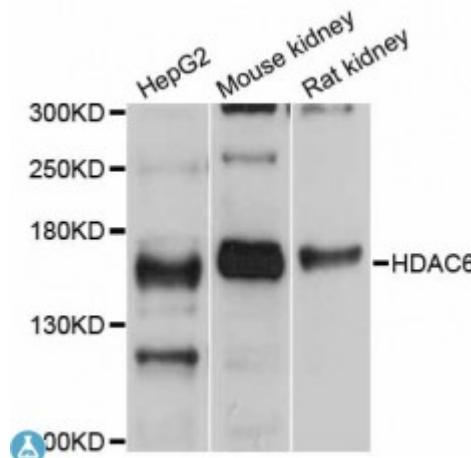




## Anti-HDAC6 Antibody



### 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:14064</a> <a href="#">OMIM:300272</a> <a href="#">Reactome: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,  |

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