

**Anti-Pin1 Picoband Antibody**  
Catalog # ABO12007**Specification****Anti-Pin1 Picoband Antibody - Product Information**

Application	WB
Primary Accession	<a href="#">Q13526</a>
Host	Rabbit
Reactivity	Human, Mouse, Rat
Clonality	Polyclonal
Format	Lyophilized

**Description**

Rabbit IgG polyclonal antibody for Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1(PIN1) detection. Tested with WB in Human;Mouse;Rat.

**Reconstitution**

Add 0.2ml of distilled water will yield a concentration of 500ug/ml.

**Anti-Pin1 Picoband Antibody - Additional Information**

**Gene ID** 5300

**Other Names**

Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1, 5.2.1.8, Peptidyl-prolyl cis-trans isomerase Pin1, PPlase Pin1, Rotamase Pin1, PIN1

**Calculated MW**

18243 MW KDa

**Application Details**

Western blot, 0.1-0.5 µg/ml, Human, Mouse, Rat

**Subcellular Localization**

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).

**Tissue Specificity**

The phosphorylated form at Ser-71 is expressed in normal breast tissue cells but not in breast cancer cells.

**Protein Name**

Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1

**Contents**

Each vial contains 5mg BSA, 0.9mg NaCl, 0.2mg Na<sub>2</sub>HPO<sub>4</sub>, 0.05mg Na<sub>3</sub>.

**Immunogen**

E.coli-derived human Pin1 recombinant protein (Position: M1-E163). Human Pin1 shares 95%

amino acid (aa) sequence identity with mouse Pin1.

#### Purification

Immunogen affinity purified.

#### Cross Reactivity

No cross reactivity with other proteins

#### Storage

**At -20°C for one year. After reconstitution, at 4°C for one month. It can also be aliquotted and stored frozen at -20°C for a longer time. Avoid repeated freezing and thawing.**

#### Sequence Similarities

Contains 1 PpiC domain.

### Anti-Pin1 Picoband Antibody - Protein Information

#### Name PIN1

#### Function

Peptidyl-prolyl cis/trans isomerase (PPIase) that binds to and isomerizes specific phosphorylated Ser/Thr-Pro (pSer/Thr-Pro) motifs (PubMed: [21497122](http://www.uniprot.org/citations/21497122), PubMed: [23623683](http://www.uniprot.org/citations/23623683), PubMed: [29686383](http://www.uniprot.org/citations/29686383)). By inducing conformational changes in a subset of phosphorylated proteins, acts as a molecular switch in multiple cellular processes (PubMed: [21497122](http://www.uniprot.org/citations/21497122), PubMed: [22033920](http://www.uniprot.org/citations/22033920), PubMed: [23623683](http://www.uniprot.org/citations/23623683)). Displays a preference for acidic residues located N-terminally to the proline bond to be isomerized. Regulates mitosis presumably by interacting with NIMA and attenuating its mitosis-promoting activity. Down-regulates kinase activity of BTK (PubMed: [16644721](http://www.uniprot.org/citations/16644721)). Can transactivate multiple oncogenes and induce centrosome amplification, chromosome instability and cell transformation. Required for the efficient dephosphorylation and recycling of RAF1 after mitogen activation (PubMed: [15664191](http://www.uniprot.org/citations/15664191)). Binds and targets PML and BCL6 for degradation in a phosphorylation-dependent manner (PubMed: [17828269](http://www.uniprot.org/citations/17828269)). Acts as a regulator of JNK cascade by binding to phosphorylated FBXW7, disrupting FBXW7 dimerization and promoting FBXW7 autoubiquitination and degradation: degradation of FBXW7 leads to subsequent stabilization of JUN (PubMed: [22608923](http://www.uniprot.org/citations/22608923)). May facilitate the ubiquitination and proteasomal degradation of RBBP8/CtIP through CUL3/KLHL15 E3 ubiquitin-protein ligase complex, hence favors DNA double-strand repair through error-prone non-homologous end joining (NHEJ) over error-free, RBBP8-mediated homologous recombination (HR) (PubMed: [23623683](http://www.uniprot.org/citations/23623683), PubMed: [27561354](http://www.uniprot.org/citations/27561354)). Upon IL33-induced lung inflammation, catalyzes cis-trans isomerization of phosphorylated IRAK3/IRAK-M, inducing IRAK3 stabilization, nuclear translocation and expression of pro-inflammatory genes in dendritic cells (PubMed: [29686383](http://www.uniprot.org/citations/29686383)). Catalyzes cis-trans isomerization of phosphorylated phosphoglycerate kinase PGK1 under hypoxic conditions to promote its binding to the TOM complex and targeting to the mitochondrion (PubMed: [29686383](http://www.uniprot.org/citations/29686383)).

href="http://www.uniprot.org/citations/26942675" target="\_blank">26942675</a>).

#### Cellular Location

Nucleus. Nucleus speckle. Cytoplasm Note=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)

#### Tissue Location

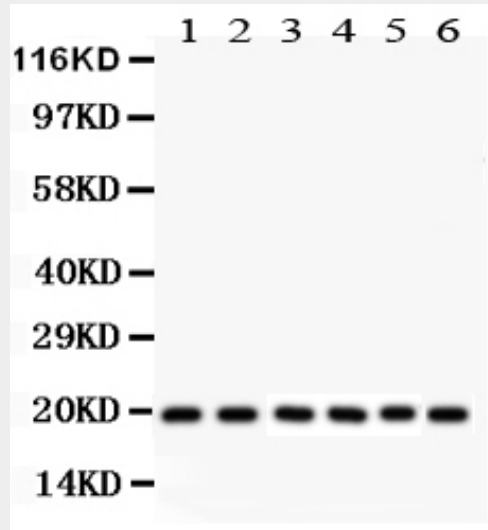
Expressed in immune cells in the lung (at protein level) (PubMed:29686383). The phosphorylated form at Ser-71 is expressed in normal breast tissue cells but not in breast cancer cells

### Anti-Pin1 Picoband Antibody - 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](#)

### Anti-Pin1 Picoband Antibody - Images



Anti- PIN1 Picoband antibody, ABO12007, Western blotting All lanes: Anti PIN1 (ABO12007) at 0.5ug/ml Lane 1: Rat Lung Tissue Lysate at 50ug Lane 2: Mouse Lung Tissue Lysate at 50ug Lane 3: NRK Whole Cell Lysate at 40ug Lane 4: A549 Whole Cell Lysate at 40ug Lane 5: 293T Whole Cell Lysate at 40ug Lane 6: MCF-7 Whole Cell Lysate at 40ug Predicted bind size: 19KD Observed bind size: 19KD

### Anti-Pin1 Picoband Antibody - Background

Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1, also called DOD, is an enzyme that in humans is encoded by the PIN1 gene. It is mapped to 19p13.2. The enzyme binds to a subset of proteins and thus plays a role as a post phospho PIN1rol in regulating protein function. Studies have

shown that the deregulation of PIN1 may play a pivotal role in various diseases. Notably, the up-regulation of PIN1 may be implicated in certain cancers, and the down-regulation of Pin1 may be implicated in Alzheimer's disease. Inhibitors of PIN1 may have therapeutic implications for cancer and immune disorders. PIN1 activity regulates the outcome of proline-directed kinase (e.g. MAPK, CDK or GSK3) signalling and consequently regulates cell proliferation (in part through control of cyclin D1 levels and stability) and cell survival. PIN1 also has an essential role in maintaining cell proliferation and regulating cyclin D1 function.