

Product datasheet for SC331145

TRAF1 (NM_001190945) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: TRAF1 (NM_001190945) Human Untagged Clone

Tag: Tag Free
Symbol: TRAF1

Synonyms: EBI6; MGC:10353

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC331145 representing NM_001190945.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGGCCTCCAGCTCAGGCAGCCGTCCTCGCCCGGCCCCTGATGAGAATGAGTTTCCCTTTGGGTGCCCT CCCACCGTCTGCCAGGACCCAAAGGAGCCCAGGGCTCTCTGCTGTGCAGGCTGTCTCTCTGAGAACCCG AGGAATGGCGAGGATCAGATCTGCCCCAAATGCAGAGGGGAAGACCTCCAGTCTATAAGCCCAGGAAGC CGTCTTCGAACTCAGGAGAAGGCTCACCCCGAGGTGGCTGAGGCTGGAATTGGGTGCCCCTTTGCAGGT GTCGGCTGCTCCTTCAAGGGAAGCCCACAGTCTGTGCAAGAGCATGAGGTCACCTCCCAGACCTCCCAC CTAAACCTGCTGTTGGGGTTCATGAAACAGTGGAAGGCCCGGCTGGGCTGTGGCCTGGAGTCTGGGCCC GTCGATTGCTACCGGGCACCCTGCTCCGAGAGCCAGGAGGAGCTGGCCCTGCAGCACTTCATGAAGGAG AAGCTTCTGGCTGAGCTGGAGGGGAAGCTGCGTGTTTTTGAGAACATTGTTGCTGTCCTCAACAAGGAG GTGGAGGCCTCCCACCTGGCCCTGGCCACCTCTATCCACCAGAGCCAGCTGGACCGTGAGCGCATCCTG AGCTTGGAGCAGAGGGTGGTGGAGCTTCAGCAGACCCTGGCCCAGAAAGACCAGGCCCTGGGCAAGCTG GAGCAGAGCTTGCGCCTCATGGAGGAGGCCTCCTTCGATGGCACTTTCCTGTGGAAGATCACCAATGTC ACCAGGCGGTGCCATGAGTCGGCCTGTGGCAGGACCGTCAGCCTCTTCTCCCCAGCCTTCTACACTGCC AAGTATGGCTACAAGTTGTGCCTGCGGCTGTACCTGAATGGAGATGGCACTGGAAAGAGAACCCATCTG TCGCTCTTCATCGTGATCATGAGAGGGGAGTATGATGCGCTGCCGTGGCCCTTCCGGAACAAGGTC ACCTTCATGCTGCTGGACCAGAACAACCGTGAGCACGCCATTGACGCCTTCCGGCCTGACCTAAGCTCA GCGTCCTTCCAGAGGCCCCAGAGTGAAACCAACGTGGCCAGTGGATGCCCACTCTTCTTCCCCCTCAGC AAACTGCAGTCACCCAAGCACGCCTACGTGAAGGACGACACAATGTTCCTCAAGTGCATTGTGGAGACC

AGCACTTAG

Restriction Sites: Sgfl-Mlul

ACCN: NM 001190945

Insert Size: 1251 bp



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TRAF1 (NM_001190945) Human Untagged Clone - SC331145

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:

- 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 001190945.1</u>

 RefSeq Size:
 4303 bp

 RefSeq ORF:
 1251 bp

 Locus ID:
 7185

 UniProt ID:
 Q13077

 Cytogenetics:
 9q33.2

Protein Families: Druggable Genome

Protein Pathways: Pathways in cancer, Small cell lung cancer

MW: 46.2 kDa

Gene Summary: The protein encoded by this gene is a member of the TNF receptor (TNFR) associated factor

(TRAF) protein family. TRAF proteins associate with, and mediate the signal transduction from various receptors of the TNFR superfamily. This protein and TRAF2 form a heterodimeric complex, which is required for TNF-alpha-mediated activation of MAPK8/JNK and NF-kappaB. The protein complex formed by this protein and TRAF2 also interacts with inhibitor-of-apoptosis proteins (IAPs), and thus mediates the anti-apoptotic signals from TNF receptors. The expression of this protein can be induced by Epstein-Barr virus (EBV). EBV infection membrane protein 1 (LMP1) is found to interact with this and other TRAF proteins; this interaction is thought to link LMP1-mediated B lymphocyte transformation to the signal transduction from TNFR family receptors. Three transcript variants encoding two different

isoforms have been found for this gene. [provided by RefSeq, Jul 2010]

Transcript Variant: This variant (2) differs in the 5' UTR compared to isoform 1. Variants 1 and 2 both encode the same isoform (a). 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.