

### **Hsp90 Antibody**

Purified Mouse Monoclonal Antibody (Mab)
Catalog # AP52820

### **Specification**

### **Hsp90 Antibody - Product Information**

Application
Primary Accession
Reactivity
Host
Clonality
Isotype
Calculated MW

WB
P07900
Human, Mouse
Mouse
Monoclonal
IgG1
90 KDa

### **Hsp90 Antibody - Additional Information**

**Gene ID 3320** 

#### **Other Names**

D6S182;FLJ26984;FLJ31884;Heat shock 86 kDa;heat shock 90kDa protein 1 alpha;Heat shock protein 90kDa alpha cytosolic class A member 1;Heat shock protein 90kDa alpha cytosolic class B member 1; Heat shock protein HSP 90 alpha;Heat shock protein HSP 90 beta;Heat shock protein HSP 90-alpha;HS90A\_HUMAN;HSP 84;HSP 86;Hsp 90;HSP84;HSP86;Hsp89;Hsp90;HSP90 Beta;HSP9 0A;HSP90AA1;HSP90AB1;HSP90B;HSP90N;HSPC1;HSPC2;HSPCA;HSPCAL1;HSPCAL4;HSPCB;HSPN;L AP2;Lipopolysaccharide associated protein2;LPS associated protein 2;NY REN 38 antigen;Renal carcinoma antigen NY-REN-38.

# **Dilution**

WB~~1:2000

### **Format**

Purified mouse monoclonal in PBS(pH 7.4) containing with 0.09% (W/V) sodium azide and 50% glycerol.

### Storage

Store at -20 °C. Stable for 12 months from date of receipt

#### **Hsp90 Antibody - Protein Information**

Name HSP90AA1 (HGNC:5253)

Synonyms HSP90A, HSPC1, HSPCA

#### **Function**

Molecular chaperone that promotes the maturation, structural maintenance and proper regulation of specific target proteins involved for instance in cell cycle control and signal transduction. Undergoes a functional cycle that is linked to its ATPase activity which is essential for its chaperone activity. This cycle probably induces conformational changes in the client proteins,



thereby causing their activation. Interacts dynamically with various co-chaperones that modulate its substrate recognition, ATPase cycle and chaperone function (PubMed: <a href="http://www.uniprot.org/citations/11274138" target="\_blank">11274138</a>, PubMed:<a href="http://www.uniprot.org/citations/12526792" target="\_blank">12526792</a>, PubMed:<a href="http://www.uniprot.org/citations/15577939" target="blank">15577939</a>, PubMed:<a href="http://www.uniprot.org/citations/15937123" target="blank">15937123</a>, PubMed:<a href="http://www.uniprot.org/citations/27353360" target=" blank">27353360</a>, PubMed:<a href="http://www.uniprot.org/citations/29127155" target="blank">29127155</a>). Engages with a range of client protein classes via its interaction with various co-chaperone proteins or complexes, that act as adapters, simultaneously able to interact with the specific client and the central chaperone itself (PubMed: <a href="http://www.uniprot.org/citations/29127155" target=" blank">29127155</a>). Recruitment of ATP and co-chaperone followed by client protein forms a functional chaperone. After the completion of the chaperoning process, properly folded client protein and co- chaperone leave HSP90 in an ADP-bound partially open conformation and finally, ADP is released from HSP90 which acquires an open conformation for the next cycle  $(PubMed:<a\ href="http://www.uniprot.org/citations/26991466" target="\_blank">26991466</a>, PubMed:<a href="http://www.uniprot.org/citations/27295069" target="\_blank">27295069</a>).$ Plays a critical role in mitochondrial import, delivers preproteins to the mitochondrial import receptor TOMM70 (PubMed:<a href="http://www.uniprot.org/citations/12526792" target=" blank">12526792</a>). Apart from its chaperone activity, it also plays a role in the regulation of the transcription machinery. HSP90 and its co-chaperones modulate transcription at least at three different levels (PubMed:<a href="http://www.uniprot.org/citations/25973397" target=" blank">25973397</a>). In the first place, they alter the steady-state levels of certain transcription factors in response to various physiological cues (PubMed:<a href="http://www.uniprot.org/citations/25973397" target="\_blank">25973397</a>). Second, they modulate the activity of certain epigenetic modifiers, such as histone deacetylases or DNA methyl transferases, and thereby respond to the change in the environment (PubMed: <a href="http://www.uniprot.org/citations/25973397" target=" blank">25973397</a>). Third, they participate in the eviction of histones from the promoter region of certain genes and thereby turn on gene expression (PubMed:<a href="http://www.uniprot.org/citations/25973397" target=" blank">25973397</a>). Binds bacterial lipopolysaccharide (LPS) and mediates LPS-induced inflammatory response, including TNF secretion by monocytes (PubMed: <a href="http://www.uniprot.org/citations/11276205" target=" blank">11276205</a>). Antagonizes STUB1-mediated inhibition of TGF-beta signaling via inhibition of STUB1-mediated SMAD3 ubiquitination and degradation (PubMed: <a href="http://www.uniprot.org/citations/24613385" target=" blank">24613385</a>). Mediates the association of TOMM70 with IRF3 or TBK1 in mitochondrial outer membrane which promotes host antiviral response (PubMed: <a href="http://www.uniprot.org/citations/20628368" target=" blank">20628368</a>, PubMed:<a href="http://www.uniprot.org/citations/25609812" target="\_blank">25609812</a>).

### **Cellular Location**

Nucleus {ECO:0000250|UniProtKB:P07901}. Cytoplasm {ECO:0000250|UniProtKB:P07901}. Melanosome. Cell membrane. Mitochondrion. Note=Identified by mass spectrometry in melanosome fractions from stage I to stage IV

#### **Hsp90 Antibody - Protocols**

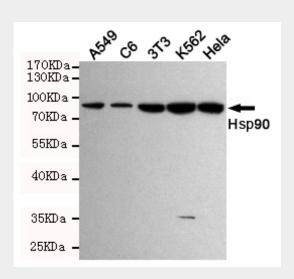
Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- <u>Immunohistochemistry</u>
- <u>Immunofluorescence</u>
- Immunoprecipitation



- Flow Cytomety
- Cell Culture

## Hsp90 Antibody - Images



Western blot detection of Hsp90 in Hela,3T3,C6,K562 and A549 cell lysates using Hsp90 mouse mAb (1:2000 diluted).Exposion time: 4min.Predicted band size:90KDa.Observed band size:90KDa.

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## **Hsp90 Antibody - References**

Soeda E.,et al.Nucleic Acids Res. 17:7108-7108(1989). Yamazaki M.,et al.Agric. Biol. Chem. 54:3163-3170(1990). Hickey E.,et al.Mol. Cell. Biol. 9:2615-2626(1989). Chen B.,et al.Genomics 86:627-637(2005). Ota T.,et al.Nat. Genet. 36:40-45(2004).