

Product datasheet for SC207568

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TCEAL1 (NM_001006640) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: TCEAL1 (NM_001006640) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: TCEAL1

Synonyms: p21; pp21; SIIR; WEX9

ACCN: NM_001006640

Insert Size: 587 bp

Insert Sequence: >SC207568 3'UTR clone of NM_001006640

The sequence shown below is from the reference sequence of NM_001006640. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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.





TCEAL1 (NM_001006640) Human 3' UTR Clone - SC207568

RefSeq: <u>NM 001006640.2</u>

Summary: This gene encodes a member of the transcription elongation factor A (SII)-like (TCEAL) gene

family. Members of this family may function as nuclear phosphoproteins that modulate transcription in a promoter context-dependent manner. The encoded protein is similar to transcription elongation factor A/transcription factor SII and contains a zinc finger-like motif as well as a sequence related to the transcription factor SII Pol II-binding region. It may exert its effects via protein-protein interactions with other transcriptional regulators rather than via

direct binding of DNA. Multiple family members are located on the X chromosome.

Alternative splicing results in multiple transcript variants encoding a single isoform. [provided

by RefSeq, Jul 2008]

Locus ID: 9338 **MW:** 22.7