

Product datasheet for **MR210511**

MIh1 (NM_026810) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	MIh1 (NM_026810) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	MIh1
Synonyms:	1110035C23Rik; AI317206; AI325952; AI561766
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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**ORF Nucleotide
Sequence:**

>MR210511 ORF sequence
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGCATCGCC**

ATGGCGTTTGTAGCAGGAGTTATTTCGGCGTCTGGACGAGACGGTGGTGAACCGCATAGCGCGGGGAAG
 TCATTCAGCGGCCGCCAATGCTATCAAAGAGATGATAGAAAACGTGTTAGATGCAAAATCTACAAATAT
 TCAAGTGGTTGTTAAGGAAGGTGGCCTGAAGCTAATTCAGATCCAAGACAATGGCACTGGAATCAGGAAG
 GAAGATCTGGATATTGTGTGTGAGAGGTTCACTACGAGTAAACTGCAGACTTTTGAGGATTTAGCCAGTA
 TTTCTACCTATGGCTTTTCGTGGTGGAGCATTGGCAAGCATAAGCCATGTGGCCCATGTCATTACAAC
 CAAAACAGCTGATGGAAATGTGCGTACAGAGCAAGTTACTCAGATGGAAAGCTGCAAGCCCTCTCTAAA
 CCCTGTGCAGGCAACCAGGGCACCTGATCACGGTGAAGACCTTTTTTACAACATAATCACAAGGAGGA
 AAGCTTTAAAAAATCCAAGTGAAGAGTACGAAAAATTTTGAAGTTGTTGGCAGGTATTCAATACACAA
 TTCAGGCATTAGTATCTCAGTAAAAAACAAGGTGAGACAGTATCTGATGTCAGAACACTGCCAATGCC
 ACAACCGTGGACAACATTCGCTCCATCTTTGAAATGCGGTTAGTCGAGAAGTATAGAAGTTGGGTGTG
 AGGATAAAACCTAGCTTTCAAATGAATGGCTATATATCGAATGCAAACTATTCAGTGAAGAAGTGCAT
 TTTCTACTCTTCATCAACCACCGTCTGGTAGAATCAGCTGCCTTGAGAAAAGCCATTGAAACTGTATAT
 GCAGCATATTTGCCAAAAACACACACCCATTCTGTACCTCAGTTTGGAAATCAGCCCTCAGAAGCTGG
 ACGTCAATGTACACCCCAAGCAGCAAGTTCACCTTTCTGCATGAGGAGAGCATTCTGCAGCGTGTGCA
 GCAGCACATTGAGAGCAAGCTGCTGGGCTCCAATTCCTCCAGGATGTATTCACCCAGACCTTGCTTCCA
 GGACTTGCTGGCCCTCTGGGGAGGCAGCTAGACCCACGACAGGGGTGGCTTCTCATCCACTATGGAA
 GTGGCGACAAGGTCTACGCTTACCAGATGGTCCGTACGGACTCCCGGGATCAGAAGCTTGACGCCTTCT
 GCAGCCTGTAAGCAGCCTTGTGCCAGCCAGCCAGGACCTGCCCTGTCCGAGGGGCCAGGACAGAG
 GGCTCTCTGAAAGGGCCACGCGGGAGGATGAGGAGATGCTTGCTCTCCAGCCCCGCTGAAGCAGCTG
 CTGAGAGTGAGAACTTGAGAGGGAATCACTAATGGAGACTTCAGACACAGCCAGAAAAGCGGCACCCAC
 TTCCAGTCCAGGAAGCTCCAGAAAAGAGACATCGGGAGGACTCTGATGTGGAAATGGTGGAAAATGCTTCC
 GGGAAAGGAAATGACAGCTGCTTGCTACCCAGGAGGAGGATCATTAACTCACCAGCGTCTTGAGTCTCC
 AGGAAGAGATTAGTGAGCGGTGCCATGAGACTCTCCGGGAGATGCTCCGTAACCATTCCTTTGTGGGCTG
 TGTGAATCCTCAGTGGGCTTGGCACAGCACCAGACCAAGCTATACCTCCTCAACACTACCAAGCTCAGT
 GAAGAGCTGTTCTACCAGATACTATTTATGATTTTGCCAACTTTGGTGTCTGAGGTTATCGGAACCAG
 CGCCACTCTTCGACCTGGCCATGCTGGCCTTAGACAGTCCCTGAAAGTGGCTGGACAGAGGACGACGCC
 GAAGGAAGGGCTTGCAGAGTACATTGTCGAGTTTCTGAAGAAGAAAGCGGAGATGCTTGCAGACTATTTT
 TCTGTGGAGATCGATGAGGAAGGAACTGATCGGATTACCTCTTCTGATTGACAGCTATGTGCCACCTT
 TGGAGGGACTGCCTATCTTCATTCTTCGACTGGCCACTGAGGTGAATTGGGATGAAGAAAAGGAGTGTTT
 TGAAAGTCTCAGTAAAGAATGTGCTATGTTTTACTCCATTCGGAAGCAGTATACTGGAGGAGTCGACC
 CTCTCAGGCCAGCAGAGTGACATGCCTGGCTCCACGTCAAAGCCCTGGAAGTGGACTGTGGAGCACATTA
 TCTATAAAGCCTCCGCTCACACCTCTACCTCCGAAGCATTTCACAGAAGATGGCAATGTCTGACGCT
 TGCCAACCTGCCAGATCTATACAAGTCTTTGAGCGGTGT

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >MR210511 protein sequence
 Red=Cloning site Green=Tags(s)

MAFVAGVIRRLDETVVNRIAAGEVIQRPANA KEM IENCLDAKSTNIQVVVKEGGLKLIQIQDNGTGIRK
 EDLDIVCERFTTSKLTQFEDLASISTYGRGEALASISHVAHVTTITKTADGKCAYRASYS DGKLQAPPK
 PCAGNQGLTITVEDLFYNIITRRKALKNPSE EYGKILEVVGRYSIHNSGISISVKKQGETVSDVRTLPNA
 TTVDNIRSIFGNAVSRELI EVGCEDKTLAFKMNGYISNANYSVKKCIFLLFINHRLVESAA LRKAIETVY
 AAYLPKNTHPFLYL SLEISPNVDVNVHPTKHEVHFLHEESILQRVQQHIESKLLGSNSSRMYFTQTLLP
 GLAGPSGEAARPTTG VASSTSGSGDKVYAYQMVRTDSRDQKLD AFLQPVSSLVPSQPQDPAPVRGARTE
 GSPERATREDEEMLALPAPAEAAA ESENLERESLMETSDTAQKAAPTSSPGSSRKRHRESDVEMVENAS
 GKEMTAACYPRRRIINL TSVLSLQEEISERCHETLREMLRNHSFVGCVNPQWALAQHQTKLYLLNTTKLS
 EELFYQILYDFANFVLR LSEPAPLFDLAMLALDSPESGWTEDDGPKEGLAEYIVEFLKKAEM LADYF
 SVEIDEEGNLIGLPLLIDS YVPPLEGLPIFILRLATEVNWDEEKECFESLSKECAMFYSIRKQYILEEST
 LSGQQSDMPGSTSKPWKWTVEHIIYKAFRSHLLPPKHFTEDGNVQLANLPDLYKVFERC

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites: SgfI-MluI

Cloning Scheme:

Cloning sites used for ORF Shuttling:



* The last codon before the Stop codon of the ORF

ACCN: NM_026810

ORF Size: 2283 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.

RefSeq: [NM_026810.1](#), [NP_081086.1](#)

RefSeq Size: 2598 bp

RefSeq ORF: 2283 bp

Locus ID: 17350

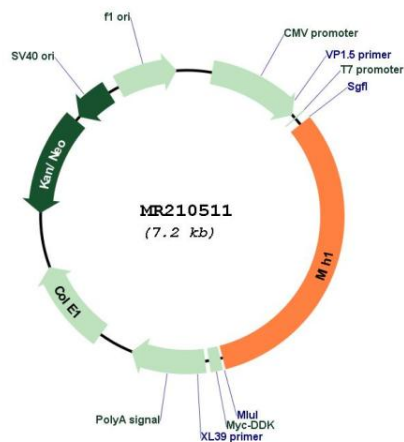
UniProt ID: [Q9JK91](#)

Cytogenetics: 9 60.92 cM

MW: 84.7 kDa

Gene Summary: Heterodimerizes with Pms2 to form MutL alpha, a component of the post-replicative DNA mismatch repair system (MMR). DNA repair is initiated by MutS alpha (Msh2-Msh6) or MutS beta (MSH2-MSH3) binding to a dsDNA mismatch, then MutL alpha is recruited to the heteroduplex. Assembly of the MutL-MutS-heteroduplex ternary complex in presence of RFC and PCNA is sufficient to activate endonuclease activity of Pms2. It introduces single-strand breaks near the mismatch and thus generates new entry points for the exonuclease EXO1 to degrade the strand containing the mismatch. DNA methylation would prevent cleavage and therefore assure that only the newly mutated DNA strand is going to be corrected. MutL alpha (Mlh1-Pms2) interacts physically with the clamp loader subunits of DNA polymerase III, suggesting that it may play a role to recruit the DNA polymerase III to the site of the MMR. Also implicated in DNA damage signaling, a process which induces cell cycle arrest and can lead to apoptosis in case of major DNA damages. Heterodimerizes with Mlh3 to form MutL gamma which plays a role in meiosis (By similarity).[UniProtKB/Swiss-Prot Function]

Product images:



Circular map for MR210511