

Product datasheet for **SC114889**

TRIM14 (NM_014788) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	TRIM14 (NM_014788) Human Untagged Clone
Tag:	Tag Free
Symbol:	TRIM14
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF:

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>OriGene sequence for NM_014788 edited
GAATTCGGCACGAGGCACCTCCCGGCTGGGCGGGCAGAGGGAGCACCTGCTGGTCCGA
GGATGCTCCGAGGCCCGACGCTGCTCCCGGCGGGCTGTCATCCTCCTAGGGGTCGGCCGG
CCGGCCTGGAGCCGCCCCGCCACGGCCGAGGTTTCGCTTCCCTTGGGAGCCCCGCCCT
GTGTCCCGGGGAGCCGGAGGAGTGGAGATGAATGGCGGGCGCGGACCCGGGAGCCGGA
CCCCTGGGAGGTCCGAGCTTGTCTGCGCCGCTGCCGCCGCTGCGTGTGCGCGCTTTCGCCGGTGC
TGGGCGCGCACCGTGGCCACCCTGTGGGCTGGCGCTGGAGGCAGCGGTGCACGTGCAGA
AACTCAGCCAAGAATGTTTAAAGCAGCTGGCAATCAAGAAGCAGCAGCACATTGACAACA
TAACCCAGATAGAAGATGCCACCGAGAAGCTCAAGGCTAATGCAGAGTCAAGTAAAACCT
GGCTGAAGGGGAAATTCAGTGAAGTCACTGACTACTTACTTACGACGAGGAAAGCGTGGCCA
AGAAATTCATTGATAAAAAACACGACGTTACCCTCCAGGTGTACAGGGAACAAGTGAAGT
CTTGACAGAGGCAACTTGACATCATGAATGATCTCTCCAACAGGGTCTGGAGTATCAGCC
AGGAGCCCGATCCTGTCCAGAGGCTTACAGCGCGCACGCCACGCTGGATCCTGACAC
GATGCACGCGCGCTGCGCCTGTCCGCCGATCGCCTGACGGTGCCTGCGGCCTGCTGGG
CAGCCTGGGGCCCGTGGCCGTGCTGCGGTTGACGCGCTCTGGCAAGTGTGGCTCGTGA
CTGCTTCGCCACCGCCGCCACTACTGGGAGGTTGACGTGCAGGAGGCGGGCGCCGGCTG
GTGGTGGGCGCGGCTACGCTCCCTTCGGCGCCGCGGGGCTCGGCCCGCCCGCCCT
GGGCTGCAACCGCAGTCTGGTGCCTCAAGCGCTACGACCTTGAAGTACTGGGCTTCCA
CGACGGCCAGCGCAGCCGCTGCGGCCCGCGACGACCTCGACCGGCTCGGCGTCTTCT
GGACTACGAGGCCGGCGTCTCGCCTTCTACGACGTGACGGGCGGATGAGCCACCTGCA
TACCTTCCGCGCCAGTTCAGGAGCCGCTTACCCGGCCCTGCGGCTCTGGGAGGGGGC
CATCAGCATCCCCGGCTGCCCTAGGGGCCAGGACCGGCTGACAGCCTCCAGGTACGCC
GCAGTGGCCAGTCTCGCCTAATCTACCTAGATCAGCGTGGCTGGTCCCTTACTGCCTG
CTTCTTAGGGCCCTCCTCCTGCCCCAGTTCCTCCGACCAATCACGCCTACAGTGTCTTG
AAGGTTTCTCTCCTAGGCTAGTTTCAAACAGGCCCTAAACAAGTCTGCTGCTGCCCTCT
CATCAGACCTCCGACCCTCACCCACCATCACTTACACTACTTTAATCCAGTTCCTTCA
AAGTGATACCCCCACAGGTAAGCCCTCAGCATCCTGAATACATCATCCGACGCTGGGAA
CCTTCTCCCTCGTACAGCACAGGAACCTGACACATAGTAGGCACACAGTAAACGTTTGTG
AATGAATGGGAGTCATCCAGTCTGACTCTTGTCTCTTGGAGTCCCTTGAATCTTCCG
CTTCTCCACCGATTTCCAGCGTGTCCACATCACAGCTCCCTCCAGAAGCTGCAAGAGC
TTCTTAGCAGTTCCTGGTCTGAACCCTCTCCAGTCTCATCTTCCACCCTAAAACCTAGA
GTGATCTTCTAAAACCTCACTTAACCCCTCAGCTATGAAAAGGCTTCCAGGAGTTTCCA
TGAAATAACAAAAAATAACAAGCCCTCACNTNNNNNTTCAAAAAAAAAAAAAAAAAAAAA
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAACTCGAC
    
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5' Read Nucleotide Sequence:	<p>>OriGene 5' read for NM_014788 unedited CTATCCCCCGCCCGTTGCCGCAAAGGGCGGTAGGCGTGTACGGTGGGAGGTCTATATAAG CAGAGCTCGTTTGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGCG AATTCCGGCAGGAGGCACCTCCCGGCTGGGCGGGCAGAGGGAGCACCTGCTGGTCGGAG GATGCTCCGAGGCCGACGCTGCTCCCGGGGCTGTCATCTCTAGGGGTCGGCCGGC CGGCCTGGAGCCGCCCGCCACGGCCGAGGTTTCGCTTTCCTTGGGAGCCCCGCCCTG TGTCCCGGGAGCCGGAGGAGGTGGAGATGAATGGCGGGCGGGCGACCCGGAGCCGGAC CCTGGGAGGTCCGAGCTTGTGCGAGGATGCGGCTGGCGCTGCCCGAGCATGGCGACCG CGTGGCTGAGCTCTTCTGTCGCCGCTGCCGCCGCTGCGTGTGCGCGCTTTGCCCGGTGCT GGGCGCGCACCGTGGCCACCCTGTGGGCTGGCGCTGGAGGCAGCGGTGCACGTGCAGAA ACTCAGCCAAGAATGTTTAAAGCAGCTGGCAATCAAGAAGCAGCAGCACATTGACAAAT AACCCAGATAGAAGATGCCACCGAGAAGCTCAAGGCTAATGCAGAGTCAAGTAAACCTG GCTGAAGGGGAAATTCAGTGAAGTCAAGTACTACTTACGCAAGAGGAAGCGCTGGCCAA GAAATTCATTGATNAANACACGAGCTTACCCTCCANGTACAGGGAACAAGCTGACTC TTGAGAGAGCAACTTGACATCATGAATGGATCTCTCACAGGNTCTGGAGTATCAGCCAG GAGCCCGATTCTGTCCAGAGGCTTCAGACGCGCGCACGCCACGCTGGATCCTGACACGA TGCACGCGCGCTGCGCCTA</p>
3' Read Nucleotide Sequence:	<p>>OriGene 3' read for NM_014788 unedited NNCCTTCTTAGGTGNGGGCGCTTGATTTTTTTTTTTGTTATTTNATGGNAACTCCTGGAA GCCTTTTCATAGCTGAGGGGTTAAGTGAAGTTTTAGGAAGTCACTCTAGTTTTAGGGTG GAAGATGAGGACTGGGAGAGGGTTCAGACCAGGAACTGCTAAGAAGCTCTTGCAGTTCT GGAGGGAGCTGTGATGTGGACACGCTGAAATCGGTGGGGAGGAAGCGGAAGATTCAGGG ACCTCAAGAGACAGAAGAGTCAAGACTGGATGACTCCCATTCATTCACAAACGTTTACTG TGTGCCTACTATGTGTCAGGTTCTGTGCTGTACGAGGGAGAAGGTTCCAGGCTGCGGA TGATGTATTAGGATGCTGAGGGCTTACCTGTGGGGGTATCACTTTGAAGGAACTGGATT AAAGTAGTGTAAGTGTGGTGGGGTGGGGTGGGAGGCTGTATGAGAGGGCAGCAGCAG ACTTGTTAGGGCCTGTTTAAAGTACCTAGGAGAGGAACTTCAAAGCACTGTATGC GTGATTTGGTCCGGCAAAGCTGGGGCACGGAGAGGGCCCTAAGAAGCANGCAGTAAGGGGA CCAGCCACGCTGATCTAGTAGATTATGCCAGACTGGGAGCTGCGGCGTACCTGGAGGG TGTCAACCGGTTCTTGGCCCTAGGCAACCGGGGAATGCGATGGCCCCCTTCCAAATC CGCAAGGCCGGATAAAGCGGCTCTTGGAACGTGGCCCGCAAGGGCTGCAAGTGGCTTA TTCCCGCCCCGCCACGTTTCATAATGCGAGGCACCCCGCCTTTTATTTACGAAAAACCT CCACCCGGTACGGCTTGACTCGGGTCCCAAGGAGGTTTCATCTGCCCGTCCGGTAAGA CGCCACCTTCCGGGTCCAGCCGCTTCTGGCCACCAATTCGCGG</p>
Restriction Sites:	NotI-NotI
ACCN:	NM_014788
Insert Size:	1920 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).
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:	NM_014788.2 , NP_055603.2
RefSeq Size:	4454 bp
RefSeq ORF:	1329 bp
Locus ID:	9830
UniProt ID:	Q14142
Cytogenetics:	9q22.33
Domains:	zf-B_box, SPRY, PRY
Protein Families:	Druggable Genome
Gene Summary:	<p>The protein encoded by this gene is a member of the tripartite motif (TRIM) family. The TRIM motif includes three zinc-binding domains, a RING, a B-box type 1 and a B-box type 2, and a coiled-coil region. The protein localizes to cytoplasmic bodies and its function has not been determined. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2010]</p> <p>Transcript Variant: This variant (1) represents the longer transcript. Both variants encode the same protein. 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.</p>