

## Product datasheet for **SC214629**

### **TJP1 (NM\_175610) Human 3' UTR Clone**

#### **Product data:**

Product Type:	3' UTR Clones
Product Name:	TJP1 (NM_175610) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	TJP1
Synonyms:	ZO-1
ACCN:	NM_175610
Insert Size:	2000 bp



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**Insert Sequence:** >SC214629 3'UTR clone of NM\_175610  
 The sequence shown below is from the reference sequence of NM\_175610. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
TGTGTTTCTGTCCTTATTGACCACTTTAACTCTTGAAATATAGGAACTTAATAATGTGAAACTGGAT
TAAACTTAATCTAAATGGAACCACTCTATCAAGTATTATACCTTTTTAGAGTTGATACTACAGTTGT
TAGTATGAGGCATTTGTTTGAACGTATAAAGATGAGTGAGCATGCCCTGAACCATGGTCGGAAAACAT
GCTACACACTGCATGTTTGTGATTGACGGGACTGTTGGTATTGGCTAGAGGTTCAAAGATATTTTGCTT
TGTGATTTTTGTAATTTTTTATCGTCACTGCTTAACTTCACATATTGATTTCCGTTAAAATACCAGCC
AGTAAATGGGGTGCATTTGAGGTCTGTTCTTCCAAAGTACACTGTTTCAAACCTTACTATGGCCCTG
GCCTAGCATACTACACATTTTATTTTATTATGCATGAAGTAATATGCACACATTTTTTAAATGCACCT
GGAATATATAACCAAGTGTGTTGATTAAACAGAAATGTACAGCAAGGAGATTTACAACCTGGGGGAGGGT
GAAGTGAAGACAATGACTTACTGTACATGAAAACACATTTTTCTTAGGGAAGGATACAAAAGCATGTGA
GACTGGTTCCATGGCCTCTTCAGATCTCTAACTTCACCATATTACCACAGACATACTAACCCAGCAGAAA
TGCCCTACCCTCATGTTCTTAATCTTAGCTCATTCTCCTTGTGTTACTAAGTTTTATGGCTTTTGTG
CATTATCTAGATACTGTATCATGACAAAGACTGAGTACGTTGGTGCATTTGGTGGTTTCAGAAATGTGTT
ATCACCCAGAAGAAAATAGTGGTGTGATTTGGGGATATTTTTTCTTTCTTTCTTTCTTTCTTTTTTTT
TTTTTGGACAAGGGCAGTGGTGGTTTTCTGTTCTTTCTGGCTATGCATTTGAAAATTTTATGATGTTTAA
AGGATGCTTGACATAAATGCGTGCATACCACTTTTGTCTTGGTTTGTAAATTAACCTTTATAAACTTT
ACCTTTTTTATACATAAACAAGACCAGTCTTAAAGGCTACCTTTGTATTCTCTCTGTACCTCTTGA
GCCTTGAACCTTTGACCTCTGCAGCAATAAAGCAGCGTTTTCTATGACACATGCAAGGTCATTTTTTTTAA
GAAAAAGGATGCACAGAGTTGTTACATTTTTAAGTGTGCATTTAAAAGATACAGTTACTCAGAATTCT
CTAGTTTGATTAATTTCTGCAAAGTATCCCTACTGTAATTTGTGATACAATGCTGTGCCCTAAAGTGT
ATTTTTTACTAATAGACAATTTATTATGGCACATCAGCACGATTTCTGTTTAGATAATACACCACTAC
ATTCTGTTAATCATTAGGTGTGACTGAATTTCTTTGCCGTTATTAATAAATCTCAAATTTCTAAATCTC
CAAAATAAAACTTTTTAAAATAAAGTGTGGCTTGGTCTGTTGCCCACTGTTTTCTAGTTTCATGCAG
CTTTATAATCCTGTTTTAAAATCCTGCACACAAATCCCTATCACCCAGCGTCACCTACCACCTCGTCGT
CTGGTGTGCATGCAGAATTTCTCCCTTGGCCAGCATGTACAGATGGGTGGGCAGTGCTCATCTGAAG
GGCTCAGACTGAAGTGGGGCAGAAGGACCTGGAGACAGAGTGGGAGAAGGCAGCAGGCCGACTTCCCCC
TGTGGGTAACACACACCCCTGCGTGGAGAAAACCCCTGCATGGGGACACACGTGCGTTTGTGTGTGT
GCGTGTAAACATTTGTATATGGTTTTATCCCCAGATAAATAGCACGGGCATTGTTAATGTCACCCAC
ATTTGGGGGAAGAAAATGGGTTTGTGAGCATAAATATCCATTGTGGACATCCTACTTAACTGC
CACCCCTGTTGACATTTGGATTATTGCTATAGTGTTCAGTAAACACCTTTGTGGACTGAATCCATCT
ACGCGT AAGCGGCCGCGGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCACCCGCCCTTCTATGAAAGG
  
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**Restriction Sites:** SgfI-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:** [NM\\_175610.4](#)

**Summary:**

This gene encodes a member of the membrane-associated guanylate kinase (MAGUK) family of proteins, and acts as a tight junction adaptor protein that also regulates adherens junctions. Tight junctions regulate the movement of ions and macromolecules between endothelial and epithelial cells. The multidomain structure of this scaffold protein, including a postsynaptic density 95/disc-large/zona occludens (PDZ) domain, a Src homology (SH3) domain, a guanylate kinase (GuK) domain and unique (U) motifs all help to co-ordinate binding of transmembrane proteins, cytosolic proteins, and F-actin, which are required for tight junction function. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Aug 2017]

**Locus ID:**

7082

**MW:**

77