

## MAD1 (phospho Ser428) Polyclonal Antibody

Catalog No: YP1007

**Reactivity:** Human; Rat; Mouse;

**Applications:** IHC;IF;ELISA

Target: MAD1

**Fields:** >>Cell cycle;>>Oocyte meiosis;>>Progesterone-mediated oocyte

maturation;>>Human T-cell leukemia virus 1 infection;>>Viral carcinogenesis

Gene Name: MAD1L1

**Protein Name:** Mitotic spindle assembly checkpoint protein MAD1

Q9Y6D9

Q9WTX8

Human Gene Id: 8379

**Human Swiss Prot** 

No:

**Mouse Swiss Prot** 

No:

**Immunogen:** The antiserum was produced against synthesized peptide derived from human

MAD1 around the phosphorylation site of Ser428. AA range:394-443

**Specificity:** Phospho-MAD1 (S428) Polyclonal Antibody detects endogenous levels of

MAD1 protein only when phosphorylated at S428.

**Formulation :** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Source: Polyclonal, Rabbit, IgG

**Dilution:** IHC 1:100 - 1:300. ELISA: 1:5000.. IF 1:50-200

**Purification:** The antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

Concentration: 1 mg/ml

1/3



Storage Stability: -15°C to -25°C/1 year(Do not lower than -25°C)

Molecularweight: 83kD

Location:

**Cell Pathway:** Cell\_Cycle\_G1S;Cell\_Cycle\_G2M\_DNA;

**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],

**Function:** 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 or

Subcellular Nucleus . Chromosome, centromere, kinetochore . Nucleus envelope .

Cytoplasm, cytoskeleton, microtubule organizing center, centrosome. Cytoplasm, cytoskeleton, spindle. Cytoplasm, cytoskeleton, spindle pole. Co-localizes with TPR at the nucleus envelope during interphase and throughout the cell cycle (PubMed:22351768, PubMed:18981471). 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 (PubMed:9546394). Localizes to kinetochores during prometaphase (PubMed:22351768, PubMed:29162720). Does not localize to kinetochores during metaphase (PubMed:29162720).

Colocalizes with NEK2 at the kinetochore (PubMed:14978040). Colocalizes with

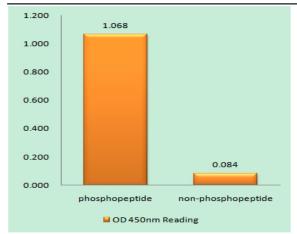
IK at spindle poles during metaphase and ana

**Expression:** [Isoform 1]: Expressed in hepatocellular carcinomas and hepatoma cell lines (at

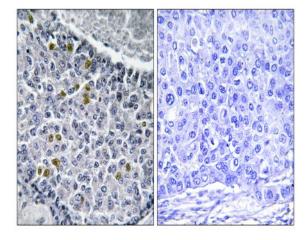
protein level).; [Isoform 3]: Expressed in hepatocellular carcinomas and

hepatoma cell lines (at protein level).

## **Products Images**



Enzyme-Linked Immunosorbent Assay (Phospho-ELISA) for Immunogen Phosphopeptide (Phospho-left) and Non-Phosphopeptide (Phospho-right), using MAD1 (Phospho-Ser428) Antibody



Immunohistochemistry analysis of paraffin-embedded human breast carcinoma, using MAD1 (Phospho-Ser428) Antibody. The picture on the right is blocked with the phospho peptide.