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Product datasheet for SC323528

MAP4K1 (NM_007181) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	MAP4K1 (NM_007181) Human Untagged Clone
Tag:	Tag Free
Symbol:	MAP4K1
Synonyms:	HPK1
Mammalian Cell Selection:	None
Vector:	pCMV6-XL4
E. coli Selection:	Ampicillin (100 ug/mL)

View online »

Fully Sequenced ORF:

>NCBI ORF sequence for NM_007181, the custom clone sequence may differ by one or more nucleotides

ATGGACGTCGTGGACCCTGACATTTTCAATAGAGACCCCCGGGACCACTATGACCTGCTACAGCGGCTGG
GTGGCGGCACGTATGGGGAAGTCTTTAAGGCTCGAGACAAGGTGTCAGGGGACCTGGTGGCACTGAAGAT
GGTGAAGATGGAGCCTGATGATGATGTCTCCACCCTTCAGAAGGAAATCCTCATATTGAAAACTTGCCGG
CACGCCAACATCGTGGCCTACCATGGGAGTTATCTCTGGTTGCAGAAACTCTGGATCTGCATGGAATTCT
GTGGGGCTGGTTCTCTCCAGGACATCTACCAAGTGACAGGCTCCCTGTCAGAGCTCCAGATTAGCTATGT
CTGCCGGGAAGTGCTCCAGGGACTGGCCTATTTGCACTCACAGAAGAAGATACACAGGGACATCAAGGGA
GCTAACATCCTCATCAATGATGCTGGGGAGGTCAGATTGGCTGACTTTGGCATCTCGGCCCAGATTGGGG
CTACACTGGCCAGACGCCTCTCTTTCATTGGGACACCCTACTGGATGGCTCCGGAAGTGGCAGCTGTGGC
CCTGAAGGGAGGATACAATGAGCTGTGTGACATCTGGTCCCTGGGCATCACGGCCATCGAACTGGCCGAG
CTACAGCCACCGCTCTTTGATGTGCACCCTCTCAGAGTTCTCTTCCTCATGACCAAGAGTGGCTACCAGC
CTCCCCGACTGAAGGAAAAAGGCAAATGGTCGGCTGCCTTCCACAACTTCATCAAAGTCACTCTGACTAA
GAGTCCCAAGAAACGACCCAGCGCCACCAAGATGCTCAGTCATCAACTGGTATCCCAGCCTGGGCTGAAT
CGAGGCCTGATCCTGGATCTTCTTGACAAACTGAAGAATCCCGGGAAAGGACCCTCCATTGGGGACATTG
AGGATGAGGAGCCCGAGCTACCCCCTGCTATCCCTCGGCGGATCAGATCCACCCACCGCTCCAGCTCTCT
GGGGATCCCAGATGCAGACTGCTGTCGGCGGCACATGGAGTTCAGGAAGCTCCGAGGAATGGAGACCAGA
CCCCCAGCCAACACCGCTCGCCTACAGCCTCCTCGAGACCTCAGGAGCAGCAGCCCCAGGAAGCAACTGT
CAGAGTCGTCTGACGATGACTATGACGACGTGGACATCCCCACCCCTGCAGAGGACACACCTCCTCCACT
TCCCCCCAAGCCCAAGTTCCGTTCTCCATCAGACGAGGGTCCTGGGAGCATGGGGGATGATGGGCAGCTG
AGCCCGGGGGTGCTGGTCCGGTGTGCCAGTGGGCCCCCACCAAACAGCCCCCGTCCTGGGCCTCCCCCAT
CCACCAGCAGCCCCCACCTCACCGCCCATTCAGAACCCTCACTCTGGAACCCACCCTCCCGGGAGCTTGA
CAAGCCCCCACTTCTGCCCCCCAAGAAGGAAAAGATGAAGAGAAAGGGATGTGCCCTTCTCGTAAAGTTG
TTCAATGGCTGCCCCCTCCGGATCCACAGCACGGCCGCCTGGACACATCCCTCCACCAAGGACCAGCACC
TGCTCCTGGGGGCAGAGGAAGGCATCTTCATCCTGAACCGGAATGACCAGGAGGCCACGCTGGAAATGCT
CTTTCCTAGCCGGACTACGTGGGTGTACTCCATCAACAACGTTCTCATGTCTCTCTCAGGAAAGACCCCC
CACCTGTATTCTCATAGCATCCTTGGCCTGCTGGAACGGAAAGAGACCAGAGCAGGAAACCCCATCGCTC
ACATTAGCCCCCACCGCCTACTGGCAAGGAAGAACATGGTTTCCACCAAGATCCAGGACACCAAAGGCTG
CCGGGCGTGCTGTGTGGCGGAGGGTGCGAGCTCTGGGGGCCCGTTCCTGTGCGGTGCATTGGAGACGTCC
GTTGTCCTGCTTCAGTGGTACCAGCCCATGAACAAATTCCTGCTTGTCCGGCAGGTGCTGTTCCCACTGC
CGACGCCTCTGTCCGTGTTCGCGCTGCTGACCGGGCCAGGCTCTGAGCTGCCCGCTGTGTGCATCGGCGT
GAGCCCCGGGCGGCCGGGGAAGTCGGTGCTCTTCCACACGGTGCGCTTTGGCGCGCTCTCTTGCTGGCTG
GGCGAGATGAGCACCGAGCACAGGGGACCCGTGCAGGTGACCCAGGTAGAGGAAGATATGGTGATGGTGT
TGATGGATGGCTCTGTGAAGCTGGTGACCCCGGAGGGGTCCCCAGTCCGGGGACTTCGCACACCTGAGAT
CCCCATGACCGAAGCGGTGGAGGCCGTGGCTATGGTTGGAGGTCAGCTTCAGGCCTTCTGGAAGCATGGA
GTGCAGGTGTGGGCTCTAGGCTCGGATCAGCTGCTACAGGAGCTGAGAGACCCTACCCTCACTTTCCGTC
TGCTTGGCTCCCCCAGGCTGGAGTGCAGTGGCACGATCTCGCCTCACTGCAACCTCCTCCTCCCAGGTTC
AAGCAATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGATTACAGGCCTGTAG

5' Read Nucleotide Sequence:

>OriGene 5' read for mutant NM_007181 unedited
ACGCCGTTGAGCAATGGGCGGTAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGCTCGTTTAGTGAAC
CGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGCGAATTCGGCACGAGGCTCAGGCCCCCAGGG
ATGGACGTCGTGGACCCTGACATTTTCAATAGAGACCCCCGGGACCACTATGACCTGCTACAGCGGCTGG
GTGGCGGCACGTATGGGGAAGTCTTTAAGGCTCGAGACAAGGTGTCAGGGGACCTGGTGGCACTGATGAT
GGTGAAGATGGAGCCTGATGATGATGTCTCCACCCTTCAGAAGGAAATCCTCATATTGAAAACTTTGCCG
GCACGCCAACATCGTGGGCCTACCATGGGAGTTATCTCTGGTTGCAGAAAACTCTGGATCTGCATGGAAT
TCTGTGGGGCTGGTTCTCTCCAGACCATCTACCAGTTGACAGGCTCCCTGGTCGAACCTCCGATAACCTA
TGTCTGCCGGGAGTTGCTCAGGGACTGCCCTATTTGCCCTACCGAAGAAGAATACAAGGGCCCTCAAGGG
ACTTACCTTCCTAATCATTGATGCTGGGAAGGTCAGATTGGCTGACTTGGACTCTCGCCCAAATTGGCTC
ACCCTGCCGAACCCCTCTTTTCTTGGGCACCCTATCTGAAGGGCTCGAATGGGCCCTTGGCCCGAGAGGA
GAGAACAATAGCCTGTGTGACTGTGTCCGGGGTACAGGACATAAAGGGCACGACTACCCACGCTTGTGTG
TGCCCCTAGATCTCTCTCTTACACAATGTGTCAGCGTCGCACATGAAAAGACAAAGTGTCGCTTCCATCT
CAGTCTCGCGCAAGGTGCGGAGGACTCGCGAAGCCTAACTGTCACTCGTTAAGCGACTGATCTATGAA

Kinase Domain Sequence:

>SC323528 kinase domain raw sequence. By performing BLASTX analysis with this sequence against NCBI refernce protein database, you can confirm the presence of the kinase-deficient mutation
AKKATGMGCAATGGGCGGTAGSGTGTACGGTGGGAGGTCTATATAAGCAGAGCTCGTTTAGTGAACCGTC
AGAATTTTGTAATACGACTCACTATAGGGCGGCCGCGAATTCGGCACGAGGCTCAGGCCCCCAGGGATGG
ACGTCGTGGACCCTGACATTTTCAATAGAGACCCCCGGGACCACTATGACCTGCTACAGCGGCTGGGTGG
CGGCACGTATGGGGAAGTCTTTAAGGCTCGAGACAAGGTGTCAGG

Restriction Sites: Please inquire

ACCN: NM_007181

Insert Size: 2900 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 kinase-deficient mutant clone was generated by created by site-directed mutagenesis from the corresponding wild-type clone. See details in "Application of active and kinase-deficient kinome collection for identification of kinases regulating hedgehog signaling." Cell. 2008 May p536-548.

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_007181.3, NP_009112.1

RefSeq Size: 2732 bp

RefSeq ORF:	2502 bp
Locus ID:	11184
UniProt ID:	Q92918
Cytogenetics:	19q13.2
Domains:	pkinase, CNH, TyrKc, S_TKc
Protein Families:	Druggable Genome, Protein Kinase
Protein Pathways:	MAPK signaling pathway
Gene Summary:	Serine/threonine-protein kinase, which may play a role in the response to environmental stress. Appears to act upstream of the JUN N-terminal pathway. May play a role in hematopoietic lineage decisions and growth regulation. Able to autophosphorylate. [UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (2) represents the longer transcript and encodes the longer isoform (2).