

Product datasheet for **MC224085**

Magi1 (NM_001083320) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Magi1 (NM_001083320) Mouse Untagged Clone
Tag: Tag Free
Symbol: Magi1
Synonyms: AIP3; Baiap1; BAP1; Gukmi1; Magi-1; MAGI1c; mKIAA4129; TNRC19; WWP3
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC224085 representing NM_001083320
Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGGATCGCC**

ATGCAAAATGCTGGCATAGTCCACCCGGAGAATGAGGAGGAGGAGGATGTCCTGAAATGAACAGTAGCT
TTACAGCCGACTCTGGAGACCAGGACGAGCACACTCTCCAAGAAGCAACGCTCCCGCCTGTGAATAGTAG
CATCCTCGCTGCTCCCATCACGGACCCTTCTCAGAAGTCCCTCAGTACCTACCTCTTTCTGCAGAGGAT
AATTTAGGTCCTCTACCTGAAAACCTGGGAGATGGCCTATACTGAAAATGGAGAAGTCTATTTTCATAGACC
ACAACACGAAAACAACATCATGGTTAGACCCTCGGTGCCTGAACAAACAGCAGAAGCCTCTGGAAGAATG
TGAAGATGATGAAGGGGTACACACCGAGGAGCTGGACAGTGAAGTAGAGTTGCCTGCTGGCTGGGAAAAG
ATTGAAGACCCTGTCTACGGTGTCTACTATGTAGACCACATCAACAGGAAGACGCAATATGAAAACCCAG
TCCTAGAAGCCAAACGGAAGAAACAGCTTGAACAGCAGCAGCAACAGCAGCAGCCTCAGCCACCCGAGCC
AGAAGAGTGGACAGAGGATCATGCATCTGTTGTGCCTCCTGTTGCTCCTCCCATCCCCGAGCAATCCG
GAGCCAGCCAGGGAAACTCCACTTCAGGGCAAACCTTTTTTTACAAGAAACCCCTCTGAGTGAAGGCA
AGTTCATTCACACGAAGCTACGAAAAGCAGCCGAGGCTTTGGCTTCACGGTGGTTGGAGGAGACGAGCC
TGATGAGTTCCTGCAGATCAAGAGCCTCGTCTCGATGGTCTGCCGCACTGGATGGCAAGATGGAGACA
GGGGATGTAATTGTGAGTGTGAATGACACCTGTGTTTTGGGACACACATGCTCAAGTTGTGAAAATCT
TCCAGTCCATTCCATTGGTGCCAGTGTGGACCTTGAAGTCTGCAGAGGTTATCCATTGCCTTTTGACCC
GGATGACCCTAATAACAAGTTTAGTGACCTCGGTGGCCATTTTGGCAAAGAACCAATTATTGTAATGGA
CAAGAGACCTACGATTACCAGCGAGCCACAGTAGTAAAACAGGCAAAGTCAAGCAGCATGAAGGATGCCA
GGCCAAGCAGCCCTGCTGATGTGGCTTCCAACAGCTCTCATGGTTATCCCAACGACACAGTCTCCTTGGC
TTCTCCATAGCCACCCAGCCAGAGCTAATAACTGTTACATAGTCAAAGGGCCAATGGGATTTGGCTTT
ACGATCGCAGACAGTCCCGGTGGGGTGGCCAAAGAGTGAACAGATTGTTGACAGTCCAGCTGCAGAG
GCCTCAAAGAAGGGGATCTTATCGTGGAGGTGAATAAGAAGAACGTGCAGGCCCTGACGCACAATCAAGT
CGTGGATATGCTGATTGAATGTCCAAAGGGAAGTGAAGTCAACTGTTGGTGCAGCGAGGAGGGCTACCA
GTTCCAAAGAAGAGCCAAAGTCGCCACTGGAGAGGAAAGACAGCCAGAATAGCTCCAGCACAGCGTCT



CCAGCCACCGGAGCCTGCACACTGCGTCCCCGAGCCACGGCATACAGGTGCTCCCTGAGTACCTACCTGC
 AGACGCCCTGCTCCAGATCAGACCCGACAGCTCTGGGCAGAAAAAGCCAGATCCTTTTAAAAATCTGGGCC
 CAGTCCAGGAGCATGTATGAAAACCGACCTATGTCACCTTCGCCTGCATCAGGATTGAGCAAGGGTGAAA
 GAGACAGAGAAATCAATTCCACGAATTTTGAGAAATGTCAGATTCAGATTACCAGGAACAGGACATCTT
 CCTCTGGAGAAAAGAAACCGGATTTGGATTTAGGATTCTGGGTGGAATGAACCAGGGGAACCCATTTAT
 ATCGGTACATCGTACCGCTGGGTGCTGCTGACACAGACGGCCGCTGAGGTCTGGAGATGAATTAATCT
 GTGTGGATGGGACACCAGTAATTGGGAAATCACACCAGCTCGTGGTCCAGCTTATGCAACAGAGTGCCAA
 GCAAGGCCATGTCAATCTCACAGTGAGGCGGAAAGTGGTCTTTGCCGTCCCCAAAGCAGAGAATGAGGTG
 CCCTACCCAGCCTCATCACACCAGTAGCAACCAGCCGCGTCCCTGACGGAGGAGAAACGCACACCCG
 AAGGCAGCCAGAACTCTCTGAACACTGTGAGCTCTGGCAGCGGCAGCACCAGTGGCATTGGCAGTGGTGG
 CGGCGGGGGCAGCGGTGTGGTGAAGCCTGTGCTGCAGCCCTATGATGTGGAGATTCGGCGTGGGAGAAC
 GAGGGCTTTGGGTTTGCATCGTGTCTCCGTGAGCAGACCCGAAGCGGGCACAACCTTCGCAGGCAATG
 CATGTGTGGCTATGCCTCACAAAATAGTCCGATTATTGAGGGGAGCCCTGCTGACCCTGTGGCAAGCT
 GAAAGTAGGAGACCGGATCTTGGCAGTAAATGGATGTTCCATACCAACAAATCCCATTCTGACATTGTC
 AACCTAATCAAAGAAGCGGGCAACACAGTGACTCTCCGCATCATCCCCGGGATGAGTCTCAAATGCCA
 CGCTGCTGACTAATGCTGAGAAGATTGCCACCATCACCACCCTCATGCCCCCTCTCAGCAGGGGACCCA
 GAAACAAGGACCACCACAAACCAAGCAGGATTCTCAGTTTGAAGTCAAAAGGACCCGAGGCTGCACAG
 GAGCAAGATTTCTACTGTGGAATTGGAAAGAGGGGCCAAGGGATTTGGCTTTAGTCTTCGAGGGGGCC
 GAGAATAAATGATGATCTTTATGTTCTGCGCTTGGCAGAGGATGGTCTGCAGAAAGATGTGGGAAGAT
 GAGGATTGGCGATGAAATTTAGAGATCAATGGTGAAGCACCACAAAACATGAAACACTCTCGGGCCATA
 GAACTGATCAAGAATGGCGGCCGAGGGTCCGTCTGTTTCTGCGGGGGAGACGGCTCAGTCCCAGAAT
 ATGACCCAGCAGCGACAGGAACGGCCCTCCACCGGTGCACAAGGTGTTCCGGAAGTGAGGCCGGGGCC
 GCCAGACCACAGACCGCATCCAGCCTTGGAGTCCAGTTACCCACCCGAACCTCACAAATCATCAACAT
 GCCGAAAAGCGAGCACACGCGAAGGATCCAAAAGGCAACAGGGAGCACAGCAAAACAACCCAAACGAAATC
 ACACCTGGAATGGAATTCTAGAAAACAGGACAGCGGGGCATGCCGGCCAAAGACCGGCCACCCGACGC
 ATGGAGAGAGGCGCAGCCAGAGCGGACAGCCCAATGTTCAAAGAGGAGGTACCAGGAGAAGCGCAGG
 GAAGGCACCCGACGCTGACAACACTTTGAAAGAAGGGAGAAGCATGAGAAGAGAAGAGATATCTC
 CCGAGAGGAAGCGAGAGCGTTACCCACCCGGAGAAAAGATAGCTCCCCGAGCCGCGGGCAGGTCCT
 CGAAAGACTCCTGGATCAAAGACGGTCCCAGAGCGCAGAAGAGGGGGCTCCCCGAGAGGAGAGCCAAA
 TCCACCGACAGGAGGAGGCCAGGTCCCCTGAGCGCAGGCGAGAGCGGTCACTGGACAAAAGGAACCGGG
 ACGACAAGGTTGGCCACCAGAAAAGAGAGGAGGCTGGTCTGAAGCTGGAAGCGGGGAGAAGCCCCGAAA
 TCCCCAGAGCAGAGAAGGCGGCCTTACAAAGAATGTAGCACCGACCTCAGCATCTGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

Sgfl-MluI

ACCN:

NM_001083320

Insert Size:

3768 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_001083320.2](#), [NP_001076789.1](#)

RefSeq Size: 7181 bp

RefSeq ORF: 3768 bp

Locus ID: 14924

Cytogenetics: 6 D1

Gene Summary: May play a role as scaffolding protein at cell-cell junctions. May regulate acid-induced ASIC3 currents by modulating its expression at the cell surface.[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (2) uses an alternate 5' exon structure and has multiple differences in the coding region, compared to variant 3. It encodes isoform b, which is shorter than isoform c. 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.