

## **Product datasheet for SC207424**

## TAPA1 (CD81) (NM 004356) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: TAPA1 (CD81) (NM\_004356) Human 3' UTR Clone

Symbol: TAPA

**Synonyms:** CVID6; S5.7; TAPA1; TSPAN28

**Mammalian Cell** 

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_004356

**Insert Size:** 561 bp

Insert Sequence: >SC207424 3'UTR clone of NM\_004356

The sequence shown below is from the reference sequence of NM\_004356. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

**GGCATGCTA** 

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

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).



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## TAPA1 (CD81) (NM\_004356) Human 3' UTR Clone - SC207424

**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 004356.4</u>

**Summary:** The protein encoded by this gene is a member of the transmembrane 4 superfamily, also

known as the tetraspanin family. Most of these members are cell-surface proteins that are characterized by the presence of four hydrophobic domains. The proteins mediate signal transduction events that play a role in the regulation of cell development, activation, growth and motility. This encoded protein is a cell surface glycoprotein that is known to complex with

integrins. This protein appears to promote muscle cell fusion and support myotube maintenance. Also it may be involved in signal transduction. This gene is localized in the tumor-suppressor gene region and thus it is a candidate gene for malignancies. Two transcript variants encoding different isoforms have been found for this gene. [provided by

RefSeq, Jul 2014]

**Locus ID:** 975 **MW:** 19.9