

## Product datasheet for **RC401866**

### MSH2 (NM\_000251) Human Mutant ORF Clone

#### Product data:

Product Type:	Mutant ORF Clones
Product Name:	MSH2 (NM_000251) Human Mutant ORF Clone
Mutation Description:	G692R
Affected Codon#:	692
Affected NT#:	2074
Nucleotide Mutation:	MSH2 Mutant (G692R), Myc-DDK-tagged ORF clone of Homo sapiens mutS homolog 2, colon cancer, nonpolyposis type 1 (E. coli) (MSH2) as transfection-ready DNA
Effect:	Colorectal cancer, non-polyposis
Symbol:	MSH2
Synonyms:	COCA1; FCC1; hMSH2; HNPCC; HNPCC1; LCFS2; MMRCS2
E. coli Selection:	Kanamycin (25 ug/mL)
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
Tag:	Myc-DDK
ACCN:	NM_000251
ORF Size:	2802 bp
Restriction Sites:	SgfI-MluI



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**ORF Nucleotide Sequence:**

>RC401866 representing NM\_000251  
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**CGGATCGCC**

ATGGCGGTGCAGCCGAAGGAGACGCTGCAGTTGGAGAGCGCGGCCGAGGTCGGCTTCGTGCGCTTCTTT  
 AGGGCATGCCGGAAGAAGCCGACCACCACAGTGCCTTTTTCGACCGGGCGACTTCTATACGGCGCACGG  
 CGAGGACGCGCTGCTGGCCGCCGGGAGGTGTTCAAGACCCAGGGGTGATCAAGTACATGGGGCCGCA  
 GGAGCAAAGAATCTGCAGAGTGTGTGCTTAGTAAATGAATTTGAATCTTTGTAAAAGATCTTCTTC  
 TGGTTCGTAGTATAGAGTTGAAGTTTATAAGAATAGAGCTGAAATAAGGCATCCAAGGAGAATGATTG  
 GTATTTGGCATATAAGGCTTCTCCTGGCAATCTCTCAGTTTGAAGACATTCTTTGGTAACAATGAT  
 ATGTCAGCTTCCATTGGTGTGTGGGTGTTAAATGTCCGAGTTGATGGCCAGAGACAGGTTGGAGTTG  
 GGTATGTGGATTCCATACAGAGGAACTAGGACTGTGTGAATCCCTGATAATGATCAGTTCTCCAATCT  
 TGAGGCTCTCCTCATCCAGATTGGACCAAAGGAATGTGTTTTACCCGGAGGAGAGACTGTGGAGACATG  
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 CAAAAGACATTTATCAGGACCTCAACCGTTGTTGAAAGGCAAAAAGGGAGAGCAGATGAATAGTGTGT  
 ATTGCCAGAAATGGAGAATCAGGTTGCAGTTTCATCACTGTCTGCGGTAATCAAGTTTTAGAACTCTTA  
 TCAGATGATTCCAACCTTTGGACAGTTTGAAGTACTACTTTTACTTTCAGCCAGTATATGAAATTTGATA  
 TTGACAGCTCAGAGCCCTTAACCTTTTTAGGGTCTGTTGAAGATACCACTGGCTCTCAGTCTCTGGC  
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 AGCAGCAAACCTTACAAGATTGTTACCGACTCTATCAGGGTATAAATCAACTACCTAATGTTATACAGGCT  
 CTGAAAAACATGAAGGAAAAACACCAGAAATTATTGTTGGCAGTTTTTGTGACTCCTTACTGATCTTC  
 GTTCTGACTTCTCCAAGTTTCAGGAAATGATAGAAACAACTTTAGATATGGATCAGGTGGAAAACCATGA  
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 AAGATGCAGTCAACATTAATAAGTGCAGCCAGAGATCTGGCTTGGACCCTGGCAAACAGATTAACCTGG  
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 AAATTTAGTACTGTAGATATCCAGAAGAATGGTGTAAATTTACCAACAGCAAATGACTTCTTTAAAT  
 GAAGAGTATACCAAAAAATAAACAGAATATGAAGAAGCCAGGATGCCATTGTTAAAGAAATGTCAATA  
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 AGAATTATATTAAGCATCCAGGCATGCTTGTGTTGAAGTTCAAGATGAAATTCATTTATTCCTAATG  
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 TACCAACTGTTAATAATCTACATGTCACAGCACTACCACTGAAGAGACCTTAACTATGCTTTATCAGGT  
 GAAGAAAGGTGTCTGTGATCAAAGTTTTGGATTGATGTTGCAGAGCTTGCTAATTTCCCTAAGCATGTA  
 ATAGAGTGTGCTAAACAGAAAGCCCTGGAACCTGAGGAGTTTCAAGTATATTGGAGAATCGCAAGGATG  
 ATATCATGGAACCAGCAGCAAAGAAGTCTATCTGAAAGAGAGCAAGGTGAAAAAATTTTCAAGGAGTT  
 CCTGTCCAAGGTGAAACAAATGCCCTTACTGAAATGTCAGAAGAAAACATCACAAATAAAGTTAAACAG  
 CTAAGGCTGAAGTAATAGCAAAGAATAATAGCTTTGAAATGAAATCATTTACGAATAAAAGTTACTA  
 CG

AG**CGGACCG**ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC  
 TGGATTACAAGGATGACGACGA TAAGGTTTAA

**Protein Sequence:** >RC401866 representing NM\_000251  
 Red=Cloning site Green=Tags(s)

MAVQPKETLQLESAAEVGFVRFVFGMPEKPTTTVRLFDRGDFYTAHGEDALLAAREVFKTQGVIKYMGPA  
 GAKNLQSVVL SKMNFESFVKDLLLVRQYRVEVYKNRAGNKASKENDWYLAYKASPGNLSQFEDILFGNND  
 MSASIGVVGVKMSAVDQGRQVGVGYVDSIQRKLGLCEFPDNDQFSNLEALLIQIGPKECVLPGGETAGDM  
 GKLRIIQRGGILITERKKADFSTKDIYQDLNRLLLKGGKGEQMNSAVLPEMENQVAVSSLSAVIKFLELL  
 SDDSNFGQFELTTFFDSQYMKLDIAAVRALNLFQGSVEDTTGSQSLAALLNKCKTPQGGRLVNQWIKQPL  
 MDKNRIEERLNLVEAFVEDAELRQTLQEDLLRRFPDLNRLAKKFQQAANLQDCYRLYQGINQLPNVIQA  
 LEKHEGHQKLLLAVFVTPLTDLRSDFSKFQEMIEITLMDQVENHEFLVKPSFDPNLSELREIMNDLEK  
 KMQSTLISAARDLGLDPGKQIKLDSSAQFGYYFRVTCKEEKVLRNNKNFSTVDIQKNGVKFTNSKLTSLN  
 EEYTKNKTEYEEAQDAIVKEIVNISSGYVEPMQTLNDVLAQLDAVVSFAHVSNGAPVPPYVRPAILEKGGQ  
 RIILKASRHACVEVQDEIAFIPNDVYFEKDKQMFHIITGPNMGGKSTYIRQTGVIVLMAQIRCFVPCESA  
 EVSIVDCILARVGAGDSQLKGVSTFMAEMLETASILRSATKDSLIIIDELGRGTSTYDGFGLAWAISEYI  
 ATKIGAFCMFATHFHELTALANQIPTVNNLHVTALTTEETLTMLYQVKKGVCDQSFGIHVAELANFPKHV  
 IECAKQKALELEEFQYIGESQGYDIMEPAAKKCYLEREQGEKIIQEFLSKVKQMPFTEMSEENITIKLKV  
 LKAEVIAKNNSFVNEIISRIKVT

SGPTRRRL**EQLISEEDLAANDILDYKDDDDKV**

**Restriction Sites:**

SgfI-MluI

**Cloning Scheme:**



<b>OTI Disclaimer:</b>	<p>Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <a href="mailto:custsupport@origene.com">custsupport@origene.com</a> or by calling 301.340.3188 option 3 for pricing and delivery.</p> <p>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. <a href="#">More info</a></p>
<b>OTI Annotation:</b>	<p>This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.</p>
<b>Components:</b>	<p>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).</p>
<b>RefSeq:</b>	<p><a href="#">NP_000242</a></p>
<b>RefSeq Size:</b>	<p>2802 bp</p>
<b>RefSeq ORF:</b>	<p>2805 bp</p>
<b>Locus ID:</b>	<p>4436</p>
<b>Cytogenetics:</b>	<p>2p21-p16.3</p>
<b>Domains:</b>	<p>MutS_V, MutS_I, MutS_III, MutS_II, MutS_IV</p>
<b>Protein Families:</b>	<p>Druggable Genome, Stem cell - Pluripotency</p>
<b>Protein Pathways:</b>	<p>Colorectal cancer, Mismatch repair, Pathways in cancer</p>
<b>MW:</b>	<p>102.7 kDa</p>
<b>Gene Summary:</b>	<p>This locus is frequently mutated in hereditary nonpolyposis colon cancer (HNPCC). When cloned, it was discovered to be a human homolog of the E. coli mismatch repair gene mutS, consistent with the characteristic alterations in microsatellite sequences (RER+ phenotype) found in HNPCC. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Apr 2012]</p>