

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
Vector: pMirTarget (PS100062)
Symbol: MTMR1
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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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
TCCGCCACCTCCGTCCACACCTCGGTCTGATGGGCGAGAAATATGTAATCCCTGGCTGACTAGGACTG
TTAAACATAGTGTGGACTGGATGATGCCTTCGACAAACCAGAGAAGCCAAGTTGGGGGGAGCTGGTGCC
TGGAGTGGGCCCTGTGCACCTCACCTGGCGGAGGCTGGGGGGGGCTCTGTTCAGCAGGACCCCTAGAGGAG
ACTCTCATTTCGATTTTAAAGAAGCACAACGGGTCAATTTTCCTTTGTATGTTCTAGCGCAGAAGTGTTC
CTAAAACAACCTGAAGTATAGTTTTGTTATCTAAGCAATTTTGTTTAAGTAAGTAAGTACTAGAA
TGCGAAGCCGTTATGGTTCAGGTTTTTAAAACTGGTACAGTATTGTATTTGTCTCATCTGTTGCACTG
TATTTCAATCATCTGTAATTAATGATCATATGTTTGCCTCCCTGGTCTTTTTAAGTAAGTAAGTAAG
TATCTTAGTAGATTTTTCCTTTGAGGAAAATCGGTAATAAAATAACATGGATTGAATGTTTACTGTGCG
TCAAGCACAGTTAATATATGATGATGTAAGTAA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTTTCGATTCCACCGCCCTTCTATGAAAGG
```

Restriction Sites: Sgfl-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 µg 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](#)



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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