

Product datasheet for SC331335

PMF1 (NM 001199653) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: PMF1 (NM_001199653) Human Untagged Clone

Tag: Tag Free
Symbol: PMF1

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC331335 representing NM_001199653.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

 $\tt CCCAGCAGCAGGCCTGGCAGGCTCTACACAGAGAACAGAGGGAGCTGGTTGCTGCTGA$

Restriction Sites: Sgfl-Mlul

ACCN: NM_001199653

Insert Size: 543 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).

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.



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com

PMF1 (NM_001199653) Human Untagged Clone - SC331335

RefSeq: NM 001199653.1

RefSeq Size: 1061 bp
RefSeq ORF: 543 bp
Locus ID: 11243
UniProt ID: Q6P1K2
Cytogenetics: 1q22

Protein Families: Stem cell - Pluripotency

MW: 20.2 kDa

Gene Summary: Part of the MIS12 complex which is required for normal chromosome alignment and

segregation and kinetochore formation during mitosis. May act as a cotranscription partner

of NFE2L2 involved in regulation of polyamine-induced transcription of SSAT.

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) lacks an alternate exon but includes another in-frame exon in the 5' coding region, and uses an alternate splice site that causes a frameshift in the 3' coding region, compared to variant 1. The resulting isoform (3) has a distinct C-terminus and

is shorter than isoform 1.