

Product datasheet for **MC224348**

Tnik (NM_001163009) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Tnik (NM_001163009) Mouse Untagged Clone
Tag: Tag Free
Symbol: Tnik
Synonyms: 1500031A17Rik; 4831440119Rik; AI451411; C530008O15Rik; C630040K21 Rik
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC224348 representing NM_001163009
Red=Cloning site Blue=ORF

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGC**C

ATGGCGAGCGACTCCCCAGCTCGCAGCCTGGATGAAATCGATCTCTCCGCCCTGAGGGACCCTGCAGGGA
TCTTTGAGTTGGTGAACCTGTGCGAAATGGCAGCTATGGTCAAGTTTATAAGGGTCGTCATGTCAAAC
GGCCAGCTTGTGCCATTAAGTTATGGATGTCACAGGGGATGAAGAGGAAGAAATCAAACAAGAAAT
AACATGTTGAAGAAATTTCTCATCACAGGAACATTGCTACATACTACGGTGCTTTTATCAAAAAGAAC
CTCCTGGCATGGATGACCAACTCTGGTTGGTTATGGAGTTCTGTGGTGGCTCTGTCACTGACCTGAT
CAAGAACACGAAAGGCAACACATTGAAAGAGGAGTGGATTGCATACATCTGCAGGGAGATCTTACGGGGC
CTGAGTCACCTGCACCAGCACAAAGTGATTCATCGAGATATCAAAGGGCAGAACGCTTGTGGTACTGAAA
ATGCAGAGGTTAAGCTAGTGGATTTGGAGTGAGTGCCAGCTTGACCGAACTGTGGCAGGAGGAACAC
GTTTCATCGGGACTCCCTACTGGATGGCACCAGAAGTCATTGCCTGTGATGAGAACCCGGATGCCACAT
GTGACATGCATCCCATGAGAGCCCTCTCCTCATCCACGGAACCTGCACCTCGGCTCAAGTCTAAGAA
GTGGTCAAAAAAATCCAGTCATTTATCGAGAGCTGCTTGGTAAAGAATCACAGCCAGCGGCCAGCCAGC
GAGCAGTTGATGAAGCACCCATTATACAGAGCAACCTAATGAGAGGCAGGTCCGCATCCAGCTGAAGG
ACCACATTGATCGAACAAAGAAGAAGCGAGGAGAAAAAGATGAGACTGAGTATGAATACAGCGGAAGTGA
GGAAGAAGAGGAAGAGAATGACTCTGGGAACCCAGCTCCATTCTGAACCTACCAGGGGAGTCAACACTG
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AGCAGCAGCAGCGGGAGAATGAAGAACAAGCGGAGCTACTGGCTGAGCGCCAGAAGCGCATCGAAGA
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TGAGCAAGCTCTACTTCTGGAATATAAGCGCAAACAATTGGAAGAACAGAGACAAGCAGAAAGACTGCAG
AGGCAGCTAAAGCAAGAGCGGGACTATCTGTTTCCCTCCAGCATCAGCGGCAGGAGCAGAGGCCCTGG



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AGAAGAAGCCACTGTACCATTACAAGGAGGGCATGAGTCCTAGTGAGAAGCCGGCCTGGGCCAAGGAGGT
 AGAAGAACGCTCAAGACTCAACCGACAGAGTTCACCTGCCATGCCTCACAAGGTTGCCAACAGGATCTCG
 GACCCCAACCTGCCCCCAAGATCAGAGTCTTCAGCATTAGTGGGGTTCAGCCTGCAAGGACACCCCAA
 TGCTCAGACCTGTTGACCCCAAGATCCCGCAGCTGGTAGCTGTCAAATCCCAGGGACCTGCCTTGACGGC
 CTCCCAGTCAGTACATGAGCAACCCACAAGGGCCTGTCTGGGTCCAGGAGGCTCTGAATGTGACCTCT
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 AGAAGTTTGACAGAAGCTCTTGTTACGACAGGAAGAAGACATTCCACCAAGGTGCCTCAAAGAACAAC
 TTCTATATCCCCAGCACTAGCCAGAAAGAATTCCCTGGCAATGGCAGTGCTCTGGGCCCCAGACTTGGA
 TCTCAGCCATCAGAGCAAGCAACCCTGATCTGCGCAGGACAGAGCCAGTCTGGAGAGTTCCTGCAGC
 GGACAAGCAGTGGCAGTTCCTCCAGCTCCAGCACTCCCAGCTCCCAGCCAGCTCCCAAGGAGGCTCTCA
 ACCTGGCTCCCAAGCAGGATCTAGTGAGCGGTCCAGAGTGCAGGCAACAGTAAGTCCGAAGGATCACCC
 GTGCTCCCCATGAGCCTTCCAAGGTGAAACCAGAAGAATCCAGAGACATCACACGGCCAGTCGGCCAG
 CTGATCTGACGGCATTAGCCAAAGAATTACGAGAACTCCGATTGAAGAAACAACCGCCCTGAAGAA
 AGTGACTGATTACTTCTCTCCAGCGAGGAGTCGGAGAGCAGTGAGGAAGAAGAGGAAGATGGAGAGAGT
 GAGACACATGACGGGACGGTGGCTGTGAGTACATACCCAGACTAATACCCACCGGAGCTCCAGGGAACA
 ATGAGCAGTACAACATGGGGATGGTCGGGACACATGGGCTGGAACCTTCGCATGGGACACCTTTGGCGG
 CAGCATTTCAAGAGAAGGAACCTTGATGATCAGAGAGACGGCTGAAGAGAAGAAGCGATCTGGCCACAGT
 GACAGTAATGGATTCCGCGGTACATCAATCTCCAGACCTTGTACAGCAGAGCCATTGCCAGCTGGAA
 CTCCCAGTGAAGGGCTGGGCCGGTCTCCACTCATTCCCAGGAGATGGACTCTGGGGCTGAATATGGTAT
 AGGGAGCAGCACCAGCCTCTTTCACCCCTTCGTGGACCCTCGAGTGTACCAGACATCGCCCACTGAT
 GAAGATGAAGAGGATGATGAGTCTCAGCTGCTGCCCTGTTTACTAGCGAACTTCTTAGGCAAGAACAGG
 CCAAATCAATGAAGCGAGGAAGATTCAGTGGTAAATGTGAACCAACAACATTCGCCCTCATAGTGA
 CACACCGGAAATCAGAAAATACAAGAAACGCTTCAATTCAGAAATACTTTGTGCAGCTCTATGGGGTGTG
 AACCTTCTGGTGGGACTGAAAATGGCCTGATGCTTTTGGACAGAAGTGGCCAAGGCAAAGTCTACAACC
 TAATCAACCGGAGGCGGTTTTCAGCAGATGGATGTGCTAGAAGGACTAAATGTTCTGTACAGATATCAGG
 AAAGAAGAACAAGCTCCGTGTGACTATCTCTCATGGTTAAGAAACAGAATCCTGCACAATGACCCAGAA
 GTGGAAAAGAAGCAGGGCTGGATCACTGTCCGTGACTTGGAAAGGCTGCATCCATTACAAAGTCGTTAAAT
 ATGAAAGAATCAAGTTCCTGGTATTGCCTTAAAGAATGCAGTAGAGATATATGCGTGGGCCCTAAACC
 TTACCATAAGTTCATGGCATTAAAGTCTTTTGCAGATCTTCAGCATAAGCCTCTGCTCGTTGACCTCACA
 GTAGAAGAAGTCAAAGGTTAAAGTTCATTTGGCTCACACACTGGTTCCATGTAATTGATGTTGATT
 CTGGAACTCCTACGATATCTATATACCATCCCATATTCAGGGCAATATCACTCCTCATGCTATCGTCAT
 CTTGCCTAAAACAGATGGAATGGAGATGCTTGTCTGCTATGAGGATGAAGGGGTGACGTGAACACCTAC
 GGCCGGATCACTAAGGATGTGGTCTCCATGGGAGAAATGCCACATCTGTGGCTACATTCATTCCA
 ATCAGATAATGGGCTGGGGCGAGAAAGCTATTGAGATCCGGTCAGTGGAACAGGACATTTGGATGGAGT
 GTTTATGCATAAACGAGCTCAAAGGTTAAAGTTTCTATGTGAAAGAAATGATAAGGTAATCCGTTTCAA
 CTTTGCTTTAGTGTTCATTTTAAAGAGACCACCGGTACCTGTGAAACCTTCATTTTCTTTATACTGA
 CTCAAGTCACATCTGTGTTCAATGATCAATGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

Sgfl-Mlul

ACCN:

NM_001163009

Insert Size:

4092 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:	<u>NM_001163009.1, NP_001156481.1</u>
RefSeq Size:	4625 bp
RefSeq ORF:	4092 bp
Locus ID:	665113
Cytogenetics:	3 A3
Gene Summary:	<p>Serine/threonine kinase that acts as an essential activator of the Wnt signaling pathway. Recruited to promoters of Wnt target genes and required to activate their expression. May act by phosphorylating TCF4/TCF7L2. Appears to act upstream of the JUN N-terminal pathway. May play a role in the response to environmental stress. Part of a signaling complex composed of NEDD4, RAP2A and TNIK which regulates neuronal dendrite extension and arborization during development. More generally, it may play a role in cytoskeletal rearrangements and regulate cell spreading (By similarity).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (4) lacks an internal, in-frame coding exon, and differs at the 3' end compared to variant 1, resulting in a longer isoform (4) with a distinct C-terminus compared to 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.</p>