

Product datasheet for **SC321430**

ZBTB8 (ZBTB8A) (NM_001040441) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ZBTB8 (ZBTB8A) (NM_001040441) Human Untagged Clone
Tag:	Tag Free
Symbol:	ZBTB8
Synonyms:	BOZF1; ZBTB8; ZNF916A
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC (PS100020)
E. coli Selection:	Ampicillin (100 ug/mL)



[View online »](#)

Fully Sequenced ORF: >OriGene sequence for NM_001040441.1
 AAAGGAACGGCCTCGAAAGGGGGTCTTCCCCAGCCGAGAACCAGGAATCTTCCCTCGGC
 CCGCGCGCGACCGGCCGGTGCGCCGCGCCCGGACAATCCTCCAAACACAGACTGT
 TTGACTTAGAATCTTCCCTTTTTACTCCATTCAAGAGCAGCATTCAAATTTCCAGAA
 TGGAGATCTCCTCATCAGTCTCACCTCTGCAGCACTGAACGAGCAGCGCAGGCAAG
 ATGTATTTGTGACTGCAGTATTCTAGTTGAAGGAAGGTCTTCAAAGCACATCGAAATG
 TATTATTCGCTAGTAGCGCTACTTTAAAATGCTTCTTTCTCAGAATCAAAGGAGACGA
 GTCAGCCAACCACAGCTACATTTTCAGGCTTTCTCCCTGACACTTTTACAGTTATCTTGG
 ACTTCGTATATTCTGGCAAAGTGTCTTACTGGTCAGAATGTCATAGAAGTGATGTCGG
 CTGCTAGCTTCCCTCAGATGACTGATGCATAAGTGATGTAAGACTTTTATTAATCTT
 CCTTAGACATTAGTGAGAAAGAAAAGATCGCTATTTTCAGTCTCTCAGATAAAGATGCCA
 ATTCTAATGGTGTAGAACGTTCCCTTTTTATAGTGGTGGCTGGCAAGAAGGAAGCAGTT
 CTCACGTTCTCACCTAAGCCCAGAGCAAGGAACAGGTATAATAAGTGGAAAATCTTGGA
 ATAAGTATAATTATCATCCAGCCTCCAGAGAATACTCAACAACCTTTGGCCAAGCATG
 AACCAAGGAAAGAGTCCATTAAGAACCAACATTTGAGATTGTCACAGCCTTCTGAAG
 TTAATCTATAAGTCAAGCAAACGAGAAGTACGAACATCTGATTCTTCCAGCCATGTTT
 CCCAGTCTGAAGAACAAGCACAGATTGATGCTGAAATGGACTCTACTCCTGTTGGCTATC
 AGTACGGTCAAGGATCTGATGTCACATCCAAAAGCTTTCCAGATGATCTGCCTCGGATGC
 GATTCAAGTGCCCGTACTGCACACATGTGGTGAAGCGGAAGGCAGACCTAAAGCGCCACC
 TTCGTTGTACACAGGAGAAAGGCCCTATCCATGTCAAGCTTGTGGAAAAAGATTTAGCA
 GGCTAGACCATCTAAGTAGCCATTTTCGAACAATTCACCAGGCATGTAAGTCACTGCA
 GAAAATGTAACGTCATGTGACAGATCTAACAGGGCAAGTGGTACAGGAGGAACCAGGC
 GCTACAGACTGTGTAATGAGTGTCTTGCAGAATTTGGCATAGACAGCCTCCCCATTGACT
 TGGAAAGCTGAACAACATCTTATGTCCCATCAGATGGAGATAAGGATTCCAGATGGCACT
 TGAGTGAAGATGAGAATAGATCCTATGTGGAGATTGTAGAAGATGGGTCTGCTGATCTGG
 TCATCCAACAGGTTGATGATAGTGAAGAAGAAGAAGAAAAGAAATTAAGCCCAACATTA
 GGTAGCTGTAATGTGAACCAACAGAGCTGGCATGTCTGCAATTTACATTGACTTCTGTGA
 TCTCTCTTTCTATGGTCGGAGTCTAGTTAACAAATTTATCACACTGACTTTAAAGAAA
 TGATCTGATATAACTTGCATGCTTTCTGAACAGGCCATTGTATCCATTCTGAACCTTGT
 GCCCTAAAATGTGTTGCTGCACTGGAAAGAAAGAACTAGCTTGAGTTGGCTTGATGGATG
 AACAAATATCCTCTTACTTTTATTACAATCCTCTTACTTTATTCCAGAGATACCTTTTTTC
 TTTAATATTGGAGGATCTACTTAGCAAACCTTTTTTCAAAGAAAATACTTTAAATAAATT
 CACCACAACCAAACTTTAAAAAAAAAAAAAAAAAAAA

Restriction Sites: Please inquire

ACCN: NM_001040441

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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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:

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: [NM_001040441.1](#), [NP_001035531.1](#)

RefSeq Size: 7333 bp

RefSeq ORF: 1326 bp

Locus ID: 653121

UniProt ID: [Q96BR9](#)

Cytogenetics: 1p35.1

Gene Summary: May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (1) represents the longest transcript and encodes the longer isoform (1). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.