

Product datasheet for SC210515

OriGene Technologies, Inc.

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smooth muscle Myosin heavy chain 11 (MYH11) (NM 002474) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: smooth muscle Myosin heavy chain 11 (MYH11) (NM_002474) Human 3' UTR Clone

Symbol: smooth muscle Myosin heavy chain 11
Synonyms: AAT4; FAA4; SMHC; SMMHC; VSCM2

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_002474

Insert Size: 886 bp

Insert Sequence: >SC210515 3'UTR clone of NM_002474

The sequence shown below is from the reference sequence of NM_002474. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GTTTACACCCTGACTGTGCTTAAAAACACTTTCACTAATAAATGGTTATAAATCACAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul





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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 002474.3</u>

Summary: The protein encoded by this gene is a smooth muscle myosin belonging to the myosin heavy

chain family. The gene product is a subunit of a hexameric protein that consists of two heavy chain subunits and two pairs of non-identical light chain subunits. It functions as a major contractile protein, converting chemical energy into mechanical energy through the hydrolysis of ATP. The gene encoding a human ortholog of rat NUDE1 is transcribed from the reverse strand of this gene, and its 3' end overlaps with that of the latter. The pericentric inversion of chromosome 16 [inv(16)(p13q22)] produces a chimeric transcript that encodes a protein consisting of the first 165 residues from the N terminus of core-binding factor beta in a fusion with the C-terminal portion of the smooth muscle myosin heavy chain. This chromosomal rearrangement is associated with acute myeloid leukemia of the M4Eo subtype. Alternative splicing generates isoforms that are differentially expressed, with ratios changing during muscle cell maturation. Alternatively spliced transcript variants encoding different isoforms

have been identified. [provided by RefSeq, Jul 2008]

Locus ID: 4629 MW: 33.4