

Product datasheet for **MC229336**

Tek (NM_001290551) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Tek (NM_001290551) Mouse Untagged Clone
Tag: Tag Free
Symbol: Tek
Synonyms: AA517024; Cd202b; Hyk; STK1; Tie-2; Tie2
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC229336 representing NM_001290551
Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGCC**

ATGGACTCTTTAGCCGGCTTAGTTCTCTGTGGAGTCAGCTTGCTCCTTTATGGAGTAGTAGAAGGCGCCA
TGGACCTGATCTTGATCAATCCCTACCTCTTGCTGTGATGCCGAAACATCCCTCACCTGCATTGCCTC
TGGGTGGCACCCCATGAGCCCATCACCATAGGAAGGGACTTTGAAGCCTTAATGAACCAGCACCAAGAT
CCACTGGAGGTTACTCAAGATGTGACCAGAGAATGGGCGAAAAAGTTGTTTGGAAAGAGAGAAAAAGGCCA
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ATATCTTTCAAAAAGGTGTTAATTAAGAAGAAGATGCAGTGATTTACAAAAATGGATGTGAAGCTCAGA
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AGGACCTGTAAGAAAAGGTGTAGTGGACCAGAAGGATGCAAGTCTTATGTGTTCTGTCTCCAGACCCTT
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ACCAGACTGTAAGCTCAGGTGCCACTGTACCAATGAAGAGATATGTGATCGGTTCCAAGGATGCCTCTGC
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GTGTGAACACAGTGGCTGGGATGGTGGAAAAGCCTTTCAACATTTCCGTCAAAGTTCTTCCAGAGCCCT
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TGACGAATGAGATTTTCACTCTCAACTACTTGGAGCCCGGACTGACTACGAGCTGTGTGTGCAGCTGGC
CCGTCTGGAGAGGGTGGAGAAGGGCATCTGGGCTGTGAGACGATTTACAACAGCGTCTATCGGACTC
CCTCTCCAAGAGGTCTCAGTCTCTGCCAAAAAGCCAGACAGCTCTAAATTTGACTTGGCAACCGATAT



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TTACAAACTCAGAAGATGAATTTTATGTGGAAGTCGAGAGGCGATCCCTGCAAACAACAAGTGATCAGCA
GAACATCAAAGTGCCTGGGAACCTGACCTCGGTGCTACTGAGCAACTTAGTCCCCAGGGAGCAGTACACA
GTCCGAGCTAGAGTCAACACCAAGGCGCAGGGGAGTGGAGTGAAGAAGTCAAGGCTGGACCCTTAGTG
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GAGGACGCTTCCACATTCAGCCTCTGCAGACCTCGGAGGGGAAAGATGCTACTCATAGCCATCCTT
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GACATCAAGTTTCAAGACGTGATCGGAGAGGGCAACTTTGGCCAGGTTCTGAAGGCACGCATCAAGAAGG
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CGCAGGAGAACTGGAGGTTCTTTGTAACCTTGACACCATCCAACATCATTAACTCTTGGGAGCATGT
GAACACCGAGGCTATTTGTACCTAGCTATTGAGTATGCCCGCATGGAAACCTCCTGGACTTCTGCGTA
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AGGGACCTGGCTGCCAGAAACATTTTAGTTGGTAAAACCTACATAGCCAAAATAGCAGATTTTGGATTGT
CACGAGGTCAAGAAGTGTATGTGAAAAAGACAATGGGAAGGCTCCCAGTGCCTGGATGGCAATCGAATC
ACTGAACTATAGTGTCTATAACAACAACAGTGTGTCTGGTCTATGGTGTATTGCTCTGGGAGATTGTT
AGCTTAGGAGGCCCCCTACTGCGGCATGACGTGCGCGGAGCTCTATGAGAAGCTACCCAGGGCTACA
GGCTGGAGAAGCCCTGAACTGTGATGATGAGGTGTATGATCTAATGAGACAGTGTGGAGGGAGAAGCC
TTATGAGAGACCATCATTTGCCAGATATTGGTGCCTTAAACAGGATGCTGGAAGAACGGAAGACATAC
GTGAACACCACACTGTATGAGAAGTTTACCTATGCAGGAATTGACTGCTCTGCGGAAGAAGCAGCCTAG
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ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001290551
- Insert Size:** 3219 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:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- 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_001290551.1](#), [NP_001277480.1](#)

RefSeq Size: 4549 bp

RefSeq ORF: 3219 bp

Locus ID: 21687

Cytogenetics: 4 43.34 cM

Gene Summary: Tyrosine-protein kinase that acts as cell-surface receptor for ANGPT1, ANGPT2 and ANGPT4 and regulates angiogenesis, endothelial cell survival, proliferation, migration, adhesion and cell spreading, reorganization of the actin cytoskeleton, but also maintenance of vascular quiescence. Has anti-inflammatory effects by preventing the leakage of proinflammatory plasma proteins and leukocytes from blood vessels. Required for normal angiogenesis and heart development during embryogenesis. Required for post-natal hematopoiesis. After birth, activates or inhibits angiogenesis, depending on the context. Inhibits angiogenesis and promotes vascular stability in quiescent vessels, where endothelial cells have tight contacts. In quiescent vessels, ANGPT1 oligomers recruit TEK to cell-cell contacts, forming complexes with TEK molecules from adjoining cells, and this leads to preferential activation of phosphatidylinositol 3-kinase and the AKT1 signaling cascades. In migrating endothelial cells that lack cell-cell adhesions, ANGPT1 recruits TEK to contacts with the extracellular matrix, leading to the formation of focal adhesion complexes, activation of PTK2/FAK and of the downstream kinases MAPK1/ERK2 and MAPK3/ERK1, and ultimately to the stimulation of sprouting angiogenesis. ANGPT1 signaling triggers receptor dimerization and autophosphorylation at specific tyrosine residues that then serve as binding sites for scaffold proteins and effectors. Signaling is modulated by ANGPT2 that has lower affinity for TEK, can promote TEK autophosphorylation in the absence of ANGPT1, but inhibits ANGPT1-mediated signaling by competing for the same binding site. Signaling is also modulated by formation of heterodimers with TIE1, and by proteolytic processing that gives rise to a soluble TEK extracellular domain. The soluble extracellular domain modulates signaling by functioning as decoy receptor for angiopoietins. TEK phosphorylates DOK2, GRB7, GRB14, PIK3R1, SHC1 and TIE1.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) lacks an alternate in-frame exon in the 5' coding region, compared to variant 1. It encodes isoform 3, which lacks an internal segment and is shorter, compared to isoform 1. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.