

Product datasheet for SC206272

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TBX1 (NM_080647) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: TBX1 (NM 080647) Human 3' UTR Clone

Symbol: TBX1

Synonyms: CAFS; CATCH22; CTHM; DGCR; DGS; DORV; TBX1C; TGA; VCF; VCFS

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_080647

Insert Size: 491 bp

Insert Sequence: >SC206272 3' UTR clone of NM_080647

The sequence shown below is from the reference sequence of NM_080647. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site

Blue=Stop Codon

CAATTGGCAGAGCTCAGAATTCAAGCGATCGC

G

ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCG

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).





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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 080647.1</u>

Summary: This gene is a member of a phylogenetically conserved family of genes that share a common

DNA-binding domain, the T-box. T-box genes encode transcription factors involved in the regulation of developmental processes. This gene product shares 98% amino acid sequence identity with the mouse ortholog. DiGeorge syndrome (DGS)/velocardiofacial syndrome (VCFS), a common congenital disorder characterized by neural-crest-related developmental defects, has been associated with deletions of chromosome 22q11.2, where this gene has been mapped. Studies using mouse models of DiGeorge syndrome suggest a major role for this gene in the molecular etiology of DGS/VCFS. Several alternatively spliced transcript variants encoding different isoforms have been described for this gene. [provided by RefSeq,

Jul 2008]

Locus ID: 6899