

Product datasheet for SC200685

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com

OriGene Technologies, Inc.

EU: info-de@origene.com CN: techsupport@origene.cn

TCTN1 (NM_001082537) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: TCTN1 (NM 001082537) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: TCTN1

Synonyms: JBTS13; TECT1

ACCN: NM 001082537

Insert Size: 419 bp

Insert Sequence: >SC200685 3'UTR clone of NM_001082537

The sequence shown below is from the reference sequence of NM_001082537. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

 $\mathsf{A}\mathsf{G}\mathsf{A}\mathsf{A}\mathsf{G}$

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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).

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 001082537.3</u>





TCTN1 (NM_001082537) Human 3' UTR Clone - SC200685

Summary: This gene encodes a member of a family of secreted and transmembrane proteins. The

orthologous gene in mouse functions downstream of smoothened and rab23 to modulate hedgehog signal transduction. This protein is a component of the tectonic-like complex, which forms a barrier between the ciliary axoneme and the basal body. A mutation in this gene was found in a family with Joubert syndrome-13. Alternative splicing results in multiple

transcript variants. [provided by RefSeq, Feb 2016]

Locus ID: 79600

MW: 15.8