

Product datasheet for SC309934

MTR (NM_000254) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: MTR (NM_000254) Human Untagged Clone
Tag: Tag Free
Symbol: MTR
Synonyms: cbIG; HMAG; MS
Mammalian Cell Selection: Neomycin
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM_000254 edited
 ACGTCGCTCCTCTGCGGTTTTCTCTTGGGTCCTTTCCGTGCCGTCCTCCGCGACTCCG
 CCTCTGGCCGCGGTGTCTGGCTGCTAGGCCGACACCAAGGACTGGCCGGTACCCGGGA
 AGAAAGCACGTGCTCCAGCAGTTGCCGCGCCAGCCCCGAGAGAGGCCCTAGGGCGCTGC
 GGGCTTTCGGGGTCCGCGAGTCCCCCGCGACGCGAGCCAACGGGAGGCGTCAAAAGACCC
 GGGCCTTGTGTGGCAGGCTCGCCTGGCGCTGGCTGGCGTGGCCCTTGGCCGTCGTACCT
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CAAAGTGCAGGTGAAATGGGAGCCCAGGTGTTGGATGTCAACATGGATGATGGCATGCT
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 ACTTACCTGCTTCTGGTTTTTGAAGACTATTTAGTGAACCTTGTAGAGGAGCAGGGTCT
 TCCTGCAGTGCCTGGAACAGGCGCTGTTTTTTTTTGGGACCTTGCCTGAAGAGCAGTG
 AGCAGGGTTCCTGTGTTTTCCCTGGTCCCTCTGAGATGGGGACAGACTGAAGACAGAGGT
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 GGTGAGGAAATATGAATTAGGACCACAATGAGATTCCATTTTATATCCATAAGATTTGC
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 TGTGTAACAGGCACCCTGCTTTAAAAACAATTATCCCTTACAGACTTGAACATTTGC
 AGACGTTATGATCTTGTCCCACTCCACCTGTATGTCCAGCAAACTCTTGATGTGGC
 CACTAGGAGGAATGTGTAAGAATGTTTATAGTTACATATTTATAATAGTTAATAACTGGA
 AAAAGTGAATGTATGTCTGTCTACAGGAAAAAGGTGAATAATTAGATATATATATTCA
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 TATTTTTATTGAGCCAAAAACAAGCAAAACCAAGAATATGTAGTCTAAGCATACGTAT
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 CAGCTGTTCTGGTCTCCTGGTAGGCTTACAAGTGTACTATATGCTATTAATACATT
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 TTATAAAAAATAAAGGCAAGTGAATGAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

Restriction Sites: Please inquire
ACCN: NM_000254
Insert Size: 7200 bp

OTI Disclaimer:	<p>Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.</p> <p>The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. More info</p>
OTI Annotation:	The open reading frame of this TrueClone was fully sequenced and found to differ from the protein associated to this reference by two amino acids.
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_000254.1 , NP_000245.1
RefSeq Size:	7122 bp
RefSeq ORF:	3798 bp
Locus ID:	4548
UniProt ID:	Q99707
Cytogenetics:	1q43
Domains:	Pterin_bind, S-methyl_trans, B12-binding_2, Met_synt_B12, B12-binding
Protein Families:	Druggable Genome
Protein Pathways:	Cysteine and methionine metabolism, Metabolic pathways, One carbon pool by folate

Gene Summary:

This gene encodes the 5-methyltetrahydrofolate-homocysteine methyltransferase. This enzyme, also known as cobalamin-dependent methionine synthase, catalyzes the final step in methionine biosynthesis. Mutations in MTR have been identified as the underlying cause of methylcobalamin deficiency complementation group G. Alternatively spliced transcript variants encoding distinct isoforms have been found for this gene. [provided by RefSeq, May 2014]

Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1). Sequence Note: This RefSeq record was created from transcript and genomic sequence data because no single transcript was available for the full length of the gene. The extent of this transcript is supported by transcript alignments.