

Product datasheet for **MR224465**

Magel2 (NM_013779) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Magel2 (NM_013779) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Magel2
Synonyms:	Mage-l2; NDNL1; nM15; ns7
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>MR224465 representing NM_013779, codon optimized. Due to the complexity of NM_013779, the ORF clone is codon optimized for mammalian Expression. The nucleotide sequence differs from the reference sequence, yet the amino acid sequence remains identical.

Red=Cloning site Blue=ORF Green=Tags(s)

TTTGTGAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGCATCGC**

ATGTCCCAGCTTTCCACCAACCTGGGTGATTCATCACCTCCGGAATCACCAGTGCCCGCAGTCCACTCCA
GGCCACAGTACTGATGCGGGCTCCTCCCGCTAGTTCAAGGGCTCCTCCTGTCCCGTGGGATCCTCCCC
CGTTGATCTCCAGGCCCTATGGCTGCATGGCAGGCCCCAGCCAGCATGGGAAGCTCCGGAGGGTCAG
CTGCCTGCGCCTGTGCCCCAGCTGGCTCAGCCTCCAGGGCTGGGCGCACCTATGGTTTCAGGCGCCCCCTC
TGGGTGGCGCATGGCTAAACCACTACTCCAGGAGTCTGATGGTGCACAGCCACCCCTGGCGCGCC
TATGGCTCAGTCAAGCAGCCCCGGCGTGTGATGCTGCACCTTCTGTGACTGGCGCCCCCTGGCTCAC
CCCCCCCCACCGGTACCCCATGACACACCCACCCGGGACATCCATGGCCATCCCCCTCTCTCCGC
CACCTCCGCTCCCCCCCCACCTCCGGGACACCCATGACTACCCACCTCCTCCGGGAACACCTATGGG
ACACACCCCCACCGGAAATCCGATGACCCACCCCTCCTGGCAATCCTATGGTTTCACCCCTCAC
CATGGCGCGCCTATGGTGCATGGAGTCCCCATGGGACTCCTATGCCACATGTGCCATCACGGGACCC
CTATTGCTCAGCAACCGACACCCGGTGTCTGATGGCGCAGCAGTTACTCCCGAGTGCTCATGGTGCA
GCCTCCTGCTCCTGGAGCACAATGGTGCAGCCCCCTCAGGCTGCCCTCATGACCAACAGCCCCA
AGTATAACACCAATGGCAAAGCCTCCTGGACCCGGCGTGGTAATGATACATCCTCCAGGCGCTCGAGGCC
CCATTATTCAGACACCCGTATCCGGCGCTCCTATGGCTCAAACCGTGCTCCCCCGGGTCAGCCTTTGGC
CACTTGGGCACACAGGCCAGCCCTCATCTGCAGATCCAAAGCCAGGTCATCCGCGCCCCCTCAG
GTGCCAGCGTCCCCAGGCACCTCAGGTTAGCTTGCCACACCTCCTGGATGGCAGGCGACAACGCCAA



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ATTGGCAGGTCACCCCCAGGGATGGCCGGCTACTCCTCTGACGTGGCAGGCAACACAAGTGACCTGGCA
 GGCGCCAAACCATTCATGGCAGGCCACACAGCCAGGCAGGCGGCACTCCACAATCAGGACAGGGCAT
 ACACCCATCCGCCAGGCCCCGCACCGCTCCTGCGGCAGATCCCGCCTATGATCAGGCAAATTCAGCCCG
 TCATGAGGCAGGCCCCCACTTATTGCGCAGGTGCCAATCCGCCCGCGCCACATGGGATCGCCAGCCA
 GCCTCAGCTGTGGCAGGTGCTCCCTCCACCACCACCTTGCGGCAGGCCCCCAGGCACGGCTTCTCCTT
 CCACGAGTGCCCGGTACGGGGCAGGTGAGCAGGTCCCTCCTGTTGCCAGATTACCTGGTGCCCCAAA
 GCGGACCACAGGTTCCCAAACAGTGCTGCCAGCCAGCTTAGTATTCATCCCTGTACCCCAAGCAGC
 CGCCAGTCCGCCCGCCGCACTGTGCATTGCCCGCGATCATTTGGCAAGCTCCCAAAGGTCAGGCTCCC
 GTCCCTCAGGAATTCCCGTGCCACAAGAGCTGCCAGTACCCAGGAGCTGCCCGTTCCCGAGGAGTGC
 CAGTTCCACAGGAGATCCCGTGCTCAGGAGATCCAGTTTCTCAGGAAGTCCAGTACCCCAAGAACT
 TCCGGTGCTCAGGAGCTGCCGGTGCTCAAGAGCTGCCTGTCCACAGGAATTCCTGTCCCTCAGGAG
 CTGCCAGTGCCACAGGAGCTGCCGTACCACAGGAGCTGCCAGTCCCACTGGAGTTCCAGGAGGTCAAC
 AGGCTCAGGCAGTCGGATGGCGCGCTCTAAAGTCCACCCCACTTTTGGCAGCCTGTGAGCGCACAAGA
 GGCCAGGAACAAGCAACCCAGATTGCGCACGTGGAACAACAGCAGCCTTCCAGGGAGCCCCCGCATCC
 TCTAAAGCGCTGCAAAACCAACTCCCACTCACAAGCCAGGCCTCCGGCTCCAGGCGGAGCTGCCTT
 CCGTGACGCTGCAACCCAGCTGGCAGGGTCTCTGCCCATGTTGCAGGCCAGCCCGGGGCTTCAGCAAC
 CCTTGCTAACTTTCTAGGGGTCCACCAGAAGCAGAATGGCCCCAAGCGGTGAACCGGGTCCATCTTCC
 CTTGAGCCGCGGGGACCCACGAGAACGGCGAGCACCCGCCCGCAGAAAAAGGGCCCTCCGAAAGAAA
 GGATGTTTCATCGAGCGACCTTTTGTGCCCCAGGGGCGCAAGCGCTCCCGCGCATACGTGCCAACGGC
 ATGGAAGAATCTCCCTGTACCAGCGAAACCTTCCCGGCCACAAGCAGGGTATTCCTTCCACAAGTCAC
 TTTAGCCCGCTCATCTAATGCCTTTCGCGGTCTAGCGCCGCTAGTGAGTCTCCTAAAAGCTTGCCCT
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 CAACGCTAGTTCTGCATGCAAACTCGTCCAGCAATACTGATGGTGCCGCTGCTGCCCGCAGGCATCT
 GCTACGGCCGCGGAAGCCTCTAAGAGCTGTAGCCCCCTCGGCGGCCCGCAAGGCTACAAGGAAGAAGA
 AGCATCTGGAGCCAAAGGAGGACAACCTGTGGACATAGGCTGAGTAGCAGAGACTGGAGAGTCCCGCAC
 ATGGGGCAATCTAGTCACAGCGACTGGGAAATCCAACGCGCAATGCAACTGCTCGGGGACCGGAGTCA
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 AGTTCTAGCGTATCCAGAGGCTCCTCTGCGGCACAAGAAGATCCTGACCGGAGAGCCAGCCACTGAGT
 CCCCTGGATGAAAGAGCTAATGCTCTCGTGCAATTTCTGTTGGTTAAGGACCAGGCTAAGGTTCCGGTGC
 AGCTCAGTGAGATGGTCAATGTGGTCATCCGGGAGTACAAGGACGACTACTGGACATCATCAACCGCGC
 CAATACGAAACTCGAATGCACATTCCGGGTGTCAACTCAAGGAGGTTGATACCAAAACGCACACTTACATC
 ATAGTTAATAAGATGGCCTACCCACAGTGCAACCTGTTGGCTTCTATCTGGAACGCCCTAAATTTAGCC
 TCCTGATGGTGGTCTGTCACTCATTTTCATGAAGGGATATTGTATTCCGGGAGAACTGTCTTTCAGCTT
 TCTGTTCCAGCTGGGTCTGGATGTACAAGAGACAAGCGGACTCTTTCGAATTACAAGAACTGATTACT
 AGCGTCTTTGTACGGCATCGATACCTGGAGTACCGACAGATCCCCTTTACTGAGCCTGCTGAGTACGAGC
 TTCTCTGGGGCCCCGAGCGTTTTTGGAGACTAACAGAGTGCACATTCTGAGATTTTGGCGGCGCTTTA
 CGAAAAACAACCTCAAATATGGTCATGTACGTACCTGGACAGCCTCGCCGAGTTGGAATACAAGGACGCG
 AACGCCGACGTGAGGAGAGCCATGATTCAGACGACGATGCACACGACCCACATCCTCCCCTCACCCCC
 AC

ACGCGTACGCGGCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >MR224465 representing NM_013779
Red=Cloning site Green=Tags(s)

MSQLSTNLGDSSPPESPVPAVHSRPTVLMRAPPASSRAPPPVWDPPPVDLQAPMAAWQAPQPAWEAPEGQ
LPAPVAQLAQPPLGAPMVQAPPLGGMAKPPTPGVLMVHQPPGAPMAQSSTPGVLMHPSVTGAPLAH
PPPPGTPMTHPPGTSMHPPPPPPPPPPPPPGTPMTHPPPPGTPMGHPPPGNPMTHPPPGNPMVHPLT
HGAPMVHGGPHGTPMPHVPIITGTPIAQQTTPGVLMQAQQLTPGVLMVQPPAPGAPMVQPPQAALMTQPAP
SITPMAKPPGPGVVMIHPPGARGPIIQTPTVSGAPMAQTVLPPGQPLATWAPQGQPLILQIQSQVIRAPPQ
VPSVPQAPQVQLATPPGWQATTNMQVTPQGW PATPLTWQATQVTWQAPTIAWQATQPGRQGHSTIRTGH
TPIRPGAPALLRQIPPMIRQIQVPMRQAPPLIRQVPIRPAPHGIASQPQLWQVLP PPPPLRQAPQARLLL
PRVPGTGQVSTVPPVAQIHLVPQSGPQVPQTVLPAQLSIPVQAAAQSAPRTVHCPII IWQAPKGQAP
VPQELPVPQELPVPQELPVPQELPVPQELPVPQELPVPQELPVPQELPVPQELPVPQELPVPQELPVPQEL
LPVPQELPVPQELPVPLEFQEVQQAQAVGWRAPKVP PHFWQPVSAQEAQEQATQIAHVEQQQPFQGAPAS
SKALQTQLPTHQAQASGLQAE LPSVQLQPSWQG LPLMLQAQPGASATLANFPRGSTRSRMAPSGEPGPSS
LEPRGPPRRRAPARDKKGPPKERMFIGATFCAPRGASASRAYVPTAWKNLPATSETFPATSRVFPSTSH
FQPASSNAFRGPSAAASESPKSLPFALQDPYACVEALPAVPWVPYPDGNASSACKSVPAILMVAAAAPQAS
ATAAEASKSSEP RRRPGKATRK KKHLEPKEDNCGHRLSSRDWRGPRTWGNPSHSDWEIQRAMQLLDRES
LYTPQGLNDWGCNTRSMPRSLEGPSTSRDQEF CGDSGGSQTWMASEVPSVSRGSSAAQEDPDRESQPLS
PLDERANALVQFLLVKDQAKVPVQLSEMVNVVIREYKDDSLDIINRANTKLECTFGCQLKEVDTKHTYI
IVNKMAYPQC NLLASYL ERPKFSLLMMVLSLIFMKG YCIRENLLFSFLFQLGLDVQETSGLF RITKKLIT
SVFVRHRYLEYRQIPFTEPAEYELLWGPRAFLETNRVHILRFLAALYENQ PQIWSQCYLDSLAELEYKDA
NAAAEESHSDDDAHDPTSSPHPH

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites:

Sgfi-MluI

Cloning Scheme:

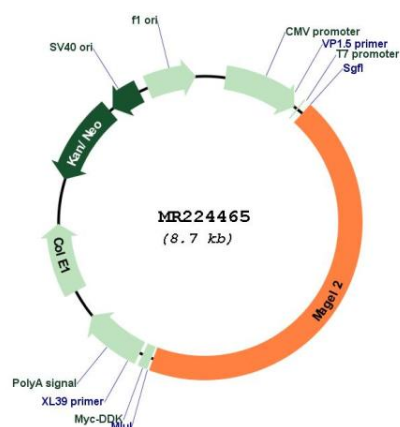


ACCN: NM_013779

ORF Size: 3852 bp

OTI Disclaimer:	The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. More info
OTI Annotation:	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_013779.2 , NP_038807.4
RefSeq Size:	4662 bp
RefSeq ORF:	3855 bp
Locus ID:	27385
UniProt ID:	Q9QZ04
Cytogenetics:	7 34.37 cM
MW:	138 kDa
Gene Summary:	Probably enhances ubiquitin ligase activity of RING-type zinc finger-containing E3 ubiquitin-protein ligases, possibly through recruitment and/or stabilization of the Ubl-conjugating enzyme (E2) at the E3:substrate complex. Acts as a regulator of retrograde transport via its interaction with VPS35. Recruited to retromer-containing endosomes and promotes the formation of 'Lys-63'-linked polyubiquitin chains at 'Lys-220' of WASHC1 together with TRIM27, leading to promote endosomal F-actin assembly (By similarity). Regulates the circadian clock by repressing the transcriptional activator activity of the CLOCK-ARNTL/BMAL1 heterodimer. Significantly promotes the cytoplasmic accumulation of CLOCK (PubMed:22208286).[UniProtKB/Swiss-Prot Function]

Product images:



Circular map for MR224465