

Product datasheet for **SC206186**

MADM (NRBP1) (NM_013392) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	MADM (NRBP1) (NM_013392) Human 3' UTR Clone
Symbol:	MADM
Synonyms:	BCON3; MADM; MUDPNP; NRBP
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_013392
Insert Size:	477 bp
Insert Sequence:	<p>>SC206186 3'UTR clone of NM_013392</p> <p>The sequence shown below is from the reference sequence of NM_013392. The complete sequence of this clone may contain minor differences, such as SNPs.</p> <p>Blue=Stop Codon Red=Cloning site</p>

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GGCAAGTTGGACGCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAACGATCGCC
AACTCAGCCGCTGTCACCGTCTCCTCTAGAGCTCACTCGGGCCAGGCCCTGATCTGCGCTGTGGCTGT
CCCTGGACGTGCTGCAGCCCTCCTGTCCCTTCCCCCAGTCAGTATTACCCTGTGAAGCCCTTCCCTC
CTTTATTATTAGGAGGCTGGGGGGGCTCCCTGGTTCTGAGCATCATCCTTTCCCTCCCTCTCTTC
CTCCCTCTGCACTTTGTTTACTTGTGTTTGCACAGACGTGGGCCTGGGCCTTCTCAGCAGCCGCTTCT
AGTTGGGGGCTAGTCGCTGATCTGCCGGCTCCCGCCAGCCTGTGTGAAAGGAGGCCACGGGCACTA
GGGGAGCCGAATTCTACAATCCCGCTGGGGCGGGCGGGGAGAGAAAGGTGGTGTGCTGCAAGTGGTGG
CCCTGGGGGGCCATTGATTGCGCTCAGTTGCTGCTGTAATAAAAGTCTACTTTTGTCTAAA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCGCCTTCTATGAAAGG
  
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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.


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RefSeq: [NM_013392.4](#)

Summary: May play a role in subcellular trafficking between the endoplasmic reticulum and Golgi apparatus through interactions with the Rho-type GTPases. Binding to the NS3 protein of dengue virus type 2 appears to subvert this activity into the alteration of the intracellular membrane structure associated with flaviviral replication.[UniProtKB/Swiss-Prot Function]

Locus ID: 29959

MW: 16.8