

## Product datasheet for RC214275L1V

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## MTR (NM\_000254) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

Product Type: Lentiviral Particles

**Product Name:** MTR (NM\_000254) Human Tagged ORF Clone Lentiviral Particle

Symbol: MTF

**Synonyms:** cblG; HMAG; MS

Mammalian Cell

Selection:

None

**Vector:** pLenti-C-Myc-DDK (PS100064)

 Tag:
 Myc-DDK

 ACCN:
 NM\_000254

 ORF Size:
 3795 bp

**ORF Nucleotide** 

OTI Disclaimer:

Sequence:

The ORF insert of this clone is exactly the same as(RC214275).

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

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**RefSeg:** NM 000254.1

 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





## MTR (NM\_000254) Human Tagged ORF Clone Lentiviral Particle - RC214275L1V

**Protein Pathways:** Cysteine and methionine metabolism, Metabolic pathways, One carbon pool by folate

MW: 140.3 kDa

**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]