

Product datasheet for RC401522

MLH1 (NM_000249) Human Mutant ORF Clone

Product data:

OriGene Technologies, Inc.

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| Product Type: | Mutant ORF Clones |
|------------------------------|---|
| Product Name: | MLH1 (NM_000249) Human Mutant ORF Clone |
| Mutation Description: | G122X |
| Affected Codon#: | 122 |
| Affected NT#: | 364 |
| Nucleotide Mutation: | MLH1 Mutant (G122X), Myc-DDK-tagged ORF clone of Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli) (MLH1), transcript variant 1 as transfection-ready DNA |
| Effect: | Colorel ner, non-polyposis |
| Symbol: | MLH1 |
| Synonyms: | COCA2; FCC2; hMLH1; HNPCC; HNPCC2; MMRCS1 |
| E. coli Selection: | Kanamycin (25 ug/mL) |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-Entry (PS100001) |
| Tag: | Myc-DDK |
| ACCN: | NM_000249 |
| ORF Size: | 363 bp |
| Restriction Sites: | Sgfl-Mlul |



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| | 11 (NM_000249) Human Mutant ORF Clone – RC401522 |
|-----------------------------|---|
| ORF Nucleotide Sequence: | <pre>>RC401522 representing NM_000249 Red=Cloning site Blue=ORF Green=Tags(s)</pre> |
| | TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC GCC <mark>GCGATCGC</mark> C |
| | ATGTCGTTCGTGGCAGGGGTTATTCGGCGGCTGGACGAGACAGTGGTGAACCGCATCGCGGCGGGGGAAG TTATCCAGCGGCCAGCTAATGCTATCAAAGAGATGATTGAGAACTGTTTAGATGCAAAATCCACAAGTAT TCAAGTGATTGTTAAAGAGGGAGGCCTGAAGTTGATTCAGATCCAAGACAATGGCACCGGGATCAGGAAA GAAGATCTGGATATTGTATGTGAAAGGTTCACTACTAGTAAACTGCAGTCCTTTGAGGATTTAGCCAGTA TTTCTACCTATGGCTTTCGAGGTGAGGCTTTGGCCAGCATAAGCCATGTGGCTCATGTTACTATTACAAC GAAAACAGCTGAT |
| | AGCGGACCGACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC TGGATTACAAGGATGACGACGA TAAG GTTTAA |
| Protein Sequence: | <pre>>RC401522 representing NM_000249 Red=Cloning site Green=Tags(s)</pre> |
| | MSFVAGVIRRLDETVVNRIAAGEVIQRPANAIKEMIENCLDAKSTSIQVIVKEGGLKLIQIQDNGTGIRK EDLDIVCERFTTSKLQSFEDLASISTYGFRGEALASISHVAHVTITTKTAD |
| | SGPTRTRRLEQKLISEEDLAANDILDYKDDDDKV |
| Restriction Sites: | Sgfl-Mlul |
| Cloning Scheme: | Cloning sites used for ORF Shuttling: Sgf1 ORF Miu I GCGATCGC C ATG YIRN ACG CGT |
| | Kozac Consensus EcoR1 BamH1 Kpn 1 RBS Sgf1 CTATAGGGGGGGGGAATTCGTCCGATCGGGTACCGGGAGAGATCTGCCGCCGCGGTACGC ATC |
| | ORF <u>Miui Noti Xhoi</u> Myc.Tag ACG CGT ACG CGG CCG CTC GAG CAG AAA CTC ATC TCA GAA GAG T R T R P L <u>E Q K L I S E E</u> |
| | EcoR V Flag.Tag Pme I Fse I GAT CTG GCA GCA AAT GAT ATC CTG GAT TAC AAG GAT GAC GAC GAT AAG GTT TAA ACGGCCGGCC D L A N D I L D Y K D D K V stop |
| | * The last codon before the Stop codon of the ORF |
| 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. <u>More info</u> |
| OTI Annotation: | This clone was angineered to express the complete OPE with an expression tag. Expression |

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

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| Service MLH1 (NM_000249) Human Mutant ORF Clone – RC401522 | |
|--|--|
| 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: | <u>NP 000240</u> |
| RefSeq Size: | 363 bp |
| RefSeq ORF: | 2271 bp |
| Locus ID: | 4292 |
| Cytogenetics: | 3p22.2 |
| Domains: | DNA_mis_repair, HATPase_c |
| Protein Families: | Druggable Genome |
| Protein Pathways | Colorectal cancer, Endometrial cancer, Mismatch repair, Pathways in cancer |
| MW: | 13.3 kDa |
| Gene Summary: | The protein encoded by this gene can heterodimerize with mismatch repair endonuclease PMS2 to form MutL alpha, part of the DNA mismatch repair system. When MutL alpha is bound by MutS beta and some accessory proteins, the PMS2 subunit of MutL alpha introduces a single-strand break near DNA mismatches, providing an entry point for exonuclease degradation. The encoded protein is also involved in DNA damage signaling and can heterodimerize with DNA mismatch repair protein MLH3 to form MutL gamma, which is involved in meiosis. This gene was identified as a locus frequently mutated in hereditary nonpolyposis colon cancer (HNPCC). [provided by RefSeq, Aug 2017] |

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