

Product datasheet for **SC207936**

MTMR1 (NM_003828) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: MTMR1 (NM_003828) Human 3' UTR Clone
Symbol: MTMR1
Mammalian Cell Selection: Neomycin
Vector: pMirTarget (PS100062)
ACCN: NM_003828
Insert Size: 586 bp
Insert Sequence: >SC207936 3'UTR clone of NM_003828
 The sequence shown below is from the reference sequence of NM_003828. The complete sequence of this clone may contain minor differences, such as SNPs.
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAACGATCGCC
TCCGCCACCTCCGTCCACACCTCGGTCTTATGGGCGAGAAATATGTAATCCCCTGGCTGACTAGGACTG
TTAAACATAGTGTGGACTGGATGATGCCTTCGACAAACCAGAGAAGCCAAGTTGGGGGAGCTGGTGCC
TGGAGTGGGCCCTGTGCACCTCACCTGGCGGAGGCTGGGGGGGCTCTGTCAGCAGGACCCTAGAGGAG
ACTCTCATTTCGATTTTAAAGAAGCACACGGGTCATTTTCCTTTGTATGTTCTAGCGCAGAACTGTTT
CTAAAACAACCTGAAGTATAGTTTGTATCTAAGCAATTTTGTTTAAGTAAGTAAGTGTACTAGAA
TGCGAAGCCGTTATGGTTCAGGTTTTTAAAACTGGTACAGTATTGTATTGTCTCATCTGTTGCACTG
TATTTCAATCATCTGTAATTAATGATCATATGTTTGCTCCCTGGTCTTTTTAAGTAAGTAAGTAAG
TATCTTAGTAGATTTTCTTTGAGGAAAATCGGTAATAAAATAACATGGATTGAATGTTTACTGTGCG
TCAAGCACAGTTAATATATGATGATGTAAAGTAA
ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG
  
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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).


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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_003828.5
Summary:	This gene encodes a member of the myotubularin related family of proteins. Members of this family contain the consensus sequence for the active site of protein tyrosine phosphatases. Alternatively spliced variants have been described but their biological validity has not been determined. [provided by RefSeq, Jul 2008]
Locus ID:	8776
MW:	21.7