

HSF1 (Phospho Ser307) Rabbit pAb

CatalogNo: YP1188

Key Features

Host Species				
 Rabbit 				

ReactivityHuman,Rat,Mouse,

Applications
• WB,IF,ELISA

MW • 30kD (Observed)

Isotype • IgG

Recommended Dilution Ratios

WB 1:500-1:2000 IF 1:200-1:1000 ELISA 1:5000 Not yet tested in other applications.

Storage

Storage*-15°C to -25°C/1 year(Do not lower than -25°C)FormulationLiquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Basic Information

Clonality Polyclonal

Immunogen Information

ImmunogenThe antiserum was produced against synthesized peptide derived from human HSF1
around the phosphorylation site of Ser307. AA range:273-322

Specificity Phospho-HSF1 (S307) Polyclonal Antibody detects endogenous levels of HSF1 protein only when phosphorylated at S307.The name of modified sites may be influenced by many factors, such as species (the modified site was not originally found in human samples) and the change of protein sequence (the previous protein sequence is incomplete, and the protein sequence may be prolonged with the development of protein sequencing technology). When naming, we will use the "numbers" in historical reference to keep the sites consistent with the reports. The antibody binds to the following modification sequence (lowercase letters are modification sites):PQsPR

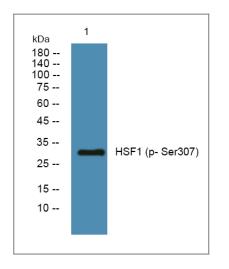
Target Information

Gene name	HSF1			
Protein Name	Heat shock factor protein 1 Organism Human	Gene ID <u>3297;</u>	UniProt ID Q00613;	
	Mouse		<u>P38532;</u>	
Cellular Localization	MouseP38532;Nucleus . Cytoplasm . Nucleus, nucleoplasm . Cytoplasm, perinuclear region . Cytoplasmcytoskeleton, spindle pole . Cytoplasm, cytoskeleton, microtubule organizing center,centrosome . Chromosome, centromere, kinetochore . The monomeric form is cytoplasrin unstressed cells (PubMed:8455624, PubMed:26159920). Predominantly nuclear protectboth unstressed and heat shocked cells (PubMed:10413683, PubMed:10359787).Translocates in the nucleus upon heat shock (PubMed:8455624). Nucleocytoplasmicshuttling protein (PubMed:26159920). Colocalizes with IER5 in the nucleus(PubMed:27354066). Colocalizes with BAG3 to the nucleus upon heat stress(PubMed:8455624, PubMed:26159920). Localizes in subnuclear granules called nuclearstress bodies (nSBs) upon heat shock (PubMed:11447121, PubMed:11514557,PubMed:10359787, PubMed:25963659, PubMed:10747973, PubMed:24581496,PubMed:10359787, PubMed:12665592, PubMed:11514557, PubMed:14707147,PubMed:10359787). Colocalizes with SYMPK and SUMO1 in nSBs upon heat shock(PubMed:10359787). Colocalizes from the nucleus ot the cytoplasm during the attenuatand recovery phase period of the heat shock response (PubMed:26159920). Translocatethe cytoplasm in a YWHAE- and XPO1/CRM1-dependent manner (PubMed:12917326).Together with histone H2AX, redistributed in discrete nuclear DNA damage-induced fociafter ionizing radiation (IR) (PubMed:26359349). Colocalizes with calcium-responsivetransactivator SS18L1 at kinetochore region on the mitotic chromosomes(PubMed:18794143). Colocalizes with gamma tubulin at centrosome (PubMed:18794143).Localizes at spindle pole in metaphase (PubMed:18794143). <t< th=""><th>lle organizing center, omeric form is cytoplasmic lominantly nuclear protein in bMed:10359787).). Nucleocytoplasmic the nucleus n heat stress granules called nuclear Med:11514557, Med:24581496, s upon heat shock bMed:14707147, s and nSBs upon heat shock asm during the attenuation l:26159920). Translocates in r (PubMed:12917326). NA damage-induced foci th calcium-responsive mosomes some (PubMed:18794143).</th></t<>		lle organizing center, omeric form is cytoplasmic lominantly nuclear protein in bMed:10359787).). Nucleocytoplasmic the nucleus n heat stress granules called nuclear Med:11514557, Med:24581496, s upon heat shock bMed:14707147, s and nSBs upon heat shock asm during the attenuation l:26159920). Translocates in r (PubMed:12917326). NA damage-induced foci th calcium-responsive mosomes some (PubMed:18794143).	

Tissue specificity Adipose tissue, Brain, Epithelium, Muscle,

Function

Function:DNA-binding protein that specifically binds heat shock promoter elements (HSE) and activates transcription. In higher eukaryotes, HSF is unable to bind to the HSE unless the cells are heat shocked., PTM: Phosphorylated on multiple serine residues, a subset of which are involved in stress-related regulation of transcription activation. Constitutive phosphorylation represses transcriptional activity at normal temperatures. Levels increase on specific residues heat-shock and enhance HSF1 transactivation activity. Phosphorylation on Ser-307 derepresses activation on heat-stress and in combination with Ser-303 phosphorylation appears to be involved in recovery after heat-stress. Phosphorylated on Ser-230 by CAMK2, in vitro. Cadmium also enhances phosphorylation at this site. Phosphorylation on Ser-303 is a prerequisite for HSF1 sumoylation. Phosphorylation on Ser-121 inhibits transactivation and promotes HSP90 binding. Phosphorylation on Thr-142 also mediates transcriptional activity induced by heat., PTM: Sumoylated BY SUMO1 AND SUMO2 on heat-shock. Heat-inducible sumoylation occurs after 15 min of heat-shock, after which levels decrease and at 4 hours, levels return to control levels. Sumoylation has no effect on HSE binding nor on transcriptional activity. Phosphorylation on Ser-303 is a prerequisite for sumoylation., similarity: Belongs to the HSF family., subcellular location:Cytoplasmic during normal growth. On activation, translocates to nuclear stress granules. Colocalizes with SUMO1 in nuclear stress granules., subunit: Monomer. Under normal conditions, interacts with HSP90AA1 in the HSP90 multichaperone complex; the interaction prevents trimerization and activation of HSF1. On activation by heat-stress or by other factors such as metal ions, HSF1 is released from the complex, homotrimerizes, is hyperphosphorylated and translocated to the nucleus where, subsequently, it can activate transcription. Binds the complex through the regulatory domain. Interacts with SYMPK and CSTF2 in heat-stressed cells. Interacts with FKBP4 in the HSP90 multichaperone complex; the interaction is independent of the phosphorylation state of HSF1. Interacts with MAPKAPK2.,



Validation Data

Western blot analysis of lysates from A431 cells, primary antibody was diluted at 1:1000, 4° over night

Contact information

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