

Product datasheet for MC224198

Mapk8ip3 (NM_001163451) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Mapk8ip3 (NM_001163451) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Mapk8ip3
Synonyms:	BB120594; D17Wsu15e; JIP-3; Jip3; JSAP1; JSAP1a; JSAP1b; JSAP1c; JSAP1d; mKIAA1066; Syd2
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
Fully Sequenced ORF:	>MC224198 representing NM_001163451 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCCCGATCGCC

ATGATGGAGATCCAGATGGACGAGGGAGGAGTGGTGGTGTACCAAGACGACTACTGCTCGGGCTCGG
TCATGTCCGAGCGTGTGTCCGGCCTGGCGGGCTCCATCTACCGCAGTTCGAGCGCCTCATTCACTGCTA
TGACGAGGAGGTGGTCAAGGAGCTCATGCCGCTGGTGGTGAACGTGCTGGAGAACCTTGACTCGGTGCTG
AGCGAGAACCAGGAGCAGGAGGTGGAGCTGAGCTCCTACGCGAGGACAACGAGCAGCTGCACGCAAT
ACGAGCGCGAGAAGGCGCTGCGCAAACAGGCCGAGGAGAAATTCATCGAATTTGAAGATGCCTTGGAACA
AGAGAAGAAAGAACTCCAGATCCAGGTAGAACATTATGAGTTTCAGACACGCCAGCTGGAGCTAAAGGCC
AAAACTATGCAGATCAGATTTCCCGACTGGAGGAACGAGAATCGGAGATGAAGAAGGAATACAATGCC
TGCACCAGCGGCACACAGAGATGATCCAGACCTATGTGGAACACATTGAAAGATCCAAGATGCAGCAAGT
TGGGGGTAGCGGCCAAACAGAAAGCAGCCTGCCCGGGCGGAGGAAGGAGCGTCCCACCTCTCTGAATGTC
TTCCCCCTGGCTGATGGCATGTGTCAAATGATGAGATGTCTGAGTCAGGCCAGTCCCTCAGCAGCTGCAA
CACCCAGTACCACAGGTACCAAGTCCAACACACCCACCTCCTCCGTGCCCTCAGCAGCAGTCAAGCCACT
CAACGAGAGCCTACAGCCCTGGGGACTATGTCAAGTGTACAAAAGAACAACAAGCAGGCCGAGAGAAG
CGCAATAGCCGTAACATGGAGGTCCAGGTACCCAAGAGATGCGGAACGTGAGTATCGGCATGGGCAGCA
GTGACGAGTGGTCCGATGTTCCAGACATTATCGACTCCACCCAGAGCTGGATGTGTGCTGAAACCCG
TCTGGAGCGCACAGGAAGCAGCCAAACCCAGGGAATTGTAACAAGCTTTTGAATCAACTGACTCC
TTGTATCACGAATCTCCACGGCGGGATCTGAGGTATCGGGGATGTGGACGAGGAGCTGATCTCCTAG
GGGAGTTTTAGTGCAGATGATTTTTTGAATGGGCAAGAAGTGGGAACTGCTGCTGGAGAACTC
ACAGCTTCTAGAGACAAAAATGCTTTAAATGTAGTGAAGAATGACCTCATTGCTAAGGTTGACCAACTG
TCAGGAGAACAGGAGTCTGAAGGTTGAGCTGGAAGCAGCAAGCAAGCAAGTCAAGCTGGAGAACC
GAATCAAAGAGCTTGAAGAAGAACTGAAGAGAGTCAAGTCAAGGAGTCAAGTCAAGGAGTCAAGTCAAGGAG
AGAAGAGGTGGAGGATGTAAGCAGCTATCTGTACAGAATTGGACAAAATCCCCATGGCCAGGCCGA
CGCTTACACGGGTGGAGATGGCCCGAGTGTCTATGGAACGCAACCAGTACAAGGAACGCCTCATGGAGC



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TGCAGGAGGCTGTGAGGTGGACTGAAATGATCAGAGCATCAAGGGAACACCCATCTGTCCAGGAGAAGAA
 GAAGTCCACCATCTGGCAGTTCTTTAGTCGCCTCTTCAGCTCCTCATCTAGCCCCCTCCGGCCAAACGA
 TCCTACCCATCTGTGAACATTCCTACAAGTCAACCACTGCAGCTGGCTTTAGCCAGCGTCGCAGCCATG
 CTTTGTGCCAGATCTCAGCCGGCAGCAGGCCCTGGAGTTCTTCCTGATGATGACTGCACCTCTTCTGC
 CCGGCGGGAGCAGAAGCGGGAGCAGTACCGCCAGGTTCTGTAACACGTGCGCAATGATGACGGGAGGCTG
 CAGGCCGTGGGTGGAGCCTGCCAAGTACAAGCAGCTGAGCCCCAATGGAGGCCAGGAAGACACCC
 GGATGAAAAATGTCCTGTCCCTGTGTACTGCCGCCCTCTGGTGGAGAAGGACCCCTCGACAAGCTGTG
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 CTGAGAAGAAGAAGGCAAGGAAACCCCTGAGGCAGATGCTACCTCCAGTCGGGTATGGATCCTCACCAG
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 GTCTGCAATGCCACGTCTGTATCTCCAGCATTCTCGGCCAGTGACAGTGACTATCCCCCTGGG
 AGATGTTCTAGACAGTGATGTGAACCTGAAGATTGAGTGTGCTGATGGTGTGCTGGCTGGCATCACCT
 GGTGGGTGTGCTACCCGCTCAATGTTCCACGTAGCAACTGTTCTCACGAGGAGACACCCAGTACTG
 GACAAGGGGCAGGGGATGTGGCGACCACTGCCAATGGGAAGGTCAACCCGTCCCAATCCACAGAAGAAG
 CCACAGAAGCCACGGAGGTGCCAGACCCCTGGTCCAGCGAGTCAAGAACGACAGTCCGGCCCGGGCC
 TCTCACAGAGCATGTCTTACTGACCCAGCACCCACCCATCCTCCAGCACCCAGCCTGCCAGTGAGAAT
 GGGTCAGAGTCCAATGGCACCATTGTACAGCCTCAGGTGGAGCCAGTGGGGAACCTCAACAACAACCA
 GTAGCGCTGCACCCTATGTGGCTAGGAGCCAGAATGGCTGGCTCTATGTGCATTACGCGGTAGCCAA
 CTGGAAGAAGTGTCTGCACTCCATCAAGCTAAAAGACTCTGTGCTGAGCCTGGTGCATGTCAAAGCCGA
 GTGCTGGTAGCTCTTGAGATGGGACCCCTGGCTATCTCCATCGTGGAGAGGATGGCCAGTGGGACCTGA
 GCAACTACCACCTAATGGACCTGGGCCACCCACCACTCCATCCGCTGCATGGCTGTTGTAATGACCC
 AGTTTGGTGTGGCTACAAGAACAAGGTGCATGTTATCCAGCCCAAGACAATGCAGATTGAGAAATCATT
 GATGCCACCCAAAGCGGGAAAGCCAGGTACGTACAGTGGCCTGGATCGGTGATGGAGTGTGGGTCTCTA
 TTCGTTGGATTCTACCCTTCGGCTCTACCATGCTCACACCCACCAAGCAGCCTGCAGGATGTGGACATTGA
 GCCCTATGTTAGCAAGATGCTAGGAACCGCAAGCTGGGCTTCTCCTTCGTGCGCATCACAGCCTTACTC
 ATTGCAGGCAACCGTCTGTGGTGGGCACTGGCAATGGGGTTGTCATCTCCATCCCCTTGACTGAGACTG
 TGGTCTGCATCGAGGCCAGCTCCTAGGCTCCGAGCCAACAAGACATCCCAACATCTGGGAGGGGAC
 CCGCCAGGGGCATCATCCATGTATGGGGACGACAGCAGTGACAAGGCCCGCAGTATTTTATCCCC
 TACTGCTCCATGGCACAGGCTCAGCTTTGCTTCCATGGGCACCGTATGCTGTCAAATTTCTTGTCTGT
 TGCCAGGAAATGTCTGGCCACTCTCAATGGCAGTGTGCTAGACAGCCCATCAGAGGGCCCTGGGCCTGC
 TGACCCGCTGCAGATGCTGAGGCCAGAAGTTGAAGAATGCACTGGTGTGAGTGGTGGTGAAGGTTAC
 ATTGACTTCCGTATCGGAGACGGAGAGGATGATGAAACTGAGGAATGTGCCGGGGACGTGAACCAGACAA
 AGCCCTCGTTGTCCAAGGCTGAGCGCAGCCACATCATCGTGTGGCAGGTGTCTACACCCCTGAGTGA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001163451
- Insert Size:** 3918 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:

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_001163451.1](#), [NP_001156923.1](#)

RefSeq Size: 5483 bp

RefSeq ORF: 3918 bp

Locus ID: 30957

UniProt ID: [Q9ESN9](#)

Cytogenetics: 17 12.53 cM

Gene Summary: The JNK-interacting protein (JIP) group of scaffold proteins selectively mediates JNK signaling by aggregating specific components of the MAPK cascade to form a functional JNK signaling module. May function as a regulator of vesicle transport, through interactions with the JNK-signaling components and motor proteins (PubMed:10523642, PubMed:10629060). Promotes neuronal axon elongation in a kinesin- and JNK-dependent manner (PubMed:23576431, PubMed:25944905, PubMed:28259553). Activates cofilin at axon tips via local activation of JNK, thereby regulating filopodial dynamics and enhancing axon elongation (PubMed:23576431, PubMed:25944905, PubMed:28259553). Its binding to kinesin heavy chains (KHC), promotes kinesin-1 motility along microtubules and is essential for axon elongation and regeneration (PubMed:23576431, PubMed:25944905, PubMed:28259553). Regulates cortical neuronal migration by mediating NTRK2/TRKB anterograde axonal transport during brain development (PubMed:23576431, PubMed:25944905, PubMed:28259553). Acts as an adapter that bridges the interaction between NTRK2/TRKB and KLC1 and drives NTRK2/TRKB axonal but not dendritic anterograde transport, which is essential for subsequent BDNF-triggered signaling and filopodia formation (PubMed:23576431, PubMed:25944905, PubMed:28259553).[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (6) uses two alternate in-frame splice sites in the 5' coding region compared to variant 1. This results in a shorter protein (isoform f) compared to isoform a. 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.