

## **Product datasheet for SC206799**

## ITGA7 (NM 002206) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: ITGA7 (NM\_002206) Human 3' UTR Clone

Symbol: ITGA7

Mammalian Cell Neomycin

Selection:

Vector:

pMirTarget (PS100062)

**ACCN:** NM 002206

**Insert Size:** 520 bp

Insert Sequence: >SC206799 3'UTR clone of NM\_002206

The sequence shown below is from the reference sequence of NM\_002206. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

GTTTCGTCTATTTATTAAAAAATATTTGAGAACAAAA

**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).

**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.



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## ITGA7 (NM\_002206) Human 3' UTR Clone - SC206799

**RefSeq:** <u>NM 002206.3</u>

**Summary:** The protein encoded by this gene belongs to the integrin alpha chain family. Integrins are

heterodimeric integral membrane proteins composed of an alpha chain and a beta chain. They mediate a wide spectrum of cell-cell and cell-matrix interactions, and thus play a role in cell migration, morphologic development, differentiation, and metastasis. This protein functions as a receptor for the basement membrane protein laminin-1. It is mainly expressed in skeletal and cardiac muscles and may be involved in differentiation and migration

processes during myogenesis. Defects in this gene are associated with congenital myopathy. Alternatively spliced transcript variants encoding different isoforms have been noted for this

gene. [provided by RefSeq, Feb 2009]

**Locus ID:** 3679 **MW:** 19