

Product datasheet for **SC205352**

BTF3 (NM_001207) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	BTF3
Synonyms:	BETA-NAC; BTF3a; BTF3b; NACB
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PSI00062)
ACCN:	NM_001207
Insert Size:	459 bp
Insert Sequence:	>SC205352 3'UTR clone of NM_001207 The sequence shown below is from the reference sequence of NM_001207. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC GATGAGGCTTCCAAGAATGAGGCAAACTGAATTGAGTCAACTTCTGAAGATAAAACCTGAAGAAGTTAC TGGGAGCTGCTATTTTATATTATGACTGCTTTTAAAGAAATTTTGTATGGATCTGATAAAATCTAG ATCTCTAATATTTTAAAGCCCAAGCCCTTGGACACTGCAGCTCTTTTCAGTTTTTGCTTATACACAAT TCATTCTTTGCAGCTAATTAAGCCGAAGAAGCCTGGGAATCAAGTTTGAAACAAAGATTAATAAAGTTC TTTGCCTAGTATACAGTTTTATTTTTTATTTTCATTGACACCGATCTGTACACAGTAAAAAAATTGCT TATAGAAAGCTAATCATGGCATGTAATATGGCTGATAACCTTTGGAATTTGATTAAAGATTTAAATCA CGGTGTAGTGTAAACAAAGGTGGTATAAAGTTCTCAGGTTTGAAAA ACGCGTAAGCGGCCGCGGCATCTAGATTCTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTTTCGATTCACCGCCGCTTCTATGAAAGG </pre>
Restriction Sites:	SgfI-MluI
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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_001207.5
Summary:	This gene encodes the basic transcription factor 3. This protein forms a stable complex with RNA polymerase IIB and is required for transcriptional initiation. Alternative splicing results in multiple transcript variants encoding different isoforms. This gene has multiple pseudogenes. [provided by RefSeq, Jul 2008]
Locus ID:	689
MW:	17.9