

Product datasheet for RC205962L3V

OriGene Technologies, Inc.

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GFM2 (NM_032380) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: GFM2 (NM_032380) Human Tagged ORF Clone Lentiviral Particle

Symbol: GFM2

Synonyms: EF-G2mt; EFG2; hEFG2; mEF-G 2; MRRF2; MST027; MSTP027; RRF; RRF2; RRF2mt

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK
ACCN: NM 032380

ORF Size: 849 bp

ORF Nucleotide

Sequence:

OTI Disclaimer:

The ORF insert of this clone is exactly the same as(RC205962).

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 032380.3

 RefSeq Size:
 3264 bp

 RefSeq ORF:
 2340 bp

 Locus ID:
 84340

 UniProt ID:
 Q969S9

 Cytogenetics:
 5q13.3

Domains: EFG_C, GTP_EFTU_D2, EFG_IV

MW: 86.2 kDa







Gene Summary:

Eukaryotes contain two protein translational systems, one in the cytoplasm and one in the mitochondria. Mitochondrial translation is crucial for maintaining mitochondrial function and mutations in this system lead to a breakdown in the respiratory chain-oxidative phosphorylation system and to impaired maintenance of mitochondrial DNA. This gene encodes one of the mitochondrial translation elongation factors, which is a GTPase that plays a role at the termination of mitochondrial translation by mediating the disassembly of ribosomes from messenger RNA . Its role in the regulation of normal mitochondrial function and in disease states attributed to mitochondrial dysfunction is not known. Alternative splicing results in multiple transcript variants encoding distinct isoforms. [provided by RefSeq, Jul 2013]