

## **Product datasheet for SC209845**

## OriGene Technologies, Inc.

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## TEA domain family member 2 (TEAD2) (NM\_003598) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: TEA domain family member 2 (TEAD2) (NM\_003598) Human 3' UTR Clone

**Symbol:** TEA domain family member 2

Synonyms: ETF; TEAD-2; TEF-4; TEF4

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_003598

**Insert Size:** 759 bp

The sequence shown below is from the reference sequence of NM\_003598. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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).





MW:

## TEA domain family member 2 (TEAD2) (NM\_003598) Human 3' UTR Clone - SC209845

**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 003598.2</u>

**Summary:** Transcription factor which plays a key role in the Hippo signaling pathway, a pathway

involved in organ size control and tumor suppression by restricting proliferation and promoting apoptosis. The core of this pathway is composed of a kinase cascade wherein MST1/MST2, in complex with its regulatory protein SAV1, phosphorylates and activates LATS1/2 in complex with its regulatory protein MOB1, which in turn phosphorylates and inactivates YAP1 oncoprotein and WWTR1/TAZ. Acts by mediating gene expression of YAP1 and WWTR1/TAZ, thereby regulating cell proliferation, migration and epithelial mesenchymal transition (EMT) induction. Binds to the SPH and GT-IIC 'enhansons' (5'-GTGGAATGT-3'). May

be involved in the gene regulation of neural development. Binds to the M-CAT motif.

[UniProtKB/Swiss-Prot Function]

**Locus ID:** 8463

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