

Product datasheet for MC212683

Eif4e1b (NM 001039683) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Eif4e1b (NM_001039683) Mouse Untagged Clone

Tag: Tag Free
Symbol: Eif4e1b

Synonyms: Eif4eloo; Gm273

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Cell Selection: Neomycin

Fully Sequenced ORF: >MC212683 representing NM_001039683

Red=Cloning site Blue=ORF Orange=Stop codon

 ${\tt TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCCC}$

GCCGCGATCGCC

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001039683

Insert Size: 552 bp

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).



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Components: 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).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

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.

RefSeq: <u>NM 001039683.2</u>, <u>NP 001034772.1</u>

RefSeq Size: 1881 bp
RefSeq ORF: 552 bp
Locus ID: 218268
UniProt ID: Q3UTA9
Cytogenetics: 13 B1

Gene Summary: Recognizes and binds the 7-methylguanosine-containing mRNA cap during an early step in

the initiation of protein synthesis and facilitates ribosome binding by inducing the unwinding

of the mRNAs secondary structures.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) uses an alternate splice site in the 3' coding region, which results in a frameshift and early stop codon, compared to variant 1. The encoded isoform (2)

has a shorter and distinct C-terminus, compared to isoform 1.