

Product datasheet for **SC114002**

TRM1 (TRMT1) (NM_017722) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TRM1 (TRMT1) (NM_017722) Human Untagged Clone
Tag:	Tag Free
Symbol:	TRM1
Synonyms:	MRT68; TRM1
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >OriGene ORF within SC114002 sequence for NM_017722 edited (data generated by NextGen Sequencing)

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ATGCAAGGATCGTCTCTGTGGCTAAGCCTCACTTCCGCTCCGCCCGGGTCTCTCTAGA
GCCCGTTTTTCGAGTGGCAGTCTCCAGGGCTGCCGAATACAGCAGCGATGGAGAACGGC
ACCGGGCCCTACGGAGAAGAACGTCCACGTGAAGTCCAGGAGACGACAGTCAACGAGGGG
GCTGCCAAAATCGCCTTTCCAGTGCCAACGAGGTCCTTTATAACCCGGTGCAGGAATTC
AATCGGGACCTGACATGTGCTGTGATCACCGAGTTTGCTCGCATTACAGTTGGGGCCAAA
GGAATCCAGATCAAGTTCCAGGAGAGAAGGACACGCAAAAAGTGGTCGTGGACTTGTCA
GAGCAAGAGGAGGAAAAGGTTGAACTGAAAGAGAGTGAAAACCTGGCCTCAGGAGACCAA
CCTCGCACAGCGCCGTGGGGGAGATCTGTGAGGAAGGCCTGCATGTGCTGGAAGGCCTG
GCAGTTCAGGCCTACGTTCCATTTCGATTTGCCCTAGAGGTGCCTGGGCTCAGATCTGTG
GTTGCAAACGATGCCTCCACCCGGGCTGTGGATCTCATAACCCGGAATGTCCAGCTCAAT
GACGTGGCCACCTGGTACAGCCGAGCCAAGCAGATGCCCGGATGCTGATGTACCAGCAC
CAGAGGGTGTGAGAGAGGTTTGACGTATCGATCTGGACCCCTATGGCAGCCAGCCACC
TTCTGGATGCAGCTGTGCAGGCTGNNNGTGCAGGAGGGTTGCTGTGTGACCTGCACA
GACATGGCGGTGTGGCGGGGAACAGCGGGGAGACGTGCTACAGCAAGTACGGGGCCATG
GCCCTCAAGAGCCGGGCCTGCCACGAGATGGCCCTGAGAATCGTCTGCACAGCCTGGAC
CTCCGCGCAACTGCTACCAGCGCTTCGTGGTGCCGCTGCTCAGCATCAGCGTGACTTC
TACGTGCGTGTTCCTGCGTCTTACCCGGCCAGGCCAAGGTCAAGGCCTCAGCCAGC
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AAAGCGTCAGGAGTCCCCAGCGGGCCGGAAGTTCTCTGAGCCTGTGGTCCCCCTGTG
ACCCCCAGTGTGAACACTGTGGGCAACGACACCAGCTTGGTGGCCCCATGTGGGAGAG
CCCATCCATGACCTGGATTTTGTGGCCGTGTCCTGGAGGCTGTGAGCGTAACCCCGGC
CGTTCCACACCTCGGAGCGGATCCGAGGGGTCCTGAGCGTCATCACTGAGGAGCTCCCG
GACGTGCCTCTGTACTACACCCTGGACCAGCTGAGCAGCACCATCCACTGCAACACACCA
AGCCTCTGCAGTTGCGGTGCGCCCTCCTCCACGCTGACTTCCGGGTCTCACTCTCCAC
GCCTGTAAGAACGCTGTGAAGACGGATGCCCTGCCTCTGCCCTCTGGGACATCATGCGT
TGCTGGGAGAAGGAATGTCGGTGAAACGGGAGCGACTATCAGAGACTAGCCAGCGTTC
CGCATTCTCAGTGTGGAGCCAGGCTGCAGGCCAACTTACCATCCGGGAAGATGCCAAC
CCCAGCTCCCAGCAGGAGGACTCAAGCGCTTCCAGGCTAACCCGGAGGCCAACTGGGGT
CCCCGGCCTCGTCCCGCCAGGGGGCAAGGCGGCCGACGAAGCTATGGAGGAGAGACGC
AGGCTGCTTCAAGAACGCGGAAGGAGCCGCGGAAGATGTGGCCAGCGGGCTGCCCGG
CTCAAGACATTTCTTGAAGAGGTTTAAAGGAGGGCACCTGTCAACGCGGGGACCAAGTGC
TGCTACTCCCACAGCCCCCGACACCCAGGGTTTCTGCTGATGCTGCCCTGACTGTCCA
GAGACCTCAACCCAGACCCCTGGACCTGGGGCTGCCGCTGGGCCAGGCATAGACTGA
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Clone variation with respect to NM_017722.3

746 t=>n;747 g=>n;748 a=>n;752 a=>c

5' Read Nucleotide Sequence:	<p>>OriGene 5' read for NM_017722 unedited</p> <p>CACATTTGTATACGACTCACTTATAGGGCGGCCGGAATTCGCACGAGGGNAAACGTGCA CCTTGACTAATTCGGTTTTCTGTGGCTACGAGAGCAGGCTTGGCGGGCGGAGGCCAGC GGATGTCTCATGCAAGGATCGTCTCTGTGGCTAAGCCTCACTTCCGCTCCGCCCGGGT CTCTCTAGAGCCCGTTTTTCGAGTGGCAGTCTCCAGGGCTGCCGAATACAGCAGCGATG GAGAACGGCACCAGGGCCCTACGGAGAAGAAGCTCCACGTGAAGTCCAGGAGACGACAGTC ACCGAGGGGGCTGCCAAAATCGCCTTTCCAGTGCCAACGAGGTCTTTTATAACCCGGTG CAGGAATTC AATCGGGACCTGACATGTGCTGTGATCACCGAGTTTGCTCGCATTACGCTT GGGGCCAAAGGAATCCAGATCAAGGTTCCAGGAGAGAAGGACACGAAAAAGTGGTCGTG GACTTGTGAGAGCAAGAGGAGGAAAAGTTGAACTGAAAGAGAGTGAAAACCTGGCCTCA GGAGACCAACCTCGCACAGCGGCCGTGGGGGAGATCTGTGAGGAAGGCCTGCATGTGCTG GAAGGCCTGGCAGCTTACGGCCTACGTTCCATTGATTTGCCCTAGAGGTGCCTGGGCTC AGATCTGTGGTTGCAAACGATGCCTCCACCCGGGCTGTGGATCTCATACGCCGAATGTC CAGCTCAATGACGTGGCCACCTGGTACAGCCGAGCCAAGCAGATGCCCGGATGCTGATG TACCAGCACCAGNAGGTGTCGGAGAAGTTGACGTCATCGATCTGGACCCTATGGCAGC CCAGCCACCTTCCGTGATGCANCTGTGCANGCTGGNTGCTGTGTGTGACCTGCACGACAT GNNCGTGTTNNGCGGNGACANCGNGNAGACGTGCTACAGCAGTACGGNGCCATGGNCCTC CAGACNNGGCCTN</p>
3' Read Nucleotide Sequence:	<p>>OriGene 3' read for NM_017722 unedited</p> <p>NNNNNTTACTTGNACCCGGCCGATNCTANNGATCGGTTTTTTTTTTTTTTTTTTTTTTT TTTTTTTTGACGTGACATCTCTTTATTGGTTCAGTCTATGCCTGGCCAGCGGCAGCCCC AGGTCCAGGGGGGTCTGGTTGGAGTCTCTGGACAGTACAGGGGCAGCATCAGCAGAAAC CCTGGGTGTGGGGGGTGTGGGAGTAGCAGCACTGGTCCCCGCGTTGACAGGTGCCCTC CTTAAACCTCTTGAAGGAAATGTCTTGAGCCGGGCAGCCGCTGGGCCACATCTCCGG CGGCTCCTCCGCTTGTCTGAAGCAGCCTGCCTCTCTCCATAGCTTCGTGGCCGC CTTGCCCTGGCCGGCAGGAGCCGGGGACCCAGTTGGCCTCCGGGTTAGCCTGGAA GCGCTTGAGTCTCGCTGTGGGAGCTGGGGTTGGCATCTCCCGGATGGTGAAGTTGGC CTGCAGCCTGGGCTCCACTGAGAATGCGGAACGCTGGGCTAGTCTCTGATAGTCGCTC CGTTTTACCCGGACATTCCTTCTCCAGCAACGCATGATGTCCAGAGGGCAAAGGCAGG GGCATCCGTCTTACAGGTTCTTACAGCGTGGGAGAGTGAGACCCGGAAGTCAGCGTG GAGGAGGGCCGACCGCAACTGCAGGAGGCTTGGTGTGTTGCAAGTGGATGGTGTGCTCAN CTGGTCCAGGGTGTAGTACAGAGGCACGTCCGGGAGCTCCTGCGATGGGGGACAGGATGG GCATGAGTGGAGATGGACCGGCACAGGTGGAGCCCCCTCCCGTACAGGCCGTTCTC ACCTTAGAGATGACGCTCAGGACCCCTCGGATCCGGTCCCGAGTGTGGAAGCGCCCGGG T</p>
Restriction Sites:	NotI-NotI
ACCN:	NM_017722
Insert Size:	2640 bp
OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:	<ol style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	NM_017722.2 , NP_060192.1
RefSeq Size:	2217 bp
RefSeq ORF:	1980 bp
Locus ID:	55621
UniProt ID:	Q9NXH9
Cytogenetics:	19p13.13
Domains:	zf-CCCH, TRM
Protein Families:	Druggable Genome
Gene Summary:	<p>This gene encodes a tRNA-modifying enzyme that acts as a dimethyltransferase, modifying a single guanine residue at position 26 of the tRNA. The encoded enzyme has both mono- and dimethylase activity when exogenously expressed, and uses S-adenosyl methionine as a methyl donor. The C-terminal region of the encoded protein has both a zinc finger motif, and an arginine/proline-rich region. Mutations in this gene have been implicated in autosomal recessive intellectual disorder (ARID). Alternative splicing results in multiple transcript variants encoding different isoforms. There is a pseudogene of this gene on the X chromosome. [provided by RefSeq, May 2017]</p> <p>Transcript Variant: This variant (1) is the longest transcript and encodes the longer isoform (1).</p>