

Product datasheet for **RC401457**

MSH6 (NM_000179) Human Mutant ORF Clone

Product data:

Product Type:	Mutant ORF Clones
Product Name:	MSH6 (NM_000179) Human Mutant ORF Clone
Mutation Description:	E1163V
Affected Codon#:	1163
Affected NT#:	3488
Nucleotide Mutation:	MSH6 Mutant (E1163V), Myc-DDK-tagged ORF clone of Homo sapiens mutS homolog 6 (E. coli) (MSH6) as transfection-ready DNA
Effect:	Colorectal cancer, non-polyposis
Symbol:	MSH6
Synonyms:	GTBP; GTMBP; HNPCC5; HSAP; MMRC3; p160
E. coli Selection:	Kanamycin (25 ug/mL)
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
Tag:	Myc-DDK
ACCN:	NM_000179
ORF Size:	4080 bp
Restriction Sites:	SgfI-MluI
ORF Nucleotide Sequence:	>RC401457 representing NM_000179 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGTGCGGACAGAGCACCTGTACAGCTTCTTCCCAAGTCTCCGGCGCTGAGTGATGCCAACAAGGCCT
CGGCCAGGGCCTCACGCGAAGGCGCCGTGCCGCGCTGCCCGGGGCCTCTCCTTCCCGAGGCGGGGA
TGCGGCCTGGAGCGAGGCTGGGCCTGGGCCAGGCCCTTGGCGCGCTCCGCGTCACCGCCCAAGGCGAAG
AACCTCAACGGAGGGCTGCGGAGATCGGTAGCGCCTGCTGCCCCACCAGTTGTGACTTCTACCAGGAG
ATTTGGTTTGGCCAAGATGGAGGGTTACCCTGGTGGCCTTGTCTGGTTTACAACCACCCCTTTGATGG
AACATTATCCGCGAGAAAGGAAATCAGTCCGTGTTTCATGTACAGTTTTTTGATGACAGCCCAACAAGG
GGCTGGGTTAGCAAAAGGCTTTAAAGCCATATACAGGTTCAAATCAAAGGAAGCCAGAAGGGAGGTC



ATTTTACAGTGCAAAGCCTGAAATACTGAGAGCAATGCAACGTGCAGATGAAGCCTTAAATAAAGACAA
 GATTAAGAGGCTTGAATTGGCAGTTTGTGATGAGCCCTCAGAGCCAGAAGAGGAAGAAGAGATGGAGGTA
 GGCACAACCTACGTAACAGATAAGAGTGAAGAAGATAATGAAATTGAGAGTGAAGAGGAAGTACAGCCTA
 AGACACAAGGATCTAGGCGAAGTAGCCGCCAAATAAAAAACGAAGGGTCATATCAGATTCTGAGAGTGA
 CATTGGTGGCTCTGATGTGGAATTTAAGCCAGACACTAAGGAGGAAGGAAGCAGTGTGAAATAAGCAGT
 GGAGTGGGGGATAGTGAAGTGAAGGCCTGAACAGCCCTGTCAAAGTTGCTCGAAAGCGGAAGAGAATGG
 TGACTGGAAATGGCTCTCTTAAAAGGAAAAGCTCTAGGAAGGAAACGCCCTCAGCCACCAACAAGCAAC
 TAGCATTTCATCAGAAACCAAGAATACTTTGAGAGCTTTCTCTGCCCTCAAATTTCTGAATCCCAAGCC
 CACGTTAGTGGAGGTGGTGTGACAGTAGTCGCCCTACTGTTTGGTATCATGAACTTTAGAATGGCTTA
 AGGAGGAAAAGAGAAGAGATGAGCACAGGAGGAGGCTGATCACCCGATTTTGTGATCTACACTCTA
 TGTGCCTGAGGATTTCTCAATTCTTGTACTCTGGGATGAGGAAGTGGTGGCAGATTAAGTCTCAGAAC
 TTTGATCTTGTCTGTACAAGGTGGGAAAATTTATGAGCTGTACCACATGGATGCTCTATTGGAG
 TCAGTGAAGTGGGCTGGTATTCATGAAAGGCAACTGGGCCATTCTGGCTTCTGAAATTGCATTTGG
 CCGTTATTCAGATTCCTGGTGCAGAAGGGCTATAAAGTAGCACGAGTGAACAGACTGAGACTCCAGAA
 ATGATGGAGGCACGATGTAGAAAGATGGCACATATATCCAAGTATGATAGAGTGGTGGAGAGGAGATCT
 GTAGGATCATTACCAAGGTACACAGACTTACAGTGTGCTGGAAGGTGATCCCTCTGAGAACTACAGTAA
 GTATCTTCTTAGCCTCAAAGAAAAGAGGAAGATTCTTCTGGCCATACTCGTGCATATGGTGTGTGCTTT
 GTTGATACTTCACTGGGAAAGTTTTTCATAGGTCAGTTTTTTCAGATGATCGCCATTGTTTCGAGATTAGGA
 CTCTAGTGGCACACTATCCCCAGTACAAGTTTTATTTGAAAAAGGAAATCTCTCAAAGGAACTAAAAC
 AATTCTAAAGAGTTCATTGTCTGTCTCTTCCAGGAAGTCTGATACCCGGCTCCAGTTTTGGGATGCA
 TCCAAAACCTTTGAGAAGTCTCCTTGAGGAAGAATTTTAGGGAAAAGCTAAGTATGGCATTGGGGTGA
 TGTTACCCAGGTGCTTAAAGGATGACTTCAGAGTCTGATCCATTGGGTTGACACCAGGAGAGAAAAG
 TGAATTTGGCCCTCTCTGCTCTAGGTGGTGTGCTCTTCTACCTCAAAAAATGCCTTATTGATCAGGACTT
 TTATCAATGGCTAATTTTGAAGAATATATTCCTTGGATTCTGACACAGTCAGCACTACAAGATCTGGTG
 CTATCTTACCAGCAAGCTATCAACGAATGGTGTAGATGCAGTGCATTAAACAAGTGGAGATTTTTCT
 GAATGGAACAAATGGTTCTACTGAAGGAACCTACTAGAGAGGGTTGATACTTGCCTACTCCTTTTGGT
 AAGCGCTCCTAAAGCAATGGCTTTGTGCCCACTCTGTAACCATTATGCTATTAATGATCGTCTAGATG
 CCATAGAAGACCTCATGGTTGTGCTGACAAAATCTCCGAAGTTGTAGAGCTTCTAAAGAAGCTTCCAGA
 TCTTGAGAGGCTACTCAGTAAAATTCATAATGTTGGGTCTCCCTGAAGAGTCAAGAACCCAGACAGC
 AGGGCTATAATGTATGAAGAACTACATACAGCAAGAAGAAGATTATTGATTTTCTTCTGCTCTGGAAG
 GATTCAAAGTAATGTGTAATATAGGGATCATGGAAGAAGTGTGCTGATGGTTTTAAGTCTAAAATCCT
 TAAGCAGGTCTCTCTGACAGCAAAAATCCTGAAGTCTGTTTTCTGATTTGACTGTAGAATTGAAC
 CGATGGGATACAGCCTTTGACCATGAAAAGGCTCGAAAGACTGGACTTATTACTCCCAAAGCAGGCTTTG
 ACTCTGATTATGACCAAGCTCTTGTGACATAAGAGAAAATGAACAGAGCCTCCTGGAATACCTAGAGAA
 ACAGCGCAACAGAATTGGCTGTAGGACCATAGTCTATTGGGGATTGGTAGGAACCGTTACCAGCTGGAA
 ATTCTGAGAATTTACCAGTCAATTTGCCAGAAGAATACGAGTTGAAATCTACCAAGAAGGGCTGTA
 AACGATACTGGACAAAACCTATTGAAAAGAAGTTGGCTAATCTCATAAATGCTGAAGAACGGAGGGATGT
 ATCATTGAAGGACTGCATGCGGCGACTGTTCTAATACTTTGATAAAAAATTACAAGGACTGGCAGTCTGCT
 GTAGAGTGTATCGCAGTGTGGATGTTTTACTGTGCTGGCTAACTATAGTCGAGGGGTGATGGTCTTA
 TGTGTCGCCAGTAATTCCTGTTGCCGGAAGATACCCCCCTTCTTAGAGCTTAAAGGATCACGCCATCC
 TTGCATTACGAAGACTTTTTTTGGAGATGATTTTATTCCTAATGACATTCTAATAGGCTGTGAGGAAGG
 GAGCAGGAAAATGGCAAAGCCTATTGTGTGCTTGTACTGGACCAAAATAGGGGGCAAGTCTACGCTTA
 TGAGACAGGCTGGCTTATTAGCTGTAATGGCCAGATGGGTTGTACGTCCTGCTGTAGTGTGCAGGCT
 CACACCAATTGATAGAGTGTACTAGACTTGGTGCCTCAGACAGAATAATGTCAGGTGAAAGTACATTT
 TTTGTTGAATTAAGTGAAGTGCAGCATACTCATGCATGCAACAGCACATTCTCTGGTCTGTGGATG
 AATTAGGAAGAGTACTGCAACATTTGATGGGACGGCAATAGCAAATGCAGTTGTTAAAGAACTGTCTGA
 GACTATAAAATGTCGTACATTATTTCAACTCACTACCATTATTAGTAGAAGATTATTCTCAAATGTT
 GCTGTGCGCCTAGGACATATGGCATGCATGGTAGAAAAATGAATGTGAAGACCCAGCCAGGAGACTATTA
 CGTTCTCTATAAATTCATTAAGGGAGCTTGTCTAAAAGCTATGGCTTTAATGCAGCAAGGCTTGTCTAA
 TCTCCCAGAGGAAGTTATTCAAAGGGACATAGAAAAGCAAGAGAATTTGAGAAGATGAATCAGTCACTA
 CGATTATTTCCGGGAAGTTTGCCTGGCTAGTGAAGGTCAACTGTAGATGCTGAAGCTGTCCATAAATTGC
 TGACTTTGATTAAGGAATTA

AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC
TGGATTACAAGGATGACGACGA TAAGGTTTAA

Protein Sequence:

>RC401457 representing NM_000179
Red=Cloning site Green=Tags(s)

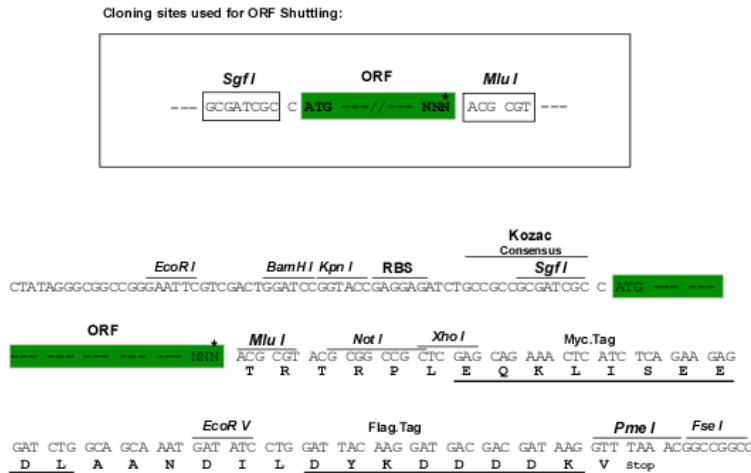
MSRQSTLYSFFPKSPALSDANKASARASREGGRAAAAPGASPSPGDDAAWSEAGPGRPLARSASPPKAK
NLNGGLRRSVAPAAPTSCDFSPGDLVWAKMEGYPWWPCLVYNHPFDGTFIREKGKSVRVHVQFFDDSPTR
GWVSKRLLKPYTGSKSKEAQKGGHFYSAPKEILRAMQRADEALNKDKIKRLELAVCDPESEPEEEEEMEV
GTTYVTDKSEEDNEIESEEEVQPKTQGSRRSSRIKRRVISEDSEDIIGGSDVEFKPDTKEEGSSDEISS
GVGDSESEGLNSPVKVARKRKRMVTGNGLKRRSSRKETPSATKQATSISSETKNTLRAF SAPQNSSEQA
HVSGGGDDSSRPTVWYHETLEWLKEEKRRDEHRRRDPDFDASTLYVPEDFLNSCTPGMRKWWQIKSQN
FDLVICYKVGKFYELYHMDALIGVSELGLVFMKGNWAHSGFPEIAFGYSDSLVQKGYKVARVEQTETPE
MMEARCRKMAHISKYDRVVRREICRIITKGTQTYSVLEGDPSENYSKYLKSLKEKEEDSSGHTRAYGVCF
VDTSLGKFFIGQFSDDRHCSTRFRTLVAHYPPVQVLFKGNLSKETKTILKSSLSLSCSLQEGLIPGSQFWD
SKTLRLLLEEEYFREKLSDGIGVMLPQVLKGMTSESDSIGLTPGEKSELALSALGGCVFYLLKCLIDQEL
LSMANFEEYIPLDSDTVSTTRSGAIFTKAYQRMVLDATLNNLEIFLNGTNGSTEGTLLELVDTCHTPF
KRLKQWLCAPLCNHYAINDRLDAIEDLMVVPDKISEVVELLKKLPDLERLLSKIHNVSPLKSNHPDS
RAIMYEETTSKKKIIDFLSALEGFKVMCKIIGIMEEVADGFKSKILKQVISLQTKNPEGRFPDLTVELN
RWDTAFDHEKARKTGLITPKAGFSDYDQALADIRENEQSLLYLEKQRNRIGCRTIVYWGIGRNRVQLE
IPENFTTRNLPEEYELKSTKKGCKRYWTKTIEKLANLANAEERRDVSLKDCMRRLFYNFDKNYKDWQSA
VECIAVLVLLCLANYSRGGDPMCRPVILLPEDTPPFLELKGSRHPCITKTFFGDDFIPNDILIGCEEE
EQENGKAYCVLVTGPNMGKSTLMRQAGLLAVMAQMGCYVPAVVCRLTPIDRVFTRLGASDRIMSGESTF
FVELSETASILMHATAHSLVLVDELGRGTATFDGTAIANAVVKELAEIKCRTLFSTHYHSLVEDYSQNV
AVRLGHMACMVENECEDPSQETITFLYKFKGACPKSYGFNAARLANLPEEVIQKGRHKAREFEKMNQSL
RLFREVLASERSTVDAEAVHKLLTLIKEL

SGPTRRRLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites:

Sgfl-MluI

Cloning Scheme:



* The last codon before the Stop codon of the ORF

OTI Disclaimer:	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 custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.
	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).
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NP_000170
RefSeq Size:	4080 bp
RefSeq ORF:	4083 bp
Locus ID:	2956
Cytogenetics:	2p16.3
Domains:	PWWP, MutS_V, MutS_I, MutS_III, MutS_II, MutS_IV
Protein Families:	Druggable Genome, Stem cell - Pluripotency
Protein Pathways:	Colorectal cancer, Mismatch repair, Pathways in cancer
MW:	149.6 kDa
Gene Summary:	This gene encodes a member of the DNA mismatch repair MutS family. In E. coli, the MutS protein helps in the recognition of mismatched nucleotides prior to their repair. A highly conserved region of approximately 150 aa, called the Walker-A adenine nucleotide binding motif, exists in MutS homologs. The encoded protein heterodimerizes with MSH2 to form a mismatch recognition complex that functions as a bidirectional molecular switch that exchanges ADP and ATP as DNA mismatches are bound and dissociated. Mutations in this gene may be associated with hereditary nonpolyposis colon cancer, colorectal cancer, and endometrial cancer. Transcripts variants encoding different isoforms have been described. [provided by RefSeq, Jul 2013]