

# HSF1 (phospho Ser303) Polyclonal Antibody

Catalog # AP67065

## Product Information

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Application	WB, IHC-P, IF
Primary Accession	<a href="#">Q00613</a>
Reactivity	Human, Mouse
Host	Rabbit
Clonality	Polyclonal
Calculated MW	57260

## Additional Information

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Gene ID	3297
Other Names	HSF1; HSTF1; Heat shock factor protein 1; HSF 1; Heat shock transcription factor 1; HSTF 1
Dilution	WB~~Western Blot: 1/500 - 1/2000. Immunohistochemistry: 1/100 - 1/300. Immunofluorescence: 1/200 - 1/1000. ELISA: 1/20000. Not yet tested in other applications. IHC-P~~N/A IF~~1:50~200
Format	Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.09% (W/V) sodium azide.
Storage Conditions	-20°C

## Protein Information

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Name	HSF1 ( <a href="#">HGNC:5224</a> )
Synonyms	HSTF1
Function	Functions as a stress-inducible and DNA-binding transcription factor that plays a central role in the transcriptional activation of the heat shock response (HSR), leading to the expression of a large class of molecular chaperones, heat shock proteins (HSPs), that protect cells from cellular insult damage (PubMed: <a href="#">11447121</a> , PubMed: <a href="#">12659875</a> , PubMed: <a href="#">12917326</a> , PubMed: <a href="#">15016915</a> , PubMed: <a href="#">18451878</a> , PubMed: <a href="#">1871105</a> , PubMed: <a href="#">1986252</a> , PubMed: <a href="#">25963659</a> , PubMed: <a href="#">26754925</a> , PubMed: <a href="#">7623826</a> , PubMed: <a href="#">7760831</a> , PubMed: <a href="#">8940068</a> , PubMed: <a href="#">8946918</a> , PubMed: <a href="#">9121459</a> , PubMed: <a href="#">9341107</a> , PubMed: <a href="#">9499401</a> , PubMed: <a href="#">9535852</a> , PubMed: <a href="#">9727490</a> ). In unstressed cells, is present in a HSP90-containing multichaperone complex that maintains it in a non-DNA-binding inactivated monomeric form (PubMed: <a href="#">11583998</a> , PubMed: <a href="#">16278218</a> , PubMed: <a href="#">9727490</a> ). Upon exposure to heat and other stress stimuli, undergoes homotrimerization and activates HSP gene transcription through binding to site-specific heat shock elements (HSEs)

present in the promoter regions of HSP genes (PubMed:[10359787](#), PubMed:[11583998](#), PubMed:[12659875](#), PubMed:[16278218](#), PubMed:[1871105](#), PubMed:[1986252](#), PubMed:[25963659](#), PubMed:[26754925](#), PubMed:[7623826](#), PubMed:[7935471](#), PubMed:[8455624](#), PubMed:[8940068](#), PubMed:[9499401](#), PubMed:[9727490](#)). Upon heat shock stress, forms a chromatin-associated complex with TTC5/STRAP and p300/EP300 to stimulate HSR transcription, therefore increasing cell survival (PubMed:[18451878](#)). Activation is reversible, and during the attenuation and recovery phase period of the HSR, returns to its unactivated form (PubMed:[11583998](#), PubMed:[16278218](#)). Binds to inverted 5'-NGAAN-3' pentamer DNA sequences (PubMed:[1986252](#), PubMed:[26727489](#)). Binds to chromatin at heat shock gene promoters (PubMed:[25963659](#)). Activates transcription of transcription factor FOXR1 which in turn activates transcription of the heat shock chaperones HSPA1A and HSPA6 and the antioxidant NADPH-dependent reductase DHRS2 (PubMed:[34723967](#)). Also serves several other functions independently of its transcriptional activity. Involved in the repression of Ras-induced transcriptional activation of the c-fos gene in heat-stressed cells (PubMed:[9341107](#)). Positively regulates pre-mRNA 3'-end processing and polyadenylation of HSP70 mRNA upon heat-stressed cells in a symplekin (SYMPK)-dependent manner (PubMed:[14707147](#)). Plays a role in nuclear export of stress-induced HSP70 mRNA (PubMed:[17897941](#)). Plays a role in the regulation of mitotic progression (PubMed:[18794143](#)). Also plays a role as a negative regulator of non-homologous end joining (NHEJ) repair activity in a DNA damage-dependent manner (PubMed:[26359349](#)). Involved in stress-induced cancer cell proliferation in a IER5-dependent manner (PubMed:[26754925](#)).

## Cellular Location

Nucleus. Cytoplasm. Nucleus, nucleoplasm. Cytoplasm, perinuclear region. Cytoplasm, cytoskeleton, spindle pole. Cytoplasm, cytoskeleton, microtubule organizing center, centrosome Chromosome, centromere, kinetochore  
 Note=The monomeric form is cytoplasmic in unstressed cells (PubMed:[26159920](#), PubMed:[8455624](#)). Predominantly nuclear protein in both unstressed and heat shocked cells (PubMed:[10359787](#), PubMed:[10413683](#)). Translocates in the nucleus upon heat shock (PubMed:[8455624](#)). Nucleocytoplasmic shuttling protein (PubMed:[26159920](#)). Colocalizes with IER5 in the nucleus (PubMed:[27354066](#)). Colocalizes with BAG3 to the nucleus upon heat stress (PubMed:[26159920](#), PubMed:[8455624](#)). Localizes in subnuclear granules called nuclear stress bodies (nSBs) upon heat shock (PubMed:[10359787](#), PubMed:[10747973](#), PubMed:[11447121](#), PubMed:[11514557](#), PubMed:[19229036](#), PubMed:[24581496](#), PubMed:[25963659](#)). Colocalizes with SYMPK and SUMO1 in nSBs upon heat shock (PubMed:[10359787](#), PubMed:[11447121](#), PubMed:[11514557](#), PubMed:[12665592](#), PubMed:[14707147](#)). Colocalizes with PRKACA/PKA in the nucleus and nSBs upon heat shock (PubMed:[21085490](#)). Relocalizes from the nucleus to the cytoplasm during the attenuation and recovery phase period of the heat shock response (PubMed:[26159920](#)). Translocates in the cytoplasm in a YWHAE- and XPO1/CRM1-dependent manner (PubMed:[12917326](#)). Together with histone H2AX, redistributed in discrete nuclear DNA damage-induced foci after ionizing radiation (IR) (PubMed:[26359349](#)). Colocalizes with calcium-responsive transactivator 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](#)). Colocalizes with PLK1 at spindle poles during prometaphase (PubMed:[18794143](#)).

## Background

Function as a stress-inducible and DNA-binding transcription factor that plays a central role in the transcriptional activation of the heat shock response (HSR), leading to the expression of a large class of

molecular chaperones heat shock proteins (HSPs) that protect cells from cellular insults' damage (PubMed:[1871105](#), PubMed:[11447121](#), PubMed:[1986252](#), PubMed:[7760831](#), PubMed:[7623826](#), PubMed:[8946918](#), PubMed:[8940068](#), PubMed:[9341107](#), PubMed:[9121459](#), PubMed:[9727490](#), PubMed:[9499401](#), PubMed:[9535852](#), PubMed:[12659875](#), PubMed:[12917326](#), PubMed:[15016915](#), PubMed:[25963659](#), PubMed:[26754925](#)). In unstressed cells, is present in a HSP90-containing multichaperone complex that maintains it in a non-DNA-binding inactivated monomeric form (PubMed:[9727490](#), PubMed:[11583998](#), PubMed:[16278218](#)). Upon exposure to heat and other stress stimuli, undergoes homotrimerization and activates HSP gene transcription through binding to site-specific heat shock elements (HSEs) present in the promoter regions of HSP genes (PubMed:[1871105](#), PubMed:[1986252](#), PubMed:[8455624](#), PubMed:[7935471](#), PubMed:[7623826](#), PubMed:[8940068](#), PubMed:[9727490](#), PubMed:[9499401](#), PubMed:[10359787](#), PubMed:[11583998](#), PubMed:[12659875](#), PubMed:[16278218](#), PubMed:[25963659](#), PubMed:[26754925](#)). Activation is reversible, and during the attenuation and recovery phase period of the HSR, returns to its unactivated form (PubMed:[11583998](#), PubMed:[16278218](#)). Binds to inverted 5'-NGAAN-3' pentamer DNA sequences (PubMed:[1986252](#), PubMed:[26727489](#)). Binds to chromatin at heat shock gene promoters (PubMed:[25963659](#)). Plays also several other functions independently of its transcriptional activity. Involved in the repression of Ras-induced transcriptional activation of the c-fos gene in heat-stressed cells (PubMed:[9341107](#)). Positively regulates pre-mRNA 3'-end processing and polyadenylation of HSP70 mRNA upon heat-stressed cells in a symplekin (SYMPK)-dependent manner (PubMed:[14707147](#)). Plays a role in nuclear export of stress-induced HSP70 mRNA (PubMed:[17897941](#)). Plays a role in the regulation of mitotic progression (PubMed:[18794143](#)). Plays also a role as a negative regulator of non-homologous end joining (NHEJ) repair activity in a DNA damage- dependent manner (PubMed:[26359349](#)). Involved in stress-induced cancer cell proliferation in a IER5-dependent manner (PubMed:[26754925](#)).

## Images



Western Blot analysis of MCF7+TNF cells using Phospho-HSF1 (S303) Polyclonal Antibody diluted at 1 : 1000

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