

## Product datasheet for **SC200407**

### ATF 4 (ATF4) (NM\_001675) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	ATF 4 (ATF4) (NM_001675) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	ATF4
Synonyms:	CREB-2; CREB2; TAXREB67; TXREB
ACCN:	NM_001675
Insert Size:	128 bp
Insert Sequence:	>SC200407 3'UTR clone of NM_001675 The sequence shown below is from the reference sequence of NM_001675. The complete sequence of this clone may contain minor differences, such as SNPs. <b>Blue</b> =Stop Codon <b>Red</b> =Cloning site  GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AAGGCAAGGGGAAGAAAAGGGTCCCC <b>TAG</b> TTGAGGATAGTCAGGAGCGTCAATGTGCTTGTACATAGA GTGCTGTAGCTGTGTGTTCCAATAAATTATTTTGTAGGGAAAGTAAAAAAAAAAAAAAAA <b>ACGCGT</b> AAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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.
RefSeq:	<u><a href="#">NM_001675.4</a></u>



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**Summary:**

This gene encodes a transcription factor that was originally identified as a widely expressed mammalian DNA binding protein that could bind a tax-responsive enhancer element in the LTR of HTLV-1. The encoded protein was also isolated and characterized as the cAMP-response element binding protein 2 (CREB-2). The protein encoded by this gene belongs to a family of DNA-binding proteins that includes the AP-1 family of transcription factors, cAMP-response element binding proteins (CREBs) and CREB-like proteins. These transcription factors share a leucine zipper region that is involved in protein-protein interactions, located C-terminal to a stretch of basic amino acids that functions as a DNA binding domain. Two alternative transcripts encoding the same protein have been described. Two pseudogenes are located on the X chromosome at q28 in a region containing a large inverted duplication. [provided by RefSeq, Sep 2011]

**Locus ID:**

468

**MW:**

4.9