

## Pin1 Polyclonal Antibody

Catalog No: YT3730

**Reactivity:** Human; Mouse; Rat

**Applications:** IHC;IF;ELISA

Target: Pin1

**Fields:** >>Viral life cycle - HIV-1;>>RIG-I-like receptor signaling pathway

Gene Name: PIN1

**Protein Name:** Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1

Human Gene Id: 5300

**Human Swiss Prot** 

No:

Mouse Gene ld: 23988

**Mouse Swiss Prot** 

No:

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

Pin1. AA range:1-50

Q13526

Q9QUR7

**Specificity:** Pin1 Polyclonal Antibody detects endogenous levels of Pin1 protein.

**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. IF 1:200 - 1:1000. ELISA: 1:5000. Not yet tested in other

applications.

**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: 18kD

Cell Pathway: RIG-I-like receptor;

**Background:** Peptidyl-prolyl cis/trans isomerases (PPlases) catalyze the cis/trans

isomerization of peptidyl-prolyl peptide bonds. This gene encodes one of the PPlases, which specifically binds to phosphorylated ser/thr-pro motifs to catalytically regulate the post-phosphorylation conformation of its substrates. The conformational regulation catalyzed by this PPlase has a profound impact on key proteins involved in the regulation of cell growth, genotoxic and other stress responses, the immune response, induction and maintenance of pluripotency, germ cell development, neuronal differentiation, and survival. This enzyme also plays a key role in the pathogenesis of Alzheimer's disease and many cancers. Multiple alternatively spliced transcript variants have been found for this

gene.[provided by RefSeq, Jun 2011],

**Function:** catalytic activity:Peptidylproline (omega=180) = peptidylproline

(omega=0).,domain:The WW domain is required for the interaction with STIL and MPHOSPH1.,function:Essential PPIase that regulates mitosis presumably by interacting with NIMA and attenuating its mitosis-promoting activity. Displays a preference for an acidic residue N-terminal to the isomerized proline bond. Catalyzing pSer/Thr-Pro cis/trans isomerizations.,PTM:Phosphorylated upon

DNA damage, probably by ATM or ATR.,similarity:Contains 1 PpiC

domain.,similarity:Contains 1 WW domain.,subunit:Interacts with STIL (By

similarity). Interacts with MPHOSPH1.,

Subcellular Location:

Nucleus . Nucleus speckle . Cytoplasm . Colocalizes with NEK6 in the nucleus (PubMed:16476580). Mainly localized in the nucleus but phosphorylation at

Ser-71 by DAPK1 results in inhibition of its nuclear localization

(PubMed:21497122)...

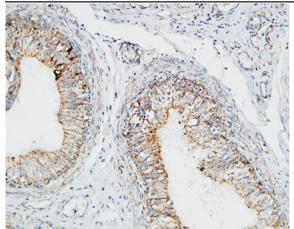
**Expression:** Expressed in immune cells in the lung (at protein level) (PubMed:29686383).

The phosphorylated form at Ser-71 is expressed in normal breast tissue cells but

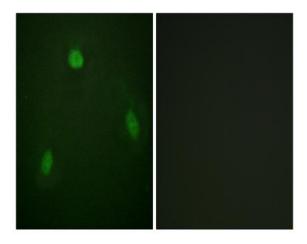
not in breast cancer cells.

## **Products Images**





Immunohistochemical analysis of paraffin-embedded Human testis. 1, Antibody was diluted at 1:100(4 $^{\circ}$  overnight). 2, Highpressure and temperature EDTA, pH8.0 was used for antigen retrieval. 3,Secondary antibody was diluted at 1:200(room temperature, 30min).



Immunofluorescence analysis of NIH/3T3 cells, using Pin1 Antibody. The picture on the right is blocked with the synthesized peptide.