

## Product datasheet for TP307795

### HSPA6 (NM\_002155) Human Recombinant Protein

#### Product data:

Product Type:	Recombinant Proteins
Description:	Recombinant protein of human heat shock 70kDa protein 6 (HSP70B') (HSPA6), 20 µg
Species:	Human
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>RC207795 protein sequence Red=Cloning site Green=Tags(s)

MQAPRELAVGIDLGTTYSCVGVFQQGRVEILANDQGNRTTPSYVAFTDTERLVGDAAKSQAALNPHNTVF  
DAKRLIGRKFADTTVQSDMKHWPFRVWSEGGKPKVRVCYRGEDKTFYPEEISSMVLKMKETAAYLGQP  
VKHAVITVPAYFNDSQRQATKDAGAIAGLNLRIINEPTAAAIAYGLDRRGAGERNVLIFDLGGGTFDVS  
VLSIDAGVFEVKATAGDTHLGGEDFDNRLVNHFMEEFRKHGKDLSGNKRALRRLRTACERAKRTLSSST  
QATLEIDSLFEGVDFYTSITRARFEELCSDLFRSTLEPVEKALRDAKLDKAQIHADVVLVGGSTRIPKVQK  
LLQDFNKGKELNKSINPDEAVAYGAAVQAAVLMGDKCEKVQDLLLLDVAPLSLGLLETAGGVMTTLIQRNA  
TIPTKQTQFTTYSNDQPGVFIQVYEGERAMTKDNNLLGRFELSGIPPAPRGVPPQIEVTFDIDANGILSV  
TATDRSTGKANKITITNDKGRLSKEEVERMVHEAEQYKAEDAQRDRVAAKNSLEAHVFHVKGSLQEESL  
RDKIPEEDRRKMQDKCREVLAWLEHNQLAEKEEYEHQKRELEQICRPIFSRLYGGPGVPGGSSCGTQARQ  
GDPSTGPIIEVD

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag:	C-Myc/DDK
Predicted MW:	70.8 kDa
Concentration:	>0.05 µg/µL as determined by microplate BCA method
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol
Preparation:	Recombinant protein was captured through anti-DDK affinity column followed by conventional chromatography steps.
Note:	For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process.
Storage:	Store at -80°C.



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**Stability:** Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.

**RefSeq:** [NP\\_002146](#)

**Locus ID:** 3310

**UniProt ID:** [P17066](#), [A0A384NKX5](#), [B3KSM6](#)

**RefSeq Size:** 2664

**Cytogenetics:** 1q23.3

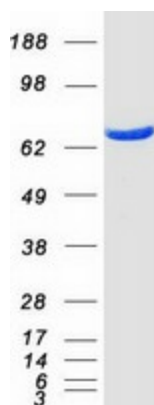
**RefSeq ORF:** 1929

**Synonyms:** HSP70B'

**Summary:** Molecular chaperone implicated in a wide variety of cellular processes, including protection of the proteome from stress, folding and transport of newly synthesized polypeptides, activation of proteolysis of misfolded proteins and the formation and dissociation of protein complexes. 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. This is achieved through cycles of ATP binding, ATP hydrolysis and ADP release, mediated by co-chaperones. The affinity 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. It goes through repeated cycles of ATP hydrolysis and nucleotide exchange, which permits cycles of substrate binding and release (PubMed:26865365).[UniProtKB/Swiss-Prot Function]

**Protein Pathways:** Antigen processing and presentation, Endocytosis, MAPK signaling pathway, Spliceosome

### Product images:



Coomassie blue staining of purified HSPA6 protein (Cat# TP307795). The protein was produced from HEK293T cells transfected with HSPA6 cDNA clone (Cat# [RC207795]) using MegaTran 2.0 (Cat# [TT210002]).