b>Storage Instruction</b>	Store at -20C. Avoid freeze / thaw cycles.
<b>Database Links</b>	<a href="#">HGNC:14064OMIM:300272Reactome:R-HSA-2122947</a>
<b>Alternative Names</b>	Histone deacetylase 6 HD6
<b>Function</b>	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 , Plays a central role in microtubule-dependent cell motility via deacetylation of tubulin, Involved in the MTA1-mediated epigenetic regulation of ESR1 expression in breast cancer,
<b>Cellular Localization</b>	Nucleus, Cytoplasm, Perikaryon,
<b>Post-translational Modifications</b>	Phosphorylated by AURKA,