

## Product datasheet for **MR209729**

### Hspa8 (NM\_031165) Mouse Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Hspa8 (NM_031165) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Hspa8
Synonyms:	2410008N15Rik; Hsc70; Hsc71; Hsc73; Hsp73; Hspa10
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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**ORF Nucleotide  
Sequence:**

>MR209729 ORF sequence  
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**GCGATCGCC**

ATGTCTAAGGACCTGCAGTTGGCATTGATCTCGCACCACTACTCCTGTGTGGGTGTCTTCCAGCATG  
 GAAAGGTGGAAATTATTGCCAATGACCAGGTAACCGCACACCAAGCTATGTTGCTTTCACGGACAC  
 AGAGAGATTAATTGGGGATGCGGCCAAGAATCAGGTTGCAATGAACCCCAACACAGTTTTTGTGACC  
 AAACGTCTGATCGGGCGTAGGTTTGTGATGCTGTTGTTCACTGATATGAAGCACTGGCCCTTATGG  
 TGGTGAATGATGCAGGCAGGCCAAGGTCCAAGTGAATACAAAGGGGAGACAAAAGTTTCTACCCAGA  
 GGAAGTGTCTCCATGTTCTGACAAAGATGAAGGAAATGCAGAAGCGTACCTCGGAAAGACTGTTACC  
 AACGCTGTGGTCACAGTCCCGCTTACTTCAATGACTCTACGCGACAGGCCAACAAAAGATGCTGAACTA  
 TTGCTGGCCTCAATGTACTTCAATCATCAATGAACCAACTGCTGCTGCTATTGCTTATGGCTTAGATAA  
 GAAGGTCGGAGCTGAAAGGAATGTGCTCATTTTTGACTTGGGAGGTGGCACTTTTGTGTCAATCCTC  
 ACTATTGAGGATGGAATTTTTGAGGTCAAATCAACAGCTGGAGACACCCACTTAGGTGGAGAAGATTTG  
 ACAACCGAATGGTCAATCATTTTCTGAGTTCAAGCGAAAGCACAAGAAAGACATCAGTGAGAACA  
 GAGAGCTGTCCGCGTCTCCGCACGGCCTGCGAGCGGGCAAGCGCACCCCTCTCCTCCAGCACCCAGGCC  
 AGTATTGAGATTGATTCTCTATGAGGGAATTGACTTCTATACCTCCATTACCCGGGCTCGATTGAGG  
 AGTTGAATGCTGACCTGTTCCGTGGCACACTGGACCCTGTAGAGAAGGCCCTTCGAGATGCCAAGCTGGA  
 CAAGTCACAGATCCATGATATTGCTTGGTGGGTGTTTCTACCAGAATCCCAAGATTGAGAACTTCTG  
 CAAGACTTCTCAATGGAAAAGAGCTGAACAAGAGCATTAAACCCGATGAAGCTGTTGCCTATGGTGCAG  
 CTGTCCAGGCAGCCATTCTATCTGGAGACAAGTCTGAGAAGCTTCAAGATTGCTGCTCTTGGATGTCAC  
 TCCTCTTTCCCTTGGTATTGAAACTGCTGGCGGAGTCATGACTGTCCTCATCAAGCGCAATACCACCATC  
 CCCACCAAGCAGACAGACTTTACCACCTACTCTGACAACCAGCCTGGTGTACTCATTCAAGGTGTATG  
 AAGGTGAAAGGGCCATGACCAAGGACAACAACCTGCTTGGAAAGTTCGAGCTCACAGGCATCCCTCCAGC  
 ACCCCGTGGGTCCCTCAGATTGAGGTTACTTTTGACATCGATGCCAATGGCATCCTCAATGTTTCTGCT  
 GTAGATAAGAGCACAGGAAAGGAGAACAAGATCACCATCACCAATGACAAGGGCCGCTTGAGTAAGGAAG  
 ATATTGAGCGCATGGTCCAAGAAGCTGAGAAGTACAAGGCTGAGGATGAGAAGCAGAGAGATAAGGTTTC  
 CTCCAAGAACTCACTGGAGTCTATGCCTCAACATGAAAGCAACTGTGGAAGATGAGAACTTCAAGGC  
 AAGATCAATGATGAGGACAAACAGAAGATTCTTGACAAGTCAATGAAATCATCAGCTGGCTGGATAAGA  
 ACCAGACTGCAGAGAAGGAAGATTTGAGCATCAGCAGAAAGAACTGGAGAAAGTCTGCAACCCTATCAT  
 TACCAAGCTGTACCAGAGTGCAGGTGGCATGCCTGGGGGAATGCCTGGTGGCTTCCAGGTGGAGGAGCT  
 CCCCCATCTGGTGGTCTTCTCAGGCCACCATTGAAGAGGTGGAT

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Protein Sequence:** >MR209729 protein sequence  
 Red=Cloning site Green=Tags(s)

MSKGPVAVGIDLGTYSVGVFQHGKVEIIANDQGNRTTPSYVAFDTERLIGDAAKNQVAMNPTNTVFDA  
 KRLIGRRFDDAVVQSDMKHWPFMVNDAGRPKVQVEYKGETKSFYPPEVSSMVLTKMKEIAEAYLGKTVT  
 NAVVTVPAYFNDSQRQATKDAGTIAGLNVLRIINEPTAAAIAAYGLDKKVGAEARNVLIIDLGGTDFDVSIL  
 TIEDGIFEVKSTAGDTHLGGEDFDRMNVNHFIAEFKRKHKKDISENKRAVRRRLTACERAKRTLSSSTQA  
 SIEIDSLYEGIDFYTSITRARFEELNADLFRGTLDPVEKALRDAKLDKSQIHDIVLVGGSTRIPKIQKLL  
 QDFNFKELNKSINPDEAVAYGAAVQAAILSGDKSENVQDLLLVDVPLSLGIETAGVMVTLIKRNTTI  
 PTKQTQFTTYSNQPGLVIQVYGERAMTKDNNLLGKFELTGIPPAPRGVQPQIEVTFDIDANGILNVSA  
 VDKSTGKENKIIITNDKGRLSKEDIERMVQEAKEYKADEKQRDKVSSKNSLESYAFNMKATVEDEKLQG  
 KINDEDKQKILDKCNELIISWLDKNQTAKEEFEHQQKELEKVCNPITIKLYQSAGMPPGGMPGGFPPGGGA  
 PPSGGASSGPTIEEVD

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

**Restriction Sites:**

Sgfl-MluI

**Cloning Scheme:**



**ACCN:** NM\_031165

**ORF Size:** 1941 bp

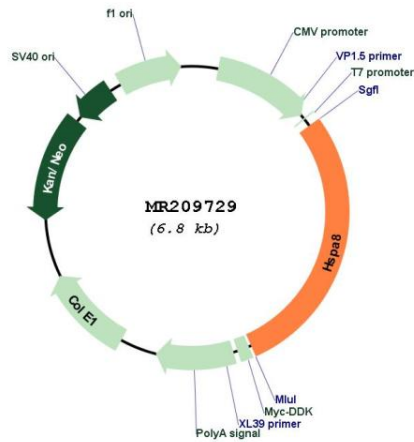
**OTI Disclaimer:** The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_031165.3</a>
<b>RefSeq Size:</b>	2104 bp
<b>RefSeq ORF:</b>	1941 bp
<b>Locus ID:</b>	15481
<b>UniProt ID:</b>	<a href="#">P63017</a>
<b>Cytogenetics:</b>	9 21.55 cM
<b>MW:</b>	70.9 kDa
<b>Gene Summary:</b>	<p>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 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. 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. 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. Participates in the ER-associated degradation (ERAD) quality control pathway in conjunction with J domain-containing co-chaperones and the E3 ligase STUB1.[UniProtKB/Swiss-Prot Function]</p>

Product images:



Circular map for MR209729