

Product datasheet for SC206146

MYOM1 (NM_003803) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: MYOM1

Synonyms: SKELEMIN

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

ACCN: NM_003803

Insert Size: 485 bp

Insert Sequence: >SC206146 3'UTR clone of NM_003803

The sequence shown below is from the reference sequence of NM_003803. The complete sequence of

this clone may contain minor differences, such as $\ensuremath{\mathsf{SNPs}}\xspace.$

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCCCTGAAAGGTGGCAAGAAGGCCAAGTGACCGGAGGTGCCAGGAGAGACCAGCCGGCCTGTGTGACTTG
GGTGTGAATGGTTTGGGTTAAGGATGAGACGTCTTCATGCTTTCTCCTCCCTATTATTTTCTGGCTTGA
GGGGAAAATAATGTCAGGTCTTTCACTCATATAAAAAAGCACCAACTAATGACACTTTAATTGTTTTTC
TTTATCTACAAAATTATGTGTTAAGAAAATACCATTCATAGCATGAAGATTAGGAAACAGTTTTAAGGA
GAAGACTTGAATGAAGGTTGGAGGGACATTGAATGATGGTCAGAGGGCAGACGAATGTGCGTGGGGCGA
ATTGGGATTTGCTGCAGCTGTGAAGCCATGGCCGTGTCTCGTGTGTTTTCACGAGAGGTGATGTCTTTT
CGACGGGCGCCCTCGTGGCCTTGGAACCTCCTCTGTATGAATAAACAGTTTTCACGTCTGTCCTCTTCCCC

GΑ

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

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.

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

filter is required.

RefSeq: <u>NM_003803.4</u>

Summary: The giant protein titin, together with its associated proteins, interconnects the major structure

of sarcomeres, the M bands and Z discs. The C-terminal end of the titin string extends into the M line, where it binds tightly to M-band constituents of apparent molecular masses of 190 kD (myomesin 1) and 165 kD (myomesin 2). This protein, myomesin 1, like myomesin 2, titin, and other myofibrillar proteins contains structural modules with strong homology to either

fibronectin type III (motif I) or immunoglobulin C2 (motif II) domains. Myomesin 1 and

myomesin 2 each have a unique N-terminal region followed by 12 modules of motif I or motif II, in the arrangement II-II-I-I-I-II-II-II-III-II. The two proteins share 50% sequence identity in this repeat-containing region. The head structure formed by these 2 proteins on one end of the titin string extends into the center of the M band. The integrating structure of the sarcomere arises from muscle-specific members of the superfamily of immunoglobulin-like proteins. Alternatively spliced transcript variants encoding different isoforms have been identified.

[provided by RefSeq, Jul 2008]

Locus ID: 8736

MW: 17.9