

## **Product datasheet for SC333869**

## TIMM23B (NM\_001290118) Human Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

Tag: Tag Free

Symbol: TIMM23B

Mammalian Cell Neomycin

Selection:

Vector: pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC333869 representing NM\_001290118.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

GGCTCCTGA

**ACGCGTACGCGGCCGCTC**GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

**Restriction Sites:** Sgfl-Mlul

ACCN: NM\_001290118

Insert Size: 423 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).



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**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.

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

filter is required.

**RefSeq:** <u>NM\_001290118.1</u>

RefSeq Size: 2389 bp

RefSeq ORF: 423 bp

**Locus ID:** 100652748

UniProt ID: <u>B4DDK6</u>

Cytogenetics: 10q11.23

**MW:** 14.6 kDa

Gene Summary: May participate in the translocation of transit peptide-containing proteins across the

mitochondrial inner membrane. the PAM complex (By similarity).[UniProtKB/Swiss-Prot

Function ]

Transcript Variant: This variant (2) lacks two exons in the coding region, compared to variant 1,

which results in a shorter protein (isoform 2), compared to isoform 1.