

Product datasheet for RC222495L1V

OriGene Technologies, Inc.

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MGAT2 (NM_002408) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: MGAT2 (NM_002408) Human Tagged ORF Clone Lentiviral Particle

Symbol: MGAT2

Synonyms: CDG2A; CDGS2; GLCNACTII; GNT-II; GNT2

Mammalian Cell

Selection:

None

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

 Tag:
 Myc-DDK

 ACCN:
 NM_002408

 ORF Size:
 1341 bp

ORF Nucleotide

Sequence:

The ORF insert of this clone is exactly the same as(RC222495).

OTI Disclaimer: 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.

verification at a reduced cost. Please contact our customer care team at

<u>custsupport@origene.com</u> or by calling 301.340.3188 option 3 for pricing and delivery.

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

Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence

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.

RefSeq: <u>NM 002408.3</u>

RefSeq Size: 2728 bp RefSeq ORF: 1344 bp





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Locus ID: 4247

UniProt ID: Q10469
Cytogenetics: 14q21.3
Domains: MGAT2

Protein Families: Transmembrane

Protein Pathways: Metabolic pathways, N-Glycan biosynthesis

MW: 51.4 kDa

Gene Summary: The product of this gene is a Golgi enzyme catalyzing an essential step in the conversion of

oligomannose to complex N-glycans. The enzyme has the typical glycosyltransferase

domains: a short N-terminal cytoplasmic domain, a hydrophobic non-cleavable signal-anchor domain, and a C-terminal catalytic domain. Mutations in this gene may lead to carbohydrate-

deficient glycoprotein syndrome, type II. The coding region of this gene is intronless.

Transcript variants with a spliced 5' UTR may exist, but their biological validity has not been

determined. [provided by RefSeq, Jul 2008]