

CDK9 Rabbit mAb
Catalog # AP75247**Specification****CDK9 Rabbit mAb - Product Information**

Application	WB, IHC, IF
Primary Accession	P50750
Reactivity	Human, Mouse, Rat
Host	Rabbit
Clonality	Monoclonal Antibody
Calculated MW	42778

CDK9 Rabbit mAb - Additional Information

Gene ID 1025

Other Names

CDK9

Dilution

WB~~1/500-1/1000

IHC~~1/50-1/100

IF~~1/50-1/200

Format

Liquid

CDK9 Rabbit mAb - Protein Information**Name** CDK9 {ECO:0000303|PubMed:10903437, ECO:0000312|HGNC:HGNC:1780}**Function**

Protein kinase involved in the regulation of transcription (PubMed:10574912, PubMed:10757782, PubMed:11145967, PubMed:11575923, PubMed:11809800, PubMed:11884399, PubMed:14701750, PubMed:16109376, PubMed:16109377, PubMed:20930849, PubMed:28426094, PubMed:29335245). Member of the cyclin-dependent kinase pair (CDK9/cyclin-T) complex, also called positive transcription elongation factor b (P-TEFb), which facilitates the transition from abortive to productive elongation by phosphorylating the CTD (C-terminal domain) of the large subunit of RNA polymerase II (RNAP

II) POLR2A, SUPT5H and RDBP (PubMed: [10574912](http://www.uniprot.org/citations/10574912)), PubMed: [10757782](http://www.uniprot.org/citations/10757782), PubMed: [11145967](http://www.uniprot.org/citations/11145967), PubMed: [11575923](http://www.uniprot.org/citations/11575923), PubMed: [11809800](http://www.uniprot.org/citations/11809800), PubMed: [11884399](http://www.uniprot.org/citations/11884399), PubMed: [14701750](http://www.uniprot.org/citations/14701750), PubMed: [16109376](http://www.uniprot.org/citations/16109376), PubMed: [16109377](http://www.uniprot.org/citations/16109377), PubMed: [20930849](http://www.uniprot.org/citations/20930849), PubMed: [28426094](http://www.uniprot.org/citations/28426094), PubMed: [30134174](http://www.uniprot.org/citations/30134174)). This complex is inactive when in the 7SK snRNP complex form (PubMed: [10574912](http://www.uniprot.org/citations/10574912), PubMed: [10757782](http://www.uniprot.org/citations/10757782), PubMed: [11145967](http://www.uniprot.org/citations/11145967), PubMed: [11575923](http://www.uniprot.org/citations/11575923), PubMed: [11809800](http://www.uniprot.org/citations/11809800), PubMed: [11884399](http://www.uniprot.org/citations/11884399), PubMed: [14701750](http://www.uniprot.org/citations/14701750), PubMed: [16109376](http://www.uniprot.org/citations/16109376), PubMed: [16109377](http://www.uniprot.org/citations/16109377), PubMed: [20930849](http://www.uniprot.org/citations/20930849), PubMed: [28426094](http://www.uniprot.org/citations/28426094)). Phosphorylates EP300, MYOD1, RPB1/POLR2A and AR and the negative elongation factors DSIF and NELFE (PubMed: [10912001](http://www.uniprot.org/citations/10912001), PubMed: [11112772](http://www.uniprot.org/citations/11112772), PubMed: [12037670](http://www.uniprot.org/citations/12037670), PubMed: [20081228](http://www.uniprot.org/citations/20081228), PubMed: [20980437](http://www.uniprot.org/citations/20980437), PubMed: [21127351](http://www.uniprot.org/citations/21127351), PubMed: [9857195](http://www.uniprot.org/citations/9857195)). Regulates cytokine inducible transcription networks by facilitating promoter recognition of target transcription factors (e.g. TNF-inducible RELA/p65 activation and IL-6-inducible STAT3 signaling) (PubMed: [17956865](http://www.uniprot.org/citations/17956865), PubMed: [18362169](http://www.uniprot.org/citations/18362169)). Promotes RNA synthesis in genetic programs for cell growth, differentiation and viral pathogenesis (PubMed: [10393184](http://www.uniprot.org/citations/10393184), PubMed: [11112772](http://www.uniprot.org/citations/11112772)). P-TEFb is also involved in cotranscriptional histone modification, mRNA processing and mRNA export (PubMed: [15564463](http://www.uniprot.org/citations/15564463), PubMed: [19575011](http://www.uniprot.org/citations/19575011), PubMed: [19844166](http://www.uniprot.org/citations/19844166)). Modulates a complex network of chromatin modifications including histone H2B monoubiquitination (H2Bub1), H3 lysine 4 trimethylation (H3K4me3) and H3K36me3; integrates phosphorylation during transcription with chromatin modifications to control co-transcriptional histone mRNA processing (PubMed: [15564463](http://www.uniprot.org/citations/15564463), PubMed: [19575011](http://www.uniprot.org/citations/19575011), PubMed: [19844166](http://www.uniprot.org/citations/19844166)). The CDK9/cyclin-K complex has also a kinase activity towards CTD of RNAP II and can substitute for CDK9/cyclin-T P-TEFb in vitro (PubMed: [21127351](http://www.uniprot.org/citations/21127351)). Replication stress response protein; the CDK9/cyclin-K complex is required for genome integrity maintenance, by promoting cell cycle recovery from replication arrest and limiting single-stranded DNA amount in response to replication stress, thus reducing the breakdown of stalled replication forks and avoiding DNA damage (PubMed: [21127351](http://www.uniprot.org/citations/21127351)).

<http://www.uniprot.org/citations/20493174> target="_blank">20493174). In addition, probable function in DNA repair of isoform 2 via interaction with KU70/XRCC6 (PubMed:http://www.uniprot.org/citations/20493174). Promotes cardiac myocyte enlargement (PubMed:http://www.uniprot.org/citations/20081228). RPB1/POLR2A phosphorylation on 'Ser-2' in CTD activates transcription (PubMed:http://www.uniprot.org/citations/21127351). AR phosphorylation modulates AR transcription factor promoter selectivity and cell growth. DSIF and NELF phosphorylation promotes transcription by inhibiting their negative effect (PubMed:http://www.uniprot.org/citations/10912001, PubMed:http://www.uniprot.org/citations/11112772, PubMed:http://www.uniprot.org/citations/9857195). The phosphorylation of MYOD1 enhances its transcriptional activity and thus promotes muscle differentiation (PubMed:http://www.uniprot.org/citations/12037670). Catalyzes phosphorylation of KAT5, promoting KAT5 recruitment to chromatin and histone acetyltransferase activity (PubMed:http://www.uniprot.org/citations/29335245).

Cellular Location

Nucleus. Cytoplasm. Nucleus, PML body. Note=Accumulates on chromatin in response to replication stress Complexed with CCNT1 in nuclear speckles, but uncomplexed form in the cytoplasm. The translocation from nucleus to cytoplasm is XPO1/CRM1- dependent. Associates with PML body when acetylated

Tissue Location

Ubiquitous.

CDK9 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](#)

CDK9 Rabbit mAb - Images



