

Product datasheet for **SC327610**

TMC7 (NM_001160364) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TMC7 (NM_001160364) Human Untagged Clone
Tag:	Tag Free
Symbol:	TMC7
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >SC327610 representing NM_001160364.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGGGAATTCGTGCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCACGATCGCC
ATGAAGTATCTCTCCGAATGGGACCACTGGAAGCGGTATAGCAGCAAGTCTTGAAGAGGTTCTAGAG
AAGGCTCGAGAGATGACGACCCACCTGGAGCTGTGGCGGGAGGACATCCGCAGCATAGAAGGAAATTT
GGCACTGGGATTCAAGTCCATTTCTCTTCTTGAGATTCTGGTGTGCTGAATTTGGTGATTTTCTG
ATCATCTTTATGCTGGTTTTGCTCCAGTCTTACTCACGAAATACAAGATCACCAACAGCAGCTTCGTG
CTCATTCTTTCAAAGACATGGATAAAACATGTACAGTCTATCCAGTAAGCAGTTCTGGACTCATTTAC
TTTACAGTTATATCATAGACTTGCTTTCTGGCACTGGTTTCTGGAGGAACTAGCCTCTTTACGGA
CATTACACCATTTGATGGGGTGAATTTTCAAGTTCACCTATGATCTGCCCTGGCGTATTTGTTAAGC
ACAATCGCTCCCTGGCCTGAGCCTTCTTTGGATAGTAAAAGGTGGTGAAGGATTCAAAATCAAC
CTGATTCGGAGTGAGGAGCACTTTCAGAGTTACTGCAACAAGATATTTGCCGGCTGGGACTTCTGCATC
ACTAACCCGAGCATGGCGGATCTGAAGCACAGCAGCTTGGGTACGAGCTCCGAGCAGATCTGGAGGAA
GAAAGAATGCGGCAGAAAATAGCAGAAAAGGACCTCAGAAGAAACAATACGCATTTACTCTTTGAGACTG
TTTTTGAAGTGTATTGTTCTGGCTGTTTTAGGGGCATGCTTTTATGCAATATACGTAGCAACTGTCTTC
TCGCAAGAGCACATGAAAAAGGAAATCGACAAGATGGTTTTTGGAGAGAACCTCTTCATATTGTATCTA
CCGTCTATTGTGATCACGCTGGCCAATTTATCACCCCAATGATCTTTGCCAAGATCATCCGCTATGAG
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TGTGTCCTGGTGTTCACGCTGGGCTCCAAGATCACATCCTGTGATGATGACACATGTGACCTTTGCGGC
TACAACCAGAACTCTACCCGTGCTGGGAGACCAAGTTGGGCAGGAAATGTACAAGCTGATGATCTTC
GACTTCATCATCATCTTGGCTGTGACACTTTCGTGGATTTTCTAGAAAGCTCCTGGTGACCTACTGT
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TTCAGAGCATCAATTCTAATTTCTTCTCCTGTTGGTGTGTTGATCGGGCTGTGTTTGCAATAATA
CCTCTGACAATCAGCATATCACGCATCCCTTCTCGAAAGCCTGTGGGCCGTTACCAACTTCAACACC
ACCTGGGAGGTATCCCCAAGACGGTGAGCACCTTCCCCAGCTCGCTGCAGTCTTCATCCATGGTGTC
ACATCCGAAGCCTTTCAGTTCCTTTCTCATGATTATTTGCCTCATCATGTTTTACTTCATTGCCTTA
GCTGGAGCACACAAACGGGTGGTCATCCAGCTCCGAGAGCAGCTATCCCTGGAAAGTCGTGACAAGTGC
TACCTAATCCAGAACTAACAGAAGCCCAAGGGACATGAGGAAC
ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
  
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Restriction Sites: SgfI-MluI

ACCN: NM_001160364

Insert Size: 1842 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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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: [NM_001160364.2](#)

RefSeq Size: 4499 bp

RefSeq ORF: 1842 bp

Locus ID: 79905

UniProt ID: [Q7Z402](#)

Cytogenetics: 16p12.3

Protein Families: Transmembrane

MW: 71.1 kDa

Gene Summary: Probable ion channel.[UniProtKB/Swiss-Prot Function]
 Transcript Variant: This variant (2) differs in both UTR's and the coding region but maintains the reading frame, compared to variant 3. This results in a protein that is shorter than isoform c. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.