

# BRD4 Rabbit mAb

Catalog # AP78960

### **Product Information**

**Application** WB, IHC-P, IP, ICC

060885 **Primary Accession** 

Reactivity Human, Mouse, Rat

Host

Clonality Monoclonal Antibody

Calculated MW 152219

### **Additional Information**

Gene ID 23476

**Other Names** BRD4

**Dilution** WB~~1/500-1/1000 IHC-P~~N/A IP~~N/A ICC~~N/A

**Format** Liquid

#### **Protein Information**

BRD4 Name

**Synonyms** HUNK1

**Function** Chromatin reader protein that recognizes and binds acetylated histones and

> plays a key role in transmission of epigenetic memory across cell divisions and transcription regulation (PubMed:20871596, PubMed:23086925, PubMed:23317504, PubMed:29176719, PubMed:29379197). Remains associated with acetylated chromatin throughout the entire cell cycle and provides epigenetic memory for postmitotic G1 gene transcription by preserving acetylated chromatin status and maintaining high-order chromatin structure (PubMed:<u>22334664</u>, PubMed:<u>23317504</u>, PubMed:<u>23589332</u>). During interphase, plays a key role in regulating the transcription of signal-inducible genes by associating with the P-TEFb complex and recruiting it to promoters (PubMed: 16109376, PubMed: 16109377, PubMed: 19596240,

> PubMed: 23589332, PubMed: 24360279). Also recruits P-TEFb complex to distal enhancers, so called anti-pause enhancers in collaboration with JMJD6

(PubMed: 16109376, PubMed: 16109377, PubMed: 19596240,

PubMed:23589332, PubMed:24360279). BRD4 and JMJD6 are required to form the transcriptionally active P-TEFb complex by displacing negative regulators such as HEXIM1 and 7SKsnRNA complex from P-TEFb, thereby transforming it into an active form that can then phosphorylate the C-terminal domain (CTD)

of RNA polymerase II (PubMed:16109376, PubMed:16109377,

PubMed: 19596240, PubMed: 23589332, PubMed: 24360279). Regulates

differentiation of naive CD4(+) T-cells into T-helper Th17 by promoting recruitment of P-TEFb to promoters (By similarity). Promotes phosphorylation of 'Ser-2' of the C-terminal domain (CTD) of RNA polymerase II (PubMed:23086925). According to a report, directly acts as an atypical protein kinase and mediates phosphorylation of 'Ser-2' of the C-terminal domain (CTD) of RNA polymerase II; these data however need additional evidences in vivo (PubMed:22509028). In addition to acetylated histones, also recognizes and binds acetylated RELA, leading to further recruitment of the P-TEFb complex and subsequent activation of NF-kappa-B (PubMed:19103749). Also acts as a regulator of p53/TP53-mediated transcription: following phosphorylation by CK2, recruited to p53/TP53 specific target promoters (PubMed:23317504).

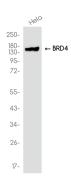
#### **Cellular Location**

Nucleus. Chromosome. Note=Associates with acetylated chromatin (PubMed:16109376, PubMed:21890894). Released from chromatin upon deacetylation of histones that can be triggered by different signals such as activation of the JNK pathway or nocodazole treatment (PubMed:16109376, PubMed:21890894). Preferentially localizes to mitotic chromosomes, while it does not localize to meiotic chromosomes (PubMed:16109376, PubMed:21890894).

#### **Tissue Location**

Ubiquitously expressed.

## **Images**



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