

Product datasheet for **SC332603**

TMEM218 (NM_001258245) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TMEM218 (NM_001258245) Human Untagged Clone
Tag:	Tag Free
Symbol:	TMEM218
Vector:	pCMV6-Entry (PS100001)
Fully Sequenced ORF:	>SC332603 representing NM_001258245. Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGGCTGGCACTGTGCTCGGAGTCGGTGCGGGCGTGTTCATCTTAGCCCTGCTCTGGGTGGCAGTGCTG
CTGCTGTGTGTGCTGCTGTCCAGAGCCTCCGGGGCGGCGAGGTTCTGTGCATTTTTTTATTCTTCGGT
GCTGTGATCATCACATCAGTTCTGTTGCTTTTCCCGCGAGCTGGTGAATCCCAGCCCCAGAAGTGGAA
GTTAAGATTGTGGATGACTTTTTTCATTGGCCGCTATGCCTGCTGGCTTTCCTTAGTGCCATCTTCCTT
GGAGGCCTCTTCTGGTTTTAATCCATTATGTTCTGGAGCCGATCTATGCCAAACCACTGCACCTCTAC
TGA
```

Restriction Sites:	Sgfl-Mlul
ACCN:	NM_001258245
Insert Size:	348 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:	<ol style="list-style-type: none">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:	NM_001258245.1



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RefSeq Size: 4068 bp

RefSeq ORF: 348 bp

Locus ID: 219854

UniProt ID: [A2RU14](#)

Cytogenetics: 11q24.2

Protein Families: Transmembrane

MW: 12.5 kDa

Gene Summary: May be involved in ciliary biogenesis or function.[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (10) has multiple differences, compared to variant 1. These differences result in a distinct 5' UTR and cause translation initiation at a downstream start codon, compared to variant 1. The encoded protein (isoform 2) is shorter than isoform 1.
Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.