# **Product Name: MAD1 Rabbit Polyclonal Antibody**

Catalog #: APRab13552



## **Summary**

**Production Name** MAD1 Rabbit Polyclonal Antibody

**Description** Rabbit Polyclonal Antibody

**Host** Rabbit

**Application** IF-P,IF-F,ICC/IF,ELISA **Reactivity** Human,Rat,Mouse

#### **Performance**

ConjugationUnconjugatedModificationUnmodified

**Isotype** IgG

Clonality Polyclonal Form Liquid

**Storage** Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.

Liquid in PBS containing 50% glycerol, 0.5% protective protein and 0.02% New type

preservative N.

**Purification** Affinity purification

## **Immunogen**

Buffer

Gene Name MAD1L1

MAD1L1; MAD1; TXBP181; Mitotic spindle assembly checkpoint protein MAD1; Mitotic

**Alternative Names** arrest deficient 1-like protein 1; MAD1-like protein 1; Mitotic checkpoint MAD1 protein

homolog; HsMAD1; hMAD1; Tax-binding protein 181

Gene ID 8379.0

Q9Y6D9. The antiserum was produced against synthesized peptide derived from SwissProt ID

human MAD1. AA range:394-443

## **Application**

**Dilution Ratio** IF-P/IF-F/ICC/IF 1:200-1:1000, ELISA 1:10000.Not yet tested in other applications.

**Molecular Weight** 

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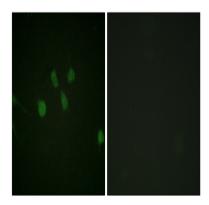
#### **Background**

MAD1L1 is a component of the mitotic spindle-assembly checkpoint that prevents the onset of anaphase until all chromosome are properly aligned at the metaphase plate. MAD1L1 functions as a homodimer and interacts with MAD2L1. MAD1L1 may play a role in cell cycle control and tumor suppression. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2015], disease: Defects in MAD1L1 are involved in the development and/or progression of various types of cancer., function: Component of the spindle-assembly checkpoint that prevents the onset of anaphase until all chromosomes are properly aligned at the metaphase plate. May recruit MAD2L1 to unattached kinetochores. Has a role in the correct positioning of the septum. Required for anchoring MAD2L1 to the nuclear periphery, induction: Increased by TP53.,PTM:Phosphorylated; by BUB1. Become hyperphosphorylated in late S through M phases or after mitotic spindle damage. Phosphorylated upon DNA damage, probably by ATM or ATR., similarity: Belongs to the MAD1 family., subcellular location: From the beginning to the end of mitosis, it is seen to move from a diffusely nuclear distribution to the centrosome, to the spindle midzone and finally to the midbody, subunit: Homodimer. Heterodimerizes with MAD2L1 in order to form a tetrameric MAD1L1-MAD2L1 core complex. Perturbation of the original MAD1L1-MAD2L1 structure by the spindle checkpoint may decrease MAD2L1 affinity for MAD1L1. CDC20 can compete with MAD1L1 for MAD2L1 binding, until the attachment and/or tension dampen the checkpoint signal, preventing further release of MAD2L1 on to CDC20. Also able to interact with the BUB1/BUB3 complex and the viral Tax protein. Interacts with TPR, tissue specificity: Expressed weakly at G0/G1 and highly at late S and G2/M phase.,

#### Research Area

Cell\_Cycle\_G1S;Cell\_Cycle\_G2M\_DNA;

#### **Image Data**



Immunofluorescence analysis of NIH/3T3 cells, using MAD1 Antibody. The picture on the right is blocked with the synthesized peptide.

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#### Note

For research use only.

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