

Product datasheet for **SC327332**

TNRC6B (NM_001162501) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TNRC6B (NM_001162501) Human Untagged Clone
Tag:	Tag Free
Symbol:	TNRC6B
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC327332 representing NM_001162501. Blue=Insert sequence Red=Cloning site Green=Tag(s)

```

GCTCGTTTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCCGATCGCC
ATGAGAGAGAAGGAGCAAGAAAGGGAAGAACAGTTAATGGAAGACAAGAAAAGGAAGAAAGAGGATAAA
AAGAAAAAAGAAGCCACTCAGAAGGTCACGGAACAAAAACCAAGTGCCCGAAGTGACGAAACCAAGT
TTAAGCCAACCAACGGCCGCCAGCCCAATTGGCAGCTCTCCATCGCCACCAGTCAATGGTGGCAACAAT
GCCAAAAGGTGGCAGTGCCGAACGGACAACGCCAAGCGCCGCCCGCTACATGCCTCGGGAGGTGCCG
CCGCGATTCCGTTGCCAGCAGGACCACAAAGTGTTACTAAAACGTGGGCAGCCCCCTCCACCGTCCTGC
ATGCTCCTTGGGGTGGGGCAGGGCCTCCTCCCTGCACAGCACCTGGAGCAAACCAACGACACAA
GTGACAGAGCGCTGCTGCAGAGTGAGAGTGGGACTGCGCCAGACTCAACCCCTGGAGGTGCTGCTGCT
TCAAAATTATGCAATTCCACTTGGGGCTCGGGAGCCTCCTCCAACAACGGCACCTCCCCAACCCAATT
CACATCTGGGACAAGGTGATTGTAGACGGGTCTGACATGGAAGAGTGGCCTTGATTGCCAGCAAAGAC
ACTGAATCTTCTCCGAAAACACCACCGATAACAACAGTGCCTCGAACCTGGCTCTGAGAAGAGCACT
CTGCCAGGAAGCACCCTAGTAACAAAGGAAAAGGAGCCAGTGCCAGTCTGCAAGTTCTGGGAACGAA
TGTAATCTTGGGGTCTGGAAATCTGACCCTAAGGCTAAATCTGTTCAATCTTCCAACCTCTACTACAGAG
AACACAATGGACTAGGAAATTTGGAGGAATGTGAGTGGTCAGGATAGAATTGGACCTGGCTCTGGCTTC
AGCAACTTTAACCCTAATAGCAACCCATCTGCCTGGCCAGCACTGGTCCAAGAAGGAACCTCTAGGAAA
GGGGCATTGGAAACAGATAATAGTAATTCAGTGACAGGTTAGCACAGTAGGTGACATCCAGGGAA
CAGCAGTCAAAGATGGAAATGCGGGTGTTAATTTGTTGTCTCTGGCAGAGAACAGGCTCAAATTCAT
AACACTGATGGACCAAAAAATGGAACACTAATCCTTGAACCTAAGTTCACCAAAACCCCATGGAGAAT
AAGGGAATGCCCTTTGGAATGGGCTTGGGGAACACCTCCAGGAGCACTGATGCCCTTCACAAAGCACT
GGAGATCGAAAGACTGGGAGTGTGGATCTTGGGTGCAGCTAGGGGGCTTCTGGAACCTGACACAGTC
TCTGGACAAAGCAATCTGGAAACAATGGGAACAATGAAAAGAGAGAGAGGACTCCTGGAAGGAGCT
TCTGTTTCAGAAATCACTGGGTCAAAAAATGACTCTTGGGACAACAATAACAGGTCTACGGGTGGGTCC
TGAACCTTTGGCCCCAGGACTCTAATGACAACAAATGGGGTGAAGGGAACAAAAATGACATCTGGGGTC
TCTCAGGAGAAATGGAAACAGCCGACTGGGTCTGATGAGTTGAAAATGGAGAATGGAGTGGTCCAAAC
CAACCAAAATCTAGCACTGGAGCATGGGACAATCAAAGGGCCACCCCTCCCTGAAAACCAAGGCAAT

```



[View online »](#)

CCCCAGGCTCCCTGTTGGGGAAGATCTTCCAGCTCCACAGGAAGTGAAGTTGGAGGTCAAAGCACTGGA
AGCAACCACAAAGCAGGAAGTAGTGACAGTCATAACTCTGGCCGTCGGTCGTACAGGCCACACATCCT
GATTGTCAGGCTGTCTTGACAGACTCTTTGAGCCGAAGTATTTGGACCCAGGGTGCTCTCAAACACT
GGCTGGGGCCAAACTCAAATTAAGCAGGACACAGTGTGGGACATTGAAGAGGTGCCAAGGCCTGAGGGG
AAATCTGACAAAGAACTGAGGGGTGGGAGAGCGCTGCCACACAGACCAAGAACTCAGGGGGCTGGGGA
GATGCACCCAGCCAAAGCAATCAAATGAAGTCTGGATGGGGGGAGCTCTCAGCCTCTACAGAGTGAAAA
GACCCCAAGAACACAGGAGGCTGGAATGACTACAAGAAACAACACTCTTCCAACCTGGGGAGGAGGACGA
CCTGATGAAAAGACACCTTCTCTTGGAAATGAGAATCCCAGCAAGGATCAGGGGTGGGGAGGTGGACGC
CAGCCCAATCAAGGATGGTCTTCTGGAAAGAATGGTTGGGGGGAGGAAGTCGATCAGACAAAAACAG
AATTGGGAAAGTTCTGCAAGTAAACCTGTGTCTGGTGGGTGAAGGAGGCGCAAGATGAAATCGGGAC
TGGGTAATAGTTGGCAATGCAAGCCTAGTCTCAAAGGTGGGTGGGAGGATTGCAAAAGATCCCCAGCA
TGGAAATGAGACGGGGCGACAGCCCAATCTCTGGAATAAACAAACACACAGCAGCAGCCCCACAGCAG
CCGCCGCCACCACAACCAGAGGCTTCTGGTTCGTGGGGAGGCCACCCCCACCACCTCCAGGCAACGTT
CGACCTTCCAATTCCAGCTGGAGCAGCGGGCCACAGCCTGCAACACCTAAGGATGAGGAACCCAGTGGT
TGGGAAGAGCCATCCCACAGTCAATTAGTCGAAAAATGGACATTGATGATGGCACTTCAGCATGGGGA
GACCTAACAGTTATAACTACAAGAATGTGAATCTGTGGGATAAGAATTTCCCAAGGGGGCCAGCAGCT
CGAGAACCAACCTGCCACCCCAATGACCAAGTAAATCGGCATCAGTCTGGAGCAAAAGCACACCACCT
GCTCCAGATAATGGTACTTCGCGTTGGGGTGAGCCAAATGAAAGCAGTCTGGGTGGGGCGAGATGGAT
GATACAGGAGCATCGACCACAGGCTGGGGGAACACGCCCGCAACGCTCCCAATGCCATGAAGCCTAAT
TCAAATCTATGCAAGACGGCTGGGGGGAGAGTGACGGGCCAGTCACAGGAGCTCGCCATCCCAGCTGG
GAAGAGGAGGAGGATGGAGGAGTCTGGAACACCACTGGCTCTCAGGGCAGTGCTTCTCCCAACCTCA
GCAAGCTGGGGACAAGGAGGAAAGAAACAAATGAAGTGCTCACTCAAAGGAGGAACAATGATTTCATGG
ATGAATCCTCTTGCCAAACAGTTTTCAAATATGGGATGCTGAGTCAGACTGAAGATAAATCCAAGCAGC
AAAATGGATTGTTGCTGTAGGAAGCCTTTCAGATAAAAAATTTGATGTGGACAGCGAGCAGTGAATCTC
GGGATTTTTAATGATATCATGAGGAAGGATCGATCTGGGTTCGCTCCACCTAATTTCAAAGACATGGGA
ACCACAGATAGTGGGCCTATTTTTGAGAAGCTGACTTTGCCTTTCTCCAATCAAGATGGGTGCCTTGGG
GATGAGGCTCCCTGCTCTCCCTTCTCCCCTTCTCCAGCTACAAGCTGTCTCCCTCTGTTTCCACACTA
CCCAACGTGAGCCTTGGAGCAATCGGCACAGGGCTCAACCCCCAAAACCTTCGCTGCTAGACAAGGCGGT
AGTCATGGTTTGTGTTGGAAACAGCACAGCACAATCGAGAGGTCTGCACACACCCGTGCAGCCACTAAAT
TCTTCTCCAGTCTCCGGGCGCAAGTGCTTCCCAGTTTATTTCCCCCAGGTTTCTGCCTCAATGCTC
AAGCAGTTTCCCAACAGTGGCTGAGTCCAGTCTTTTCAATGTGGGGCCCCAGTTATCTCTCAACAA
ATTGCCATGCTGAGCCAGCTTCCACAAATCCCCAGTTTCAAGTTGGCATGTCAGCTTCTCTTGACAGCAG
CAGCAACAGCAGCAGTTGTTACAGAACCAGAGAAAGATTTCTCAAGCTGTACGCCAACAGCAAGAGCAG
CAGCTGGCTCGAATGGTGAGTGCAGTGCAGCAGCAGCAGCAGCAGCAGGAGGCCAGCCAGGCATGAAG
CACTCGCCCTCTCATCTGTTGGGCCCAAGCCGATCTGGACAACATGGTACCAACGCATTGAATGTG
GGGCTCCCAGACCTTCAAACCAAGGGCCAATACCTGGATATGGTTCTGGCTTCAGCTCTGGCGGCATG
GACTATGGCATGGTTGGTGGGAAGAGGCTGGAACCGAGTCCGTTTAAACAGTGGACCTCATGATG
GAGGGGCTGCCCTCTGTAGCCACACAGGAAGCCAAATATGCACAAAGATGGCGCTATAGTGCCCTTGGT
AAAACCCGGGAGGGTCACCGTACAACCAAGTTTGATATCATCCCTGGTGACACACTGGGTGGCCATACG
GGTCTGCTGGTGATAGCTGGTTACCTGCCAAATCTCCACCAACAATAAAATCGGAAGTAAATCCAGC
AATGCCAGTTGGCCTCCAGAATTCCAACCAGGAGTGCCATGGAAAGGTATCCAAAACATTGACCTGAA
TCTGACCCCTATGTCACCCAGGAAGTGTGCTGGGGGTACAGCCACATCTCCATTGTAGATACTGAC
CACCAGTCTGCTGGGATAAACACCAGGGTCTAATTCTTCCCTCAACACCTCGTGCCTTCACCTGGT
GCCTGGCCCTACAGTGCCTCTGACAACTCCTTACCAACGTTTCATAGCACTTCAGCAAAGTTCCTGAT
TACAAATCAACATGGTCCCCAGATCCCATAGGACACAACCCCACTCATCTCTCAACAAGATGTGAAAA
AACCATATTTCTCCAGGAACACTACACCGCTGCCCGGCCACCTCCTGGTCTGACCAACCCCAACCA
TCATCTCCCTGGAGCAGCACAGCACCCCGATCAGTCAGGGGGTGGGGGACACAGGACTCACGGCTCGCC
TCGGCCTCTACCTGGAGTGATGGTGGCTCAGTTCGTCTAGTTACTGGCTGGTCTTTCACAACTCACCC
CCACAGATTGATGGGTCAACCTTGAAACGATCTGCATCAGCAGTGGCCCACTGCTGACATTCCATCTG
AATCTAACCCAGGCGACTGCCCTGATCCGATCAGCACCAACAGGAGCGGGCCGCAAGGCCCAACTGCA
CTGCACATGTGTGTGTTGGGAAACATACCATCTTGTGAGTTTGCCACTGATGATGAAGTCAGCCGC
TTTCTGGCACAGCTCAGCCCCCTACACCTGCAGCAACCCCAAGTGCGCCAGCTGCGGGGTGGCAGTCTG

CTGGAGACCGGCCAGAACCAGTCAGATCCCGTGGGACCTGCTCTGAATCTTTTGGTGGGTCCACTGGG
 CTCGGGCAGTGGAGCAGCAGCGCTGGTGGCAGCAGCGGGGCCGATCTTGCTGGCGCTTCATTGTGGGGG
 CCCCCAACTATTCTTCTAGCTTATGGGGAGTCCCAACGGTGGAAGATCCCCATAGGATGGGCAGCCCT
 GCTCCTTTACTACCTGGTGACCTTCTGGGAGGAGGTCGGATTCAATCTGA
 ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
 TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites:	Sgfl-MluI
ACCN:	NM_001162501
Insert Size:	5502 bp
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:	<u>NM_001162501.1</u>
RefSeq Size:	18297 bp
RefSeq ORF:	5502 bp
Locus ID:	23112
UniProt ID:	<u>Q9UPQ9</u>
Cytogenetics:	22q13.1
MW:	194 kDa

Gene Summary:

Plays a role in RNA-mediated gene silencing by both micro-RNAs (miRNAs) and short interfering RNAs (siRNAs). Required for miRNA-dependent translational repression and siRNA-dependent endonucleolytic cleavage of complementary mRNAs by argonaute family proteins. As scaffolding protein associates with argonaute proteins bound to partially complementary mRNAs and simultaneously can recruit CCR4-NOT and PAN deadenylase complexes.

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (1) encodes the longest 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.