

## Product datasheet for **SC314547**

### MTRR (NM\_024010) Human Untagged Clone

#### Product data:

|                           |                                       |
|---------------------------|---------------------------------------|
| Product Type:             | Expression Plasmids                   |
| Product Name:             | MTRR (NM_024010) Human Untagged Clone |
| Tag:                      | Tag Free                              |
| Symbol:                   | MTRR                                  |
| Synonyms:                 | cbIE; MSR                             |
| Mammalian Cell Selection: | None                                  |
| Vector:                   | <u><a href="#">pCMV6-XL5</a></u>      |
| E. coli Selection:        | Ampicillin (100 ug/mL)                |



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**Fully Sequenced ORF:** >OriGene sequence for NM\_024010 edited  
 GGAGCTTTCTATTGGTCTGGGTACCGAGCATGGGCGCTGCGTCAGTGC GCGCTGGCGCA  
 AGGTTGGTGAAGTCGCGTTGTGCGATTTCACTGTTACATGCCTTGAAGTGATGAGGAGG  
 TTTCTGTTACTATATGCTACACAGCAGGGACAGGCAAAGGCCATCGCAGAAGAAATATGT  
 GAGCAAGCTGTGGTACATGGATTTCTGCAGATCTTCACTGTATTAGTGAATCCGATAAG  
 TATGACCTAAAAACCGAAACAGCTCCTCTTGTGTTGTGGTTTCTACCACGGGCACCGGA  
 GACCCACCCGACACAGCCCGCAAGTTTGTAAAGAAATACAGAACCAAACACTGCCGGTT  
 GATTTCTTTGCTCACCTGCGGTATGGGTTACTGGGTCTCGGTGATTCAGAATACACCTAC  
 TTTTGCAATGGGGGAAGATAATTGATAAACGACTTCAAGAGCTTGGAGCCCGGCAATTC  
 TATGACACTGGACATGCAGATGACTGTGTAGGTTTAGAATTGTGGTTGAGCCGTGGATT  
 GCTGGACTCTGGCCAGCCCTCAGAAAGCATTTTAGGTCAAGCAGAGGACAAGAGGAGATA  
 AGTGGCGCACTCCCGTGGCATCACCTGCATCCTTGGAGCAGACCTTGTGAAGTCAGAG  
 CTGCTACACATTGAATCTCAAGTCGAGCTTCTGAGATTCGATGATTCAGGAAGAAAGGAT  
 TCTGAGGTTTTGAAGCAAATGCAGTGAACAGCAACCAATCCAATGTTGAATTGAAGAC  
 TTTGAGTCTCACTTACCCGTTCCGTACCCCACTCTCACAAGCCTCTCTGAATATTCCT  
 GGTTTACCCCAAGATATTTACAGGTACATCTGCAGGAGTCTCTTGGCCAGGAGGAAAGC  
 CAAGTATCTGTGACTTCAGCAGATCCAGTTTTTCAAGTGCCAATTTCAAAGGCAGTTCAA  
 CTTACTACGAATGATGCCATAAAAACTCTGCTGGTAGAATTGGACATTTCAAATACA  
 GACTTTTCTATCAGCCTGGAGATGCCTTACAGCGTGATCTGCCCTAACAGTGATTCTGAG  
 GTACAAAGCCTACTCCAAAGACTGCAGCTTGAAGATAAAAGAGAGCACTGCGCTCTTTTG  
 AAAATAAAGGCAGACACAAAGAAGAAAGGAGCTACCTTACCCAGCATATACCTGCGGGA  
 TGTCTCTCCAGTTCATTTTTACCTGGTGTCTTGAATCCGAGCAATTCCTAAAAAGGCA  
 TTTTTCGAGCCCTTGTGGACTATACCAGTGACAGTGCTGAAAAGCGCAGGCTACAGGAG  
 CTGTGCAGTAAACAAGGGCAGCCGATTATAGCCGCTTTGTACGAGATGCCTGTGCCTGC  
 TTGTTGGATCTCCTCCTCGCTTCCCTTCTTGCCAGCCCACTCAGTCTCCTGCTCGAA  
 CATCTTCTAACTTCAACCCAGACCATATTCGTGTGCAAGCTCAAGTTTATTTACCCA  
 GGAAAGCTCCATTTTGTCTTCAACATTGTGGAATTTCTGTCTACTGCCACAACAGAGGT  
 CTGCGGAAGGGAGTATGTACAGGCTGGCTGGCCTTGTGGTTGCTTCAGTTCTTCAGCCA  
 AACATACATGCATCCCATGAAGACAGCGGAAAGCCCTGGCTCCTAAGATATCCATCTCT  
 CCTCGAACAAATTTTCCACTTACCAGATGACCCCTCAATCCCATCATAATGGTG  
 GGTCCAGGAACCGGCATAGCCCGTTTATTGGGTTCTACAACATAGAGAGAACTCCAA  
 GAACAACCCAGATGGAATTTTGGAGCAATGTGGTTGTTTTTGGCTGCAGGCATAAG  
 GATAGGGATTATCTATTAGAAAAGAGCTCAGACATTTCTTAAGCATGGGATCTTAACT  
 CATCTAAAGGTTTCTTCTCAAGAGATGCTCCTGTTGGGGAGGAGGAAGCCCAAGAAAG  
 TATGTRCAAGACAACATCCAGCTTATGGCCAGCAGGTGGCAAGAATCCTCCTCCAGGAG  
 AACGGCCATATTTATGTGTGTGGAGATGCAAAGAATATGGCCAAGGATGTACATGATGCC  
 CTTGTGCAAATAAAGCAAAGAGGTTGGAGTTGAAAACTAGAAGCAATGAAAACCCCTG  
 GCCACTTTAAAAGAAGAAAAACGCTACCTTCCAGGATATTTGGTCAAAAACAGAAATTA  
 AAGAAAGAGGATTAAGCTTTTTGACTGAAAGTACTAAAAGTCAAGCTTTACTAGTGCCAA  
 ACCTTTAAATTTCAAAGAAAATTTCTTTCAACATTTCTTGAAGGACATGGAGTGGAG  
 ATTGGATCATTTAACAATAAACAATAAACAATTTCTGATTTGATTTTACGTATCTTCTATCTA  
 CGCCCTTCTGTGCCTGTGACTCTCCCA

**Restriction Sites:** Please inquire

**ACCN:** NM\_024010

**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).

|                               |  |
|-------------------------------|--|
| <b>OTI Annotation:</b>        | The open reading frame of this TrueClone was fully sequenced and found to differ from the protein associated to this reference by a single amino acid.   |
| <b>Components:</b>            | 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).   |
| <b>Reconstitution Method:</b> | <ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>  |
| <b>RefSeq:</b>                | <u><a href="#">NM_024010.1</a></u> , <u><a href="#">NP_076915.1</a></u>  |
| <b>RefSeq Size:</b>           | 3291 bp  |
| <b>RefSeq ORF:</b>            | 2178 bp  |
| <b>Locus ID:</b>              | 4552   |
| <b>UniProt ID:</b>            | <u><a href="#">Q9UBK8</a></u>  |
| <b>Cytogenetics:</b>          | 5p15.31  |
| <b>Domains:</b>               | flavodoxin, NAD_binding_1, FAD_binding_1   |
| <b>Protein Families:</b>      | Druggable Genome   |
| <b>Gene Summary:</b>          | <p>This gene encodes a member of the ferredoxin-NADP(+) reductase (FNR) family of electron transferases. This protein functions in the synthesis of methionine by regenerating methionine synthase to a functional state. Because methionine synthesis requires methyl-group transfer by a folate donor, activity of the encoded enzyme is important for folate metabolism and cellular methylation. Mutations in this gene can cause homocystinuria-megaloblastic anemia, cbl E type. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Dec 2015]</p> <p>Transcript Variant: This variant (2) differs in the 5' UTR compared to variant 6. Variants 1, 2, 6, 7, and 8 all encode the same protein. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.</p> |