

## Product datasheet for **MC222681**

### **Msh2 (NM\_008628) Mouse Untagged Clone**

#### **Product data:**

Product Type:	Expression Plasmids
Product Name:	Msh2 (NM_008628) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Msh2
Synonyms:	A1788990
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



[View online »](#)

**Fully Sequenced ORF:** >MC222681 representing NM\_008628  
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**CGGATCGCC**

ATGGCGGTGCAGCCTAAGGAGACGCTGCAGTTGGAAGGCGGCCGAGGCGGGCTTCGTGCGCTTCTTTG  
 AGGGCATGCCGGAGAAGCCGAGCACACCGTGCCTCTTCGACCGCGGGGACTTTTACACGGCGCACGG  
 AGAGGACGCGCTGCTGGCGGCCCGGAGGTGTTCAAGACCCAGGCGTGATCAAGTACATGGGGCCGGCA  
 GGGAGTAAGACCCTGCAGAGTGTGTGCTTAGCAAGATGAACTTTGAGTCTTTCGTGAAAGATCTTCTTC  
 TGGTTCGCCAGTATCGAGTTGAAGTTTATAAGAAATAAGCTGAAATAAGGCGTCTAAGGAGAATGAGTG  
 GTATCTGGCATTAAAGCTTCTCCCGCAATCTTCTCAGTTTGAAGACATCCTGTTTGGTAAACAATGAC  
 ATGTCAGCTTCCGTTGGCGTTATGGGTATTAATAAGGCGGTGTTGATGGTCAAAGACATGTTGGAGTTG  
 GGTATGTGGATCCACCCAGAGGAAGCTAGGCTTGTGTGAGTTCCCGAGAATGATCAGTTCTCCAATCT  
 CGAGGCTCTTCTGATTGAGTGGACCAAGGAATGCGTTTTACCAGGAGGAGAGACTACTGGAGACATG  
 GGGAAACTGAGGCAAGTTATCCAGAGAGGAGGATTCTGATCACAGAAAGAAAGAGAGCCGACTTTTCCA  
 CTAAGACATTTATCAGGATCTCAACCGTTACTGAAAGGCAAAAAGGAGAACAGATAAATAGTCTGC  
 CCTACCAGAGATGGAGAATCAGGTTGCAGTTTCATCACTATCTGCAGTAATCAAGTTTTTGAAGCTTTA  
 TCAGACGATCAAATTTTGGCAGTTTGAAGTGGCCACTTTTACTTCAGCCAGTACATGAAGTTGGACA  
 TGGCAGCAGTTAGAGCCCTCAACCTTTCCAGGGTTCTGTTGAAGACACCACTGGCTCTCAGTCTCTGGC  
 CGCATTATTGAATAAATGCAAACTGCTCAAGGACAAAGATTGGTTAACCAGTGGATCAAGCAGCCGCTC  
 ATGGATAGGAACAGGATAGAGGAGAGTTAAATTTAGTGAAGCTTTTGTGAGGATTGAGAAGTGGAGG  
 AGAGTTTACAGGAGGATTTGCTTCGCGGTTCCAGACTTAAACCGCTTCCAGAAATAAAGGAGGAGG  
 AGCAGCGAATTTACAAGACTGTTACCGACTGTATCAGGGTATTAACCAGCTCCCGAGCTATCCAGGCT  
 CTGGAGAAATACGAAGGAAGACACCAGGCACTGTTGTTGGCAGTTTTTGTGACTCCTCTTATTGATCTTC  
 GTTCTGATTTTTCAAATTTCAAGAAATGATAGAAACAATTTAGATATGGATCAGGTGGAAAACACGA  
 GTTCTTGTAAAACCTTCATTTGATCCTAACCTGAGTGAAGTAAAGAGAAGTATGGATGGCTGGAGAAG  
 AAGATGCAGTCCACCTTAATAAATGCAGCCGGGGCTCGGATTGGATCCTGGCAACAGATTAAATGG  
 ACTCCAGTGCACAGTTGGATATTTTCCGTGAACCTGCAAGGAAGAGAAAGTCTTCGCAACAACAA  
 GAACCTCAGCACAGTGGACATCCAGAAGAATGGCGTGAAGTTACCAACAGTGAATTGCCTCTTAAAT  
 GAAGAATACTAAGAACAAGGCGAGTATGAAGAGGCCAGGATGCCATTGTTAAAGAAATGTCAATA  
 TTTCTCAGGCTACGTAGAGCAATGCAGACGCTCAACGATGTGCTGGCTCACTTAGACGCCATTGTTAG  
 CTTGCTCATGTGTCAAACGCAGCACCCGTTCTTATGTACGACCAAGTCACTTGGAGAAAGGAAAAGGG  
 AGAATTATATTGAAAGCCTCCAGGCATGCTTGTGTTGAAGTTCAAGATGAAGTTGCATTTATTCAAATG  
 ACGTGCATTTGAAAAGATAAACAGATGTTCCACATCTACTGGTCCCAATATGGGAGGTAATCAAC  
 ATACATTCGTGACACCGGGTATTGACTCATGGCCAAATCGGGTGTGTTGTGCCCTGTGAGTCGGCA  
 GAAGTGTCCATTGTGGATTGCATCCTTGCTCGAGTCGGGGCTGGTACAGTCAACTGAAAGGCGTCTCCA  
 CATTGATGGCTGAAATGCTGGAGACTGCTTCCATCCTCAGTCAAGCAACCAAGACTCCTTAATAATCAT  
 TGATGAGCTGGGAAGAGGAACCTACCTATGATGGATTTGGTTAGCATGGGCTATATCAGATTACATT  
 GCAACGAAGATTGGTGCCTTTTGATGTTTGCACCCATTTTTCATGAACCTACTGCTTTGGCAACCAAA  
 TACCAACTGTTAATAATCTACATGTCACAGCGCTCACTACTGAGGAGACCCTAATATGCTTTACCAAGT  
 GAAAAAAGGTGTCTGTGATCAGAGTTTCGGGATTCACGTGGCTGAGCTCGCTAATTTCCCGAGGCAGTG  
 ATAGCGTGCAGCAAGCAGAAGGCTCTAGAGCTTGAAGAATTTGAAACATTGGAACCTCGCTGGGATGTG  
 ACGAAGCCGAGCCGGCTGCAAGAGACGCTGCCTGAAAGAGAGCAAGGTGAGAAAATTTCTGGAGTT  
 CCTGTGCAAGGTCAAGCAGGTGCCCTTACTGCCATGTCGGAGGAGAGCATCTCCGGAAGCTGAAGCAA  
 CTGAAAGCCGAGGTGGTGCAGAAACAACAGCTTCGTAACGAGATCATTTACGGATAAAGGCTCCGG  
 CTCCTGA

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** SgfI-MluI

<b>ACCN:</b>	NM_008628
<b>Insert Size:</b>	2808 bp
<b>OTI Disclaimer:</b>	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).
<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_008628.2</a> , <a href="#">NP_032654.1</a>
<b>RefSeq Size:</b>	3056 bp
<b>RefSeq ORF:</b>	2808 bp
<b>Locus ID:</b>	17685
<b>UniProt ID:</b>	<a href="#">P43247</a>
<b>Cytogenetics:</b>	17 57.87 cM
<b>Gene Summary:</b>	Component of the post-replicative DNA mismatch repair system (MMR). Forms two different heterodimers: MutS alpha (MSH2-MSH6 heterodimer) and MutS beta (MSH2-MSH3 heterodimer) which binds to DNA mismatches thereby initiating DNA repair. When bound, heterodimers bend the DNA helix and shields approximately 20 base pairs. MutS alpha recognizes single base mismatches and dinucleotide insertion-deletion loops (IDL) in the DNA. MutS beta recognizes larger insertion-deletion loops up to 13 nucleotides long. After mismatch binding, MutS alpha or beta forms a ternary complex with the MutL alpha heterodimer, which is thought to be responsible for directing the downstream MMR events, including strand discrimination, excision, and resynthesis. Recruits DNA helicase MCM9 to chromatin which unwinds the mismatch containing DNA strand. ATP binding and hydrolysis play a pivotal role in mismatch repair functions. The ATPase activity associated with MutS alpha regulates binding similar to a molecular switch: mismatched DNA provokes ADP-->ATP exchange, resulting in a discernible conformational transition that converts MutS alpha into a sliding clamp capable of hydrolysis-independent diffusion along the DNA backbone. This transition is crucial for mismatch repair. MutS alpha may also play a role in DNA homologous recombination repair. In melanocytes may modulate both UV-B-induced cell cycle regulation and apoptosis.[UniProtKB/Swiss-Prot Function]