

## Product datasheet for **SC202584**

### **TCEAL2 (NM\_080390) Human 3' UTR Clone**

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** TCEAL2 (NM\_080390) Human 3' UTR Clone  
**Symbol:** TCEAL2  
**Synonyms:** my048; MY0876G05; WEX1  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pMirTarget (PS100062)  
**ACCN:** NM\_080390  
**Insert Size:** 228 bp  
**Insert Sequence:** >SC202584 3'UTR clone of NM\_080390  
 The sequence shown below is from the reference sequence of NM\_080390. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

```

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAACGATCGCC
AGGGACACTGAAGACATTCCTTATGTGTAGTGTCCCTGGCAGGCATTTGTCAGGCCATATGTTTTAACC
TTATGGTAATACTTTGCTTTAGTCGTTCTCCTGCTACCATGAGCGTTTTGACCCACCTGCCAGTGTTT
GCTTGCTCTATGTTTCAGTAGCAGATTTTCACACATGTGCATTGCAGAGACGTCATGATTCTGGAATA
ATAAAGCAGCTTATAATATCA
ACGCGTAAGCGGCCGCGCATCTAGATTGGAAGAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
```

**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\\_080390.4](#)


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**Summary:** This gene encodes a member of the transcription elongation factor A (SII)-like (TCEAL) gene family. Members of this family contain TFA domains and may function as nuclear phosphoproteins that modulate transcription in a promoter context-dependent manner. Multiple family members are located on the X chromosome. [provided by RefSeq, Jul 2008]

**Locus ID:** 140597

**MW:** 8.9