

Product datasheet for SC209538

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TPM1 (NM_001018020) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: TPM1 (NM 001018020) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: TPM1

Synonyms: C15orf13; CMD1Y; CMH3; HEL-S-265; HTM-alpha; LVNC9; TMSA

ACCN: NM_001018020

Insert Size: 773 bp

Insert Sequence: >SC209538 3'UTR clone of NM_001018020

The sequence shown below is from the reference sequence of NM_001018020. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGACATTTTCTAAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).





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Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 001018020.2</u>

Summary: This gene is a member of the tropomyosin family of highly conserved, widely distributed

actin-binding proteins involved in the contractile system of striated and smooth muscles and the cytoskeleton of non-muscle cells. Tropomyosin is composed of two alpha-helical chains

arranged as a coiled-coil. It is polymerized end to end along the two grooves of actin filaments and provides stability to the filaments. The encoded protein is one type of alpha helical chain that forms the predominant tropomyosin of striated muscle, where it also functions in association with the troponin complex to regulate the calcium-dependent

interaction of actin and myosin during muscle contraction. In smooth muscle and non-muscle

cells, alternatively spliced transcript variants encoding a range of isoforms have been described. Mutations in this gene are associated with type 3 familial hypertrophic

cardiomyopathy. [provided by RefSeq, Jul 2008]

Locus ID: 7168

MW: 29.6