

## **Product datasheet for SC206544**

## Myotilin (MYOT) (NM\_006790) Human 3' UTR Clone

**Product data:** 

Product Type: 3' UTR Clones

Symbol: Myotilin

Synonyms: LGMD1; LGMD1A; MFM3; TTID; TTOD

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

**ACCN:** NM\_006790

Insert Size: 496 bp

Insert Sequence: >SC206544 3'UTR clone of NM\_006790

The sequence shown below is from the reference sequence of NM\_006790. The complete sequence of

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

**TATTTAAAACGGA** 

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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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## Myotilin (MYOT) (NM\_006790) Human 3' UTR Clone | SC206544

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\_006790.3</u>

Summary: This gene encodes a cystoskeletal protein which plays a significant role in the stability of thin

filaments during muscle contraction. This protein binds F-actin, crosslinks actin filaments, and

prevents latrunculin A-induced filament disassembly. Mutations in this gene have been associated with limb-girdle muscular dystrophy and myofibrillar myopathies. Several

alternatively spliced transcript variants of this gene have been described, but the full-length

nature of some of these variants has not been determined.[provided by RefSeq, Oct 2008]

**Locus ID:** 9499

**MW:** 19.5