

Product datasheet for **SC338114**

TAF1 (NM_001286074) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TAF1 (NM_001286074) Human Untagged Clone
Tag:	Tag Free
Symbol:	TAF1
Synonyms:	BA2R; CCG1; CCGS; DYT3; DYT3/TAF1; KAT4; MRXS33; N-TAF1; NSCL2; OF; P250; TAF(II)250; TAF2A; TAFII-250; TAFII250; XDP
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>NCBI ORF sequence for NM_001286074, the custom clone sequence may differ by one or more nucleotides

```

ATGGGACCCGGCTGCGATTTGCTGCTGCGGACAGCAGCTACCATCACTGCTGCCCCATCATGTGTCAGACA
CGGACAGCGACGAAGATTCCGCTGGAGGCGGCCATTTTCTTTAGCGGGTTTCCTTTTCGGAACATCAA
TGGAGCCGGGCGAGCTGGAGGGGAAAGCGTCTTGATGATGAATGTAAGAAGCACTTGGCAGGCTTGGGG
GCTTTGGGGCTGGGCAGCCTGATCACTGAACTCACGGCAAATGAAGAATTGACCGGGACTGACGGTGCCT
TGGTAAATGATGAAGGTGGGTTAGGAGTACAGAAGATGCTGTGGACTATTCAGACATCAATGAGGTGGC
AGAAGATGAAAGCCGAAGATACCAGCAGACGATGGGGAGCTTGCAGCCCCCTTGGCACTCAGATTATGAT
GAAGATGACTATGATGCTGATTGTGAAGACATTGATTGCAAGTTGATGCCTCCTCCACCTCCACCCCGG
GACCAATGAAGAAGGATAAGGACCAGGATTCTATTACTGGTGTGCTGAAAATGGAGAAGGCATCATCTT
GCCCTCCATCATTGCCCTTCTCTTTGGCCTCAGAGAAAAGTGGACTTCAGTAGTTCTCTGACTCAGAA
TCTGAGATGGGACCTCAGGAAGCAACACAGGCAGAATCTGAAGATGGAAAGCTGACCCCTCCATTGGCTG
GGATTATGCAGCATGATGCCACCAAGCTGTTGCCAAGTGTACAGAACTTTTTCCAGAATTCGACCTGG
AAAGGTGTTACGTTTTCTACGCTTTTTGGACCAGGGAAGAATGTCCCATCTGTTTGGCGGAGTGTCTGG
AGAAAGAGGAAGAAGAAGCACCGTGAGCTGATACAGGAAGAGCAGATCCAGGAGGTGGAGTGCTCAGTAG
AATCAGAAGTCAGCCAGAAGTCTTTGTGGAACACGACTACGCTCCACCACCACCTCCAGAGCAGTGTCT
CTCTGATGATGAAATCACGATGATGGCTCCTGTGGAGTCCAAATTTCCCAATCACTGGAGATATAGAT
AAAGTGACAGATACCAAACCAAGAGTGGCTGAGTGGCGTTATGGGCTGCCCGACTGTGGTATGATATGC
TGGGTGCCCTGAAGATGGCAGTGGGTTGACTATGGCTTCAAAGTGAAGAAGACAGAACATGAACCTGT
GATAAAATCTAGAATGATAGAGGAATTTAGGAAACTTGAGGAAAACAATGGCACTGATCTTCTGGCTGAT
GAAAATCTCTGATGGTGACACAGCTGCATTGGGAGGATGATATCATCTGGGATGGGAGGATGTCAAAC
ACAAAGGGACAAAACCTCAGCGTGCAAGCTGGCAGGCTGGCTTCTCTAGCATGACTAGGAATGCGAT
GGCTTACAATGTTTCCAGCAAGTTTTGCAGCCACTCTTGTGATGACAAACCTTGGTACTCCATTTTCCC
ATTGACAATGAGGATCTGGTATATGGACGCTGGGAGGACAATATCATTGGGATGCTCAGGCCATGCCCC

```



[View online >](#)

GGCTGTTGGAACCTCCTGTTTTGACACTTGATCCCAATGATGAGAACCTCATTTTGGAAATTCCTGATGA
GAAGGAAGAGGCCACCTCTAACTCCCCCTCCAAGGAGAGTAAGAAGGAATCATCTCTGAAGAAGAGTCGA
ATTCTCTTAGGGAAAACAGGAGTCATCAAGGAGGAACACAGCAGAACATGTCTCAGCCAGAAGTGAAAG
ATCCATGGAATCTCTCCAATGATGAGTATTATTATCCCAAGCAACAGGGTCTTCGAGGCACCTTTGGAGG
GAATATTATCCAGCATCAATTCCTGCTGTGGAATTACGGCAGCCCTTCTTTCCACCCACATGGGGCCC
ATCAAACCTCCGGCAGTTCATCGCCACCTCTGAAAAAGTACTCATTGGTGCACCTTCTCAGCCAGGTC
CCCAGTCAGTCCAACCTTTGCTAAAGCACATCAAAAAAAGGCCAAGATGAGAGAACAAGAGAGGCAAGC
TTCAGGTGGTGGAGAGATGTTTTTATGCGCACACCTCAGGACCTCACAGGCAAAGATGGTGATCTTATT
CTTGACAGAAATAGTGAGGAAAAATGGACCTTAATGATGCAGGTTGGCATGGCAACCAAGATAAAGAAGT
ATTATAAACGGAACCTGGAAAAGATCCTGGAGCACCAGATTGTAATATGGGAAAAGTGTTTACTGCCA
TACATCTCCTTCTGGGTTCTCTCCATCTGGCCAATTGCTGCAAGCATTGAGAACAACCTTTTTCTGT
GCTCCAATTTATCTTATAAGATGCCAGAACTGATTTCTTGATCATTCCGACAAGACAGGGTACTATA
TTCGGGAATTAGTGATATTTTTGTGGTGGCCAGCAGTGTCCCTGTTTGAAGTTCCTGGGCTAACTC
CAAAGGGCCAATACGCATATTCGAGACTTTCTACAGGTTTTATTTACCGCCTTTCTGGAAAAGTAAA
GATCGGCCACGGAGGATACGAATGGAAGATATAAAAAAGCCTTCTTCCCATTAGAAAAGCAGCATCC
GGAAAGAGGCTAAAGCTCTGCGCTGACTTCAAACGCACAGGGATGGACTCAAACCTGGTGGGTGCTTAAAGT
TGATTTTCGTTTACCAACGGAAGAAGAGATCAGAGCTATGGTGTACCAGAGCAGTGTGTGCTTATTAT
AGCATGATAGCTGCAGAGCAACGACTGAAGGATGCTGGCTATGGTGAGAAATCCTTTTTTGTCCAGAAG
AAGAAAATGAGGAAGATTTCCAGATGAAGATTGATGATGAAGTTCGCACTGCCCTTGGAAACCCACAAG
GGCCTTCATTGCTGCCATGAAGGGCAAGTGTCTGCTAGAGGTGACTGGGGTGGCAGATCCACCGGGTGT
GGTGAAGGATTCTCTATGTGAAGATCCAAACAAACACACAGCAGAAGGATGATAAAGAACCAGCAGC
CAGTGAAGAAGACAGTGCAGGAACAGATGCAGACCTTCGTGCGCTTTCCCTGAAAAATGCCAAGCACT
TCTACGTAATTTGGTGTGCCTGAGGAAGAGATTAAGAAAGTTGTCGCGCTGGGAAGTGTGATGATGGTGTG
CGCACAATGTCAACAGAACAGGCTCGTTTGGAGAGGGGCCATGAGTAAATTTGCCCGTGGATCAAGGT
TTTCTGTGGCTGAGCATCAAGAGCGTTACAAGAGGAATGTCAGCGCATCTTTGACCTACAGAACAAGGT
TCTGTCATCAACTGAAGTCTTATCAACTGACACAGACAGCAGCTCAGCTGAAGATAGTGACTTTGAAGAA
ATGGGAAAGAACATTGAGAACATGTTGCAGAACAAGAAAACAGCTCTCAGCTTTCAGTGAACGGGAGG
AACAGGAGCGGAAGGAACACAGCGAATGCTACTGGCAGCAGGCTCAGCAGCATCCGGAACAATCACAG
AGATGATGACACAGCTCCGTGACTAGCCTTAACCTTTCTGCCACTGGACGCTGTCTCAAGATTTATCGC
ACGTTTTGAGATGAAGAGGGGAAAGATGTTTCGCTGTGAGACAGTCCGAAAACAGCTGTCATTGATG
CCTATGTGCGCATACGGACTACAAAAGATGAGGAATTCATTGAAAATTTGCCCTTTTTGATGAACAACA
TCGGGAAGAGATCGGAAAAGAACGGCGGAGGATTAAGAGCAACTGAGGGCGCTTAAGAGGAACCCAGGAA
AAGGAGAAGCTTAAGGGTCTCCTGAGAAGAAGCCCAAGAAAATGAAGGAGCGTCTGACCTAAAAGTGA
AATGTGGGGCATGTGGTGCCATTGGACACATGAGGACTAACAAATTTGCCCCCTCTATTATCAAACAAA
TGCGCCACCTTCCAACCTGTTGCCATGACAGAAGAACAGGAGGAGGAGTTGGAAAAGACAGTCATTAT
AATGATAATGAAGAACTTATCAAGTTGAAGGGACCAAAATTTGCTTGGGGAAACAGCTAATTGAGAGTG
CGGATGAGGTTCCGAGAAAATCTCTGGTCTCAAGTTTCTAAACAGCAGCTTCTCCAAGAAGAAACG
GCGAGTTGGAACCACTGTTCACTGTGACTATTTGAATAGACCTCATAAGTCCATCCACCGCGCCGCACA
GACCCATGGTGACGCTGTCTGATCCTTGGAGTCTATCATCAATGACATGAGAGATCTTCAAATACAT
ACCCTTTCCACACTCCAGTCAATGCAAAGGTTGTAAAGGACTACTACAAAATCATCACTCGGCAATGGA
CCTACAAACACTCCGCGAAAACGTGCGTAAACGCCTCTACCCATCTCGGGAAGAGTTGAGAGAGCATCTG
GAGCTAATTGTGAAAAATAGTGCAACCTACAATGGGCCAAAACACTCATTGACTCAGATCTCTCAATCCA
TGCTGGATCTCTGTGATGAAAAACTCAAAGAGAAAAGAACAAATAGCTCGCTTAGAGAAAAGCTATCAA
CCCCTTGTGGATGATGATGACCAAGTGGCGTTTTCTTTTATTCTGGACAACATTGTCACCCAGAAAATG
ATGGCAGTTCCAGATTCTGGCCATTTTATCACCAGTTAATAAGAAATTTGTTCCAGATTATTACAAG
TGATTGTCATCAATGGATTTAGAGACCATACGTAAGAACATCTCCAAGCACAAGTATCAGAGTCGGGA
GAGCTTTCTGGATGATGTAACCTTATTCTGGCCAACAGTGTAAAGTATAATGGACCTGAGAGTCAGTAT
ACTAAGACTGCCAGGAGATTGTGAACGCTGTTACCAGACATTGACTGAGTATGATGAACATTTGACTC
AACTTGAGAAGGATATTTGACTGCTAAAGAAGCAGCTTTGGAGGAAGCAGAATTAGAAAAGCTGGACCC
AATGACCCAGGGCCCTACACGCCTCAGGCTAAGCCTCCTGATTTGATGATACCAACACATCCCTCAGT
ATGTCTCGAGATGCCTCTGATTTCAAGATGAGAGCAATATGTCTGTCTGGATATTTCCAGTGCCACTC
CAGAAAAGCAGGTAACACAGGAAGGTGAAGATGGAGATGGTGATCTTGACAGATGAAGAGGAAGGAAGTGT

ACAACAGCCTCAAGCCAGTGCCTGTATGAGGATTTGCTTATGTCTGAAGGAGAAGATGATGAGGAAGAT
GCTGGGAGTGATGAAGAAGGAGACAATCCTTTCTCTGCTATCCAGCTGAGTAAAAGTGGAAGTGACTCTG
ATGTGGGATCTGGTGAATAAGACCCAAACAACCCCGCATGCTTCAGGAGAACACAAGGATGGACATGGA
AAATGAAGAAAGCATGATGTCCTATGAGGGAGACGGTGGGGAGGCTTCCCATGGTTTGGAGGATAGCAAC
ATCAGTTATGGGAGCTATGAGGAGCCTGATCCCAAGTCGAACACCCAAGACACAAGCTTCAGCAGCATCG
GTGGGTATGAGGTATCAGAGGAGGAAGAAGATGAGGAGGAGGAAGAGCAGCGCTCTGGGCCGAGCGTACT
AAGCCAGGTCCACCTGTACAGGACGAGGAGGACAGTGAGGATTTCCACTCCATTGCTGGGGACAGTGAC
TTGGACTCTGATGAATGA

Restriction Sites:	Sgfl-Mlul
ACCN:	NM_001286074
OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
Reconstitution Method:	<ol style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM_001286074.1</u> , <u>NP_001273003.1</u>
RefSeq Size:	7740 bp
RefSeq ORF:	5688 bp
Locus ID:	6872
UniProt ID:	<u>P21675</u>
Cytogenetics:	Xq13.1
Protein Families:	Protein Kinase
Protein Pathways:	Basal transcription factors

Gene Summary:

Initiation of transcription by RNA polymerase II requires the activities of more than 70 polypeptides. The protein that coordinates these activities is the basal transcription factor 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 the largest subunit of TFIID. This subunit binds to core promoter sequences encompassing the transcription start site. It also binds to activators and other transcriptional regulators, and these interactions affect the rate of transcription initiation. This subunit contains two independent protein kinase domains at the N- and C-terminals, but also possesses acetyltransferase activity and can act as a ubiquitin-activating/conjugating enzyme. Mutations in this gene result in Dystonia 3, torsion, X-linked, a dystonia-parkinsonism disorder. Alternative splicing of this gene results in multiple transcript variants. This gene is part of a complex transcription unit (TAF1/DYT3), wherein some transcript variants share exons with TAF1 as well as additional downstream DYT3 exons. [provided by RefSeq, Oct 2013]

Transcript Variant: This variant (3) represents the longest transcript and encodes the longest isoform (3, also known as neuron specific isoform or N-TAF1).