

Product datasheet for **SC205753**

DNAJB11 (NM_016306) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: DNAJB11 (NM_016306) Human 3' UTR Clone
Vector: pMirTarget (PS100062)
Symbol: DNAJB11
Synonyms: ABBP-2; ABBP2; Dj-9; DJ9; EDJ; ERdj3; ERj3; ERj3p; PKD6; PRO1080; UNQ537
ACCN: NM_016306
Insert Size: 415 bp
Insert Sequence: >SC205753 3'UTR clone of NM_016306
The sequence shown below is from the reference sequence of NM_016306. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
AAGGTATACAATGGACTGCAAGGATATTGAGAGTGAATAAAATTGGACTTTGTTTAAATAAGTGAATA
AGCGATATTTATTATCTGCAAGTTTTTTGTGTGTTTTTTTATTTTCAATATGCAAGTTAGG
CTTAATTTTTTTATCTAATGATCATCATGAAATGAATAAGAGGGCTTAAGAATTTGTCCATTTGCATTC
GGAAAAGAATGACCAGCAAAAGTTTTACTAATACCTCTCCCTTTGGGGATTTAATGTCTGGTGCTGCCG
CCTGAGTTTCAAGAATTAAGCTGCAAGAGGACTCCAGGAGCAAAAGAAACACAATATAGAGGGTTGGA
GTTGTTAGCAATTTATTCAAAATGCCAACTGGAGAAGTCTGTTTTTAAATACATTTTGTGTTATTTTT
ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTTTCGATTCCACCGCCCTTCTATGAAAGG
```

Restriction Sites: SgfI-MluI

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_016306.6](#)



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Summary: This gene encodes a soluble glycoprotein of the endoplasmic reticulum (ER) lumen that functions as a co-chaperone of binding immunoglobulin protein, a 70 kilodalton heat shock protein chaperone required for the proper folding and assembly of proteins in the ER. The encoded protein contains a highly conserved J domain of about 70 amino acids with a characteristic His-Pro-Asp (HPD) motif and may regulate the activity of binding immunoglobulin protein by stimulating ATPase activity. [provided by RefSeq, Mar 2014]

Locus ID: 51726

MW: 16.1