

Hsp90 Antibody

Purified Mouse Monoclonal Antibody (Mab) Catalog # AM2191b

Product Information

Application WB, E **Primary Accession** P07900

Reactivity Human, Rat, Mouse

 $\begin{array}{lll} \textbf{Host} & \textbf{Mouse} \\ \textbf{Clonality} & \textbf{Monoclonal} \\ \textbf{Isotype} & \textbf{IgG1}, \lambda \\ \textbf{Clone Names} & 803\text{CT9.1.2} \\ \textbf{Calculated MW} & 84660 \\ \end{array}$

Additional Information

Gene ID 3320

Other Names Heat shock protein HSP 90-alpha, Heat shock 86 kDa, HSP 86, HSP86,

Lipopolysaccharide-associated protein 2, LAP-2, LPS-associated protein 2, Renal carcinoma antigen NY-REN-38, HSP90AA1, HSP90A, HSPC1, HSPCA

Target/SpecificityThis antibody is generated from a mouse immunized with a KLH conjugated

synthetic peptide between 150-400 amino acids from human.

Dilution WB~~1:1000 E~~Use at an assay dependent concentration.

Format Purified monoclonal antibody supplied in PBS with 0.09% (W/V) sodium azide.

This antibody is purified through a protein G column, followed by dialysis

against PBS.

Storage Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store

at -20°C in small aliquots to prevent freeze-thaw cycles.

Precautions Hsp90 Antibody is for research use only and not for use in diagnostic or

therapeutic procedures.

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: 11274138, PubMed: 12526792, PubMed: 15577939, PubMed:15937123, PubMed:27353360, PubMed:29127155). 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:29127155). 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: 26991466, PubMed: 27295069). Plays a critical role in mitochondrial import, delivers preproteins to the mitochondrial import receptor TOMM70 (PubMed:12526792). 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: 25973397). In the first place, they alter the steady-state levels of certain transcription factors in response to various physiological cues (PubMed: 25973397). 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: <u>25973397</u>). Third, they participate in the eviction of histones from the promoter region of certain genes and thereby turn on gene expression (PubMed: <u>25973397</u>). Binds bacterial lipopolysaccharide (LPS) and mediates LPS-induced inflammatory response, including TNF secretion by monocytes (PubMed: 11276205). Antagonizes STUB1-mediated inhibition of TGF-beta signaling via inhibition of STUB1-mediated SMAD3 ubiquitination and degradation (PubMed:24613385). Mediates the association of TOMM70 with IRF3 or TBK1 in mitochondrial outer membrane which promotes host antiviral response (PubMed: 20628368, PubMed: 25609812).

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

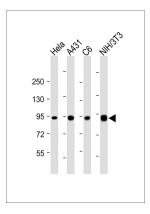
Background

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.

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).

Images



All lanes: Anti-Hsp90 at 1:4000 dilution Lane 1: Hela whole cell lysate Lane 2: A431 whole cell lysate Lane 3: C6 whole cell lysate Lane 4: NIH/3T3 whole cell lysate Lysates/proteins at 20 µg per lane. Secondary Goat Anti-mouse IgG, (H+L), Peroxidase conjugated at 1/10000 dilution. Predicted band size: 85 kDa Blocking/Dilution buffer: 5% NFDM/TBST.

Please note: All products are 'FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES'.