

Hsc70 Antibody

Rabbit mAb Catalog # AP90444

Specification

Hsc70 Antibody - Product Information

Application WB, IHC, FC, ICC, IP

Primary Accession P11142
Reactivity Rat

Clonality Monoclonal

Other Names

Heat shock cognate 71 kDa protein; HS7C; HSP73; HSP7C; HSPA10; HSPA8;

Isotype Rabbit IgG
Host Rabbit
Calculated MW 70898 Da

Hsc70 Antibody - Additional Information

Purification Affinity-chromatography

Immunogen A synthesized peptide derived from human

Hsc70

Description Acts as a repressor of transcriptional

activation. Inhibits the transcriptional coactivator activity of CITED1 on

Smad-mediated transcription. Chaperone.

Component of the PRP19-CDC5L complex

that forms an integral part of the

spliceosome and is required for activating

pre-mRNA splicing.

Storage Condition and Buffer Rabbit IgG in phosphate buffered saline,

pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid

freeze / thaw cycle.

Hsc70 Antibody - Protein Information

Name HSPA8 (HGNC:5241)

Function

Molecular chaperone implicated in a wide variety of cellular processes, including protection of the proteome from stress, folding and transport of newly synthesized polypeptides, chaperone-mediated autophagy, activation of proteolysis of misfolded proteins, formation and dissociation of protein complexes, and antigen presentation. Plays a pivotal role in the protein quality control system, ensuring the correct folding of proteins, the re-folding of misfolded proteins and controlling the targeting of proteins for subsequent degradation (PubMed:21148293, PubMed:21150129, PubMed:<a



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href="http://www.uniprot.org/citations/23018488" target=" blank">23018488</a>, PubMed:<a
href="http://www.uniprot.org/citations/24732912" target="blank">24732912</a>, PubMed:<a
href="http://www.uniprot.org/citations/27916661" target="blank">27916661</a>, PubMed:<a
href="http://www.uniprot.org/citations/2799391" target="_blank">2799391</a>, PubMed:<a
href="http://www.uniprot.org/citations/36586411" target=" blank">36586411</a>). This is
achieved through cycles of ATP binding, ATP hydrolysis and ADP release, mediated by
co-chaperones (PubMed: <a href="http://www.uniprot.org/citations/12526792"
target=" blank">12526792</a>, PubMed:<a href="http://www.uniprot.org/citations/21148293"
target="blank">21148293</a>, PubMed:<a href="http://www.uniprot.org/citations/21150129"
target="_blank">21150129</a>, PubMed:<a href="http://www.uniprot.org/citations/23018488"
target="blank">23018488</a>, PubMed:<a href="http://www.uniprot.org/citations/24732912"
target="blank">24732912</a>, PubMed:<a href="http://www.uniprot.org/citations/27916661"
target="blank">27916661</a>). The co-chaperones have been shown to not only regulate
different steps of the ATPase cycle of HSP70, but they also have an individual specificity such that
one co-chaperone may promote folding of a substrate while another may promote degradation
(PubMed:<a href="http://www.uniprot.org/citations/12526792" target=" blank">12526792</a>,
PubMed:<a href="http://www.uniprot.org/citations/21148293" target=" blank">21148293</a>,
PubMed:<a href="http://www.uniprot.org/citations/21150129" target="_blank">21150129</a>,
PubMed:<a href="http://www.uniprot.org/citations/23018488" target="blank">23018488</a>,
PubMed:<a href="http://www.uniprot.org/citations/24732912" target="blank">24732912</a>,
PubMed: <a href="http://www.uniprot.org/citations/27916661" target="blank">27916661</a>).
The affinity of HSP70 for polypeptides is regulated by its nucleotide bound state. In the ATP-bound
form, it has a low affinity for substrate proteins. However, upon hydrolysis of the ATP to ADP, it
undergoes a conformational change that increases its affinity for substrate proteins. HSP70 goes
through repeated cycles of ATP hydrolysis and nucleotide exchange, which permits cycles of
substrate binding and release. The HSP70-associated co-chaperones are of three types: J- domain
co-chaperones HSP40s (stimulate ATPase hydrolysis by HSP70), the nucleotide exchange factors
(NEF) such as BAG1/2/3 (facilitate conversion of HSP70 from the ADP-bound to the ATP-bound
state thereby promoting substrate release), and the TPR domain chaperones such as HOPX and
STUB1 (PubMed: <a href="http://www.uniprot.org/citations/24121476"
target=" blank">24121476</a>, PubMed:<a href="http://www.uniprot.org/citations/24318877"
target="blank">24318877</a>, PubMed:<a href="http://www.uniprot.org/citations/26865365"
target="blank">26865365</a>, PubMed:<a href="http://www.uniprot.org/citations/27474739"
target=" blank">27474739</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>). Acts as a
repressor of transcriptional activation. Inhibits the transcriptional coactivator activity of CITED1 on
Smad- mediated transcription. Component of the PRP19-CDC5L complex that forms an integral
part of the spliceosome and is required for activating pre- mRNA splicing. May have a scaffolding
role in the spliceosome assembly as it contacts all other components of the core complex. Binds
bacterial lipopolysaccharide (LPS) and mediates LPS-induced inflammatory response, including
TNF secretion by monocytes (PubMed: <a href="http://www.uniprot.org/citations/10722728"
target=" blank">10722728</a>, PubMed:<a href="http://www.uniprot.org/citations/11276205"
target="blank">11276205</a>). Substrate recognition component in chaperone-mediated
autophagy (CMA), a selective protein degradation process that mediates degradation of proteins
with a -KFERQ motif: HSPA8/HSC70 specifically recognizes and binds cytosolic proteins bearing a
-KFERQ motif and promotes their recruitment to the surface of the lysosome where they bind to
lysosomal protein LAMP2 (PubMed: <a href="http://www.uniprot.org/citations/11559757"
target=" blank">11559757</a>, PubMed:<a href="http://www.uniprot.org/citations/2799391"
target=" blank">2799391</a>, PubMed:<a href="http://www.uniprot.org/citations/36586411"
target=" blank">36586411</a>). KFERQ motif- containing proteins are eventually transported
into the lysosomal lumen where they are degraded (PubMed: <a
href="http://www.uniprot.org/citations/11559757" target="_blank">11559757</a>, PubMed:<a href="http://www.uniprot.org/citations/2799391" target="_blank">2799391</a>, PubMed:<a
href="http://www.uniprot.org/citations/36586411" target=" blank">36586411</a>). In
conjunction with LAMP2, facilitates MHC class II presentation of cytoplasmic antigens by guiding
antigens to the lysosomal membrane for interaction with LAMP2 which then elicits MHC class II
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presentation of peptides to the cell membrane (PubMed:15894275). Participates in the ER-associated degradation (ERAD) quality control pathway in conjunction with J domain-containing co- chaperones and the E3 ligase STUB1 (PubMed:23990462). It is recruited to clathrin-coated vesicles through its interaction with DNAJC6 leading to activation of HSPA8/HSC70 ATPase activity and therefore uncoating of clathrin-coated vesicles (By similarity).

Cellular Location

Cytoplasm. Melanosome. Nucleus, nucleolus. Cell membrane. Lysosome membrane; Peripheral membrane protein; Cytoplasmic side. Note=Localized in cytoplasmic mRNP granules containing untranslated mRNAs (PubMed:17289661). Translocates rapidly from the cytoplasm to the nuclei, and especially to the nucleoli, upon heat shock (PubMed:1586970)

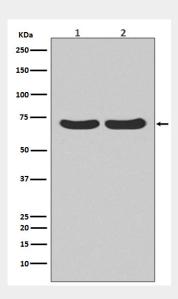
Tissue Location Ubiquitous..

Hsc70 Antibody - Protocols

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

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

Hsc70 Antibody - Images



Western blot analysis of Calreticulin expression in (1) HeLa cell lysate; (2) C6 cell lysate.