

Product datasheet for SC210518

TAF5 (NM_006951) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: TAF5 (NM_006951) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: TAF5

Synonyms: TAF(II)100; TAF2D; TAFII-100; TAFII100

ACCN: NM_006951

Insert Size: 872 bp

Insert Sequence: >SC210518 3'UTR clone of NM_006951

The sequence shown below is from the reference sequence of NM_006951. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TGATATCTTGTAAATAAAGACAACCAGCTTTTCCAGGTTCATAA

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).



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TAF5 (NM_006951) Human 3' UTR Clone - SC210518

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 006951.5</u>

Summary: Initiation of transcription by RNA polymerase II requires the activities of more than 70

polypeptides. The protein that coordinates these activities is transcription factor IID (TFIID), which binds to the core promoter to position the polymerase properly, serves as the scaffold

for assembly of the remainder of the transcription complex, and acts as a channel for regulatory signals. TFIID is composed of the TATA-binding protein (TBP) and a group of evolutionarily conserved proteins known as TBP-associated factors or TAFs. TAFs may participate in basal transcription, serve as coactivators, function in promoter recognition or

modify general transcription factors (GTFs) to facilitate complex assembly and transcription initiation. This gene encodes an integral subunit of TFIID associated with all transcriptionally competent forms of that complex. This subunit interacts strongly with two TFIID subunits that show similarity to histones H3 and H4, and it may participate in forming a nucleosome-like

core in the TFIID complex. Alternative splicing results in multiple transcript variants. [provided

by RefSeq, Dec 2015]

Locus ID: 6877

MW: 34.2