

## HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9)

Mouse Monoclonal Antibody Catalog # ALS16872

# **Specification**

# HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - Product Information

Application IHC, WB, E
Primary Accession P08238
Other Accession 3326

Reactivity Human, Mouse, Rat, Monkey

Host Mouse
Clonality Monoclonal

Isotype IgG1
Calculated MW 83264

## HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - Additional Information

# **Gene ID 3326**

# **Other Names**

HSP90AB1, D6S182, Heat shock 84 kDa, Heat shock protein beta, Heat shock protein HSP 90-beta, HSP 84, HSP84, HSP 90, HSP90-BETA, HSP90B, HSPC2, HSPCB

## Target/Specificity

Human HSP 90

### **Reconstitution & Storage**

Ascites, 0.03% sodium azide. Long term: -20°C; Short term: +4°C; Avoid freeze-thaw cycles.

#### **Precautions**

HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) is for research use only and not for use in diagnostic or therapeutic procedures.

# HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - Protein Information

Name HSP90AB1 (HGNC:5258)

Synonyms HSP90B, HSPC2, HSPCB

### **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 linked to its ATPase 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/16478993" target="\_blank">16478993</a>, PubMed:<a href="http://www.uniprot.org/citations/19696785" target="\_blank">19696785</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. 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/27295069" target="\_blank">27295069</a>, PubMed:<a href="http://www.uniprot.org/citations/26991466" target="\_blank">26991466</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. They first alter the steady-state levels of certain transcription factors in response to various physiological cues. 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. 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>). 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>). Promotes cell differentiation by chaperoning BIRC2 and thereby protecting from auto-ubiquitination and degradation by the proteasomal machinery (PubMed:<a

href="http://www.uniprot.org/citations/18239673" target="\_blank">18239673</a>). Main chaperone involved in the phosphorylation/activation of the STAT1 by chaperoning both JAK2 and PRKCE under heat shock and in turn, activates its own transcription (PubMed:<a href="http://www.uniprot.org/citations/20353823" target="\_blank">20353823</a>). Involved in the translocation into ERGIC (endoplasmic reticulum-Golgi intermediate compartment) of leaderless cargos (lacking the secretion signal sequence) such as the interleukin 1/IL-1; the translocation process is mediated by the cargo receptor TMED10 (PubMed:<a href="http://www.uniprot.org/citations/32272059" target=" blank">32272059</a>).

## **Cellular Location**

Cytoplasm. Melanosome Nucleus. Secreted. Cell membrane. Dynein axonemal particle {ECO:0000250|UniProtKB:Q6AZV1}. Cell surface. Note=Identified by mass spectrometry in melanosome fractions from stage I to stage IV (PubMed:17081065) Translocates with BIRC2 from the nucleus to the cytoplasm during differentiation (PubMed:18239673). Secreted when associated with TGFB1 processed form (LAP) (PubMed:20599762).

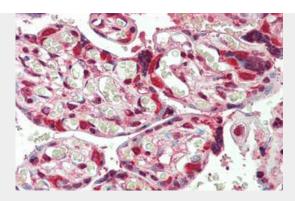
# HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - Protocols

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

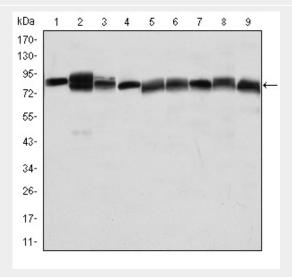
- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- <u>Immunoprecipitation</u>
- Flow Cytomety
- Cell Culture

### HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - Images

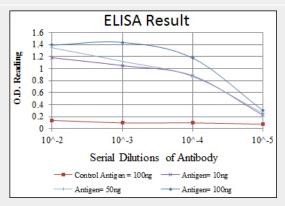




Anti-HSP90AB1 / HSP90 antibody IHC staining of human placenta.



Western blot using HSP90AB1 mouse monoclonal antibody against Jurkat (1), A431 (2), HeLa (3),...



Red: Control Antigen (100ng); Purple: Antigen (10ng); Green: Antigen (50ng); Blue: Antigen (100ng);

# HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - Background

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. 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.



# HSP90AB1 / HSP90 Alpha B1 Antibody (clone 1D9) - References

Rebbe N.F.,et al.Gene 53:235-245(1987).
Rebbe N.F.,et al.J. Biol. Chem. 264:15006-15011(1989).
Hoffmann T.,et al.Gene 74:491-501(1988).
Lu L.,et al.Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.
Wiemann S.,et al.Genome Res. 11:422-435(2001).