

## **Product datasheet for SC200876**

## GFM2 (NM\_170681) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: GFM2

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

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

**ACCN:** NM\_170681

Insert Size: 450 bp

Insert Sequence: >SC200876 3'UTR clone of NM\_170681

The sequence shown below is from the reference sequence of NM\_170681. The complete sequence of

this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GATAATCTTGTTTGACCTTTTGTTAGTAACTTTGTT

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms

(SNPs).



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Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

Note: Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um

filter is required.

**RefSeq:** <u>NM\_170681.3</u>

**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]

Locus ID: 84340

**MW:** 17.4