

## Product datasheet for **SC215315**

### **TBL2 (NM\_012453) Human 3' UTR Clone**

#### **Product data:**

|                           |                                     |
|---------------------------|-------------------------------------|
| Product Type:             | 3' UTR Clones                       |
| Product Name:             | TBL2 (NM_012453) Human 3' UTR Clone |
| Symbol:                   | TBL2                                |
| Synonyms:                 | WBSCR13; WS-betaTRP                 |
| Mammalian Cell Selection: | Neomycin                            |
| Vector:                   | pMirTarget (PS100062)               |
| ACCN:                     | NM_012453                           |
| Insert Size:              | 2000 bp                             |



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**Insert Sequence:** >SC215315 3'UTR clone of NM\_012453  
The sequence shown below is from the reference sequence of NM\_012453. The complete sequence of this clone may contain minor differences, such as SNPs.  
**Blue**=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTGAAGAGCCTGGGTGCCCTGAAGAAAGTACTCTGGGAGGGCCCGCGCAGAGGATTGAGGAGGAGGA
TCTGGCCTCCTCATGGCACTGCTGCCATCTTCTCCAGGTGGAAGCCTTTCAGAAGGAGTCTCTGG
TTTTCTTACTGGTGGCCCTGCTTCTCCATTGAACTACTCTTGTCTACTTAGGTCTCTCTCTTCTTG
CTGGCTGTGACTCCTCCCTGACTAGTGGCCAAGGTGCTTTTCTCTCCAGGCCAGTGGGTGGAATC
TGTCACCACTGGCACTGAGGAGAATGGTAGAGAGGAGAGGAGAGAGAGAGAATGTGATTTTGGCC
TTGTGGCAGCACATCCTCACACCAAGAAGTTTGTAATGTTCCAGAACAACCTAGAGAACACCTGAG
TACTAAGCAGCAGTTTTGCAAGGATGGGAGACTGGGATAGCTTCCCATCACAGAAGTGTGTCCATCAA
AAAGACACTAAGGGATTTCTTCTGGGCTCAGTTCTATTTGTAAGATGGAGAATAATCTCTCTGTGA
ACTCCTTGCAAAGATGATATGAGGCTAAGAGAATATCAAGTCCCAGGTCTGGAAGAAAAGTAGAAAAG
AGTAGTACTATTGTCCAATGTCTGAAAGTGGTAAAAGTGGGAACCAAGTGTGCTTTGAAACCAAAATTAG
AAACACATTCTTGGGAAGGCAAGTTTTCTGGGACTTGATCATACATTTTATATGGTTGGGACTTCTC
TCTTCGGGAGATGATATCTTGTTAAGGAGACCTCTTTCAGTTCATCAAGTTTCATCAGATATTTGAGT
GCCCACTCTGTGCCCAAATAATATGAGCTGGGATTAATAACGAATAAGACATGGTTTCTGCCATCAA
AGATGGCTGGTGGGAGAGAGAGATACACCTTATTAAGTGCTTTGTGTTAGTTTATTCATAGCTATTGT
TATATATGTCAAGGCAAGAGAGAGACAGAGCAGGACAGAGGCAATGGGTATTGAAGTTGAAGGGGTACA
TTTGATAAGAGAGTCGGTATTGAGAAGTTTAAATTTGTGAATTTGAAGGCTGCGTGGGTGGCAGGTA
AGGGAATGAGGTATCTGGGAAGGCTTGAGTGTAGGCAAAATTTATTCATCCCAAGTCAAACCTCTTGATC
TCTCTCACGAAAATATGACCCCTGCTTGGCCCTGACCTCTCAATGTTGGAACAAGCCACAACCGT
GGAGTTATCCTTGATGACTCCCCATTCAATATCCCTTCTGTCCACCACCACAGCTGTCTGTTCAC
CAGAACGTTCTTTTGGTTACACTCCATGTGACATCCTGAGGCTGTTGCTTCTCTCCAGCTTTTCTGTTA
CCATCCTGGTCTGGCTACTGTACCTAGTGGGCACTATAATATGGCTTCTTAATGGATCTTCTGCT
TCCTCTGTTGCCCTATTACTTTTCTATATAGCAGAATGATCTTTTGAATGTGAACCAGATCACATTA
TTCCCTAGTTAAACCCCAAGGCTTCCCATCACACTTGAATAAAATCCAAAAGAATTTTCTAAAC
CAAAAATAGTAAAAACAATTTTTTTTTCTGACAATCTCTTGTCACCCAGGCTGGAGTACAATGGCGCA
ATCTCGGCTCACTGCAACCGCCGCTCCTAGGTTCAAGCGATTCTCTGCCTCAGCTCCAGAGTAGCT
GGGATTACAGGCATGTGCCACCACGCTGGCTAATTTTTGTATTTTAGTAGAGACAGGGTTTCACGAT
GTTGGCCAGGCTGCTCTCGAACTCCTGACCTCAAATGATCCACCACCCCTTGGCTCCCCAAGTGCT
GGGATTACAGGTGTGAGCCATCGTGCCTAGCTAAAAACACTTAATCCAAAATCTTACCATGGCCTGCA
AGGTCCTGTGTAACTGGCTTCGTCCCCCTCTAAATGCTATGAATGTCAGATGGGCACCGTAAATGT
ACGCGTAAGCGGCCGCGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
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**Restriction Sites:** Sgfl-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\\_012453.4](#)

**Summary:** This gene encodes a member of the beta-transducin protein family. Most proteins of the beta-transducin family are involved in regulatory functions. This protein is possibly involved in some intracellular signaling pathway. This gene is deleted in Williams-Beuren syndrome, a developmental disorder caused by deletion of multiple genes at 7q11.23. [provided by RefSeq, Jul 2008]

**Locus ID:** 26608

**MW:** 74.8