

Product datasheet for SC330453

TMEM126B (NM 001256547) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: TMEM126B (NM_001256547) Human Untagged Clone

Tag: Tag Free
Symbol: TMEM126B

Synonyms: HT007; MC1DN29

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC330453 representing NM_001256547.

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

TAA

Restriction Sites: Sgfl-Mlul

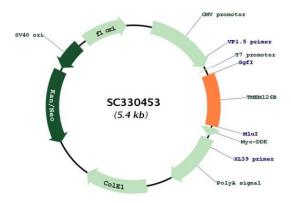
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Plasmid Map:



ACCN: NM_001256547

Insert Size: 555 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.

RefSeq: <u>NM 001256547.1</u>

RefSeq Size: 916 bp
RefSeq ORF: 555 bp
Locus ID: 55863
Cytogenetics: 11q14.1

Protein Families: Transmembrane

MW: 21.1 kDa

Gene Summary: This gene encodes a mitochondrial transmembrane protein which is a component of the

mitochondrial complex I assembly complex. The encoded protein serves as an assembly factor that is required for formation of the membrane arm of the complex. It interacts with NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 13. Naturally occurring mutations in this gene are associated with isolated complex I deficiency. A

pseudogene of this gene has been defined on chromosome 9. [provided by RefSeq, Apr 2017] Transcript Variant: This variant (4) lacks an alternate exon in the 5' coding region and initiates translation at an alternate start codon, which results in a frameshift, compared to variant 1. The encoded isoform (d) has a shorter and distinct N-terminus, compared to isoform a.