

Product datasheet for **SC204035**

HSP90AB1 (NM_007355) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: HSP90AB1 (NM_007355) Human 3' UTR Clone
Symbol: HSP90AB1
Synonyms: D6S182; HSP84; HSP90B; HSPC2; HSPCB
Mammalian Cell Selection: Neomycin
Vector: pMirTarget (PS100062)
ACCN: NM_007355
Insert Size: 315 bp
Insert Sequence: >SC204035 3'UTR clone of NM_007355
The sequence shown below is from the reference sequence of NM_007355. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GATGCGTCTCGCATGGAAGAAGTCGATTAGGTTAGGAGTTCATAGTTGGAAAAGTGTGCCCTTGTATA
GTGTCCCATGGGCTCCCACTGCAGCCTCGAGTGCCCTGTCCACCTGGCTCCCTGCTGGTGTCTA
GTGTTTTTTTCCCTCTCCTGTCTTGTGTTGAAGGCAGTAACTAAGGGTGTCAAGCCCCATTCCCTCT
CTACTCTTGACAGCAGGATTGGATGTTGTGTTATTGGTTTATTTTATTTCTTCATTTTGTCTGAAA
TTAAAGTATGCAAAATAAAGAATATGCCGTTTTTATACA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAAGAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.

RefSeq: [NM_007355.4](#)



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Summary: This gene encodes a member of the heat shock protein 90 family; these proteins are involved in signal transduction, protein folding and degradation and morphological evolution. This gene encodes the constitutive form of the cytosolic 90 kDa heat-shock protein and is thought to play a role in gastric apoptosis and inflammation. Alternative splicing results in multiple transcript variants. Pseudogenes have been identified on multiple chromosomes. [provided by RefSeq, Dec 2012]

Locus ID: 3326

MW: 11.6