

Product datasheet for **SC203268**

MYH7B (NM_020884) Human 3' UTR Clone

Product data:

| | |
|---------------------------|---|
| Product Type: | 3' UTR Clones |
| Product Name: | MYH7B (NM_020884) Human 3' UTR Clone |
| Symbol: | MYH7B |
| Synonyms: | MHC14; MYH14 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pMirTarget (PS100062) |
| ACCN: | NM_020884 |
| Insert Size: | 279 bp |
| Insert Sequence: | >SC203268 3'UTR clone of NM_020884 The sequence shown below is from the reference sequence of NM_020884. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site |
| Restriction Sites: | Sgfl-MluI |
| 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). |
| 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_020884.7</u> |

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GACGCCCTGGGCCCAAGCACAAAGGAGTGACGGCCTGACCCCTGGGCTCTAAAGAGGAATGTCTGCTG
TTGCACATCTGGCTGAGGCCACCTGCCCGATCCTGCCATCTCTGCATCGCCCCCTGCTGCCTTCAGCC
TTCCCTGGGCCCTGAATAAACACCACAGCCAGTTTCTTCTCATTCTTTCTTTGGGGTTCAGGAGGAA
AAACACAGTCCTAGGGACAAAAGCCAGGTCCACAGCAGTCATTTTAAAAATAAAGTTATTTAATAGTCT
CCA
ACGCGTAAGCGGCCGCGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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Summary:

The myosin II molecule is a multi-subunit complex consisting of two heavy chains and four light chains. This gene encodes a heavy chain of myosin II, which is a member of the motor-domain superfamily. The heavy chain includes a globular motor domain, which catalyzes ATP hydrolysis and interacts with actin, and a tail domain in which heptad repeat sequences promote dimerization by interacting to form a rod-like alpha-helical coiled coil. This heavy chain subunit is a slow-twitch myosin. Alternatively spliced transcript variants have been found, but the full-length nature of these variants is not determined. [provided by RefSeq, Mar 2010]

Locus ID:

57644

MW:

10.4