

### **CDK8 Antibody (Center)**

Purified Rabbit Polyclonal Antibody (Pab)
Catalog # AP11972c

# **Specification**

## **CDK8 Antibody (Center) - Product Information**

Application WB, FC,E Primary Accession P49336

Other Accession Q66KH9, Q8R3L8, Q8IH47, NP 001251

Reactivity Human

Predicted Zebrafish, Mouse, Xenopus

Host Rabbit
Clonality Polyclonal
Isotype Rabbit IgG
Antigen Region 249-277

### **CDK8 Antibody (Center) - Additional Information**

#### **Gene ID 1024**

### **Other Names**

Cyclin-dependent kinase 8, Cell division protein kinase 8, Mediator complex subunit CDK8, Mediator of RNA polymerase II transcription subunit CDK8, Protein kinase K35, CDK8

### Target/Specificity

This CDK8 antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 249-277 amino acids from the Central region of human CDK8.

## **Dilution**

WB~~1:1000 FC~~1:10~50

#### **Format**

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is prepared by Saturated Ammonium Sulfate (SAS) precipitation followed by dialysis against PBS.

#### Storage

Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

### **Precautions**

CDK8 Antibody (Center) is for research use only and not for use in diagnostic or therapeutic procedures.

# **CDK8 Antibody (Center) - Protein Information**

### Name CDK8





**Function** Component of the Mediator complex, a coactivator involved in regulated gene transcription of nearly all RNA polymerase II-dependent genes. Mediator functions as a bridge to convey information from gene- specific regulatory proteins to the basal RNA polymerase II transcription machinery. Mediator is recruited to promoters by direct interactions with regulatory proteins and serves as a scaffold for the assembly of a functional pre-initiation complex with RNA polymerase II and the general transcription factors. Phosphorylates the CTD (C- terminal domain) of the large subunit of RNA polymerase II (RNAp II), which may inhibit the formation of a transcription initiation complex. Phosphorylates CCNH leading to down-regulation of the TFIIH complex and transcriptional repression. Recruited through interaction with MAML1 to hyperphosphorylate the intracellular domain of NOTCH, leading to its degradation.

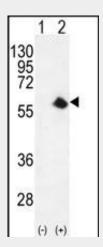
**Cellular Location** Nucleus.

### **CDK8 Antibody (Center) - Protocols**

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

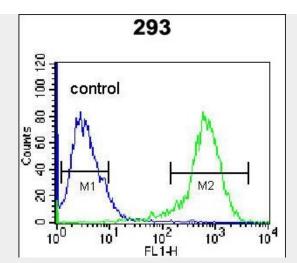
- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- <u>Immunoprecipitation</u>
- Flow Cytomety
- Cell Culture

### CDK8 Antibody (Center) - Images



Western blot analysis of CDK8 (arrow) using rabbit polyclonal CDK8 Antibody (Center) (Cat. #AP11972c). 293 cell lysates (2 ug/lane) either nontransfected (Lane 1) or transiently transfected (Lane 2) with the CDK8 gene.





CDK8 Antibody (Center) (Cat. #AP11972c) flow cytometric analysis of 293 cells (right histogram) compared to a negative control cell (left histogram).FITC-conjugated goat-anti-rabbit secondary antibodies were used for the analysis.

# CDK8 Antibody (Center) - Background

The protein encoded by this gene is a member of the cyclin-dependent protein kinase (CDK) family. CDK family members are highly similar to the gene products of Saccharomyces cerevisiae cdc28, and Schizosaccharomyces pombe cdc2, and are known to be important regulators of cell cycle progression. This kinase and its regulatory subunit cyclin C are components of the RNA polymerase II holoenzyme complex, which phosphorylates the carboxy-terminal domain (CTD) of the largest subunit of RNA polymerase II. This kinase has also been shown to regulate transcription by targeting the CDK7/cyclin H subunits of the general transcription initiation factor IIH (TFIIH), thus providing a link between the 'Mediator-like' protein complexes and the basal transcription machinery.

### CDK8 Antibody (Center) - References

Seo, J.O., et al. Oncol. Rep. 24(1):285-291(2010) Tsutsui, T., et al. Seikagaku 82(3):191-199(2010) Chattopadhyay, I., et al. Mutat. Res. 696(2):130-138(2010) Donner, A.J., et al. Nat. Struct. Mol. Biol. 17(2):194-201(2010) Alarcon, C., et al. Cell 139(4):757-769(2009)