

Product datasheet for SC328850

MLH1 (NM_001167618) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	MLH1 (NM_001167618) Human Untagged Clone
Tag:	Tag Free
Symbol:	MLH1
Synonyms:	COCA2; FCC2; hMLH1; HNPCC; HNPCC2; MMRCS1
Mammalian Cell Selection:	None
Vector:	pCMV6-XL5
E. coli Selection:	Ampicillin (100 ug/mL)

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn



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Fully Sequenced ORF:	<pre>>NCBI ORF sequence for NM_001167618, the custom clone sequence may differ by one or more nucleotides ATGAATGGTTACATATCCAATGCAAACTACTCAGTGAAGAAGTGCATCTTCTTACTCTTC ATCAACCATCGTCTGGTAGAATCAACTTCCTTGAGAAAAGCCATAGAAACAGTGTATGCA GCCTATTTGCCCAAAAACACACCCCATTCCTTGCTGACATCAGTTTAGAAATCAGTGCCCAG AATGTGGATGTTAATGTGCACCCCCACAAGCATGAAGTTCACTTCCTGCACGAGGAGAGC ATGTGGATGTTAATGTGCACCCCCACAAGCATGAAGTTCACTTCCTGGCCCCCAGGAGAGGC ATGTACTTCACCCAGACCATCGAGAGCAAGCTCGTGGGCCCCATTCCTCCCAGG ATGTACTTCACCCAGACCTCGAGAGCAAGCTCGTGGGCCCCCTCTGGGGAGATGGTTAAA TCCACAACAGTCTGCACGCCGGGACAGGACGTGCTGGGAGATGGTTAAA TCCACAACAGTCTGCACGCCGGGGGAACAGAAGCTTGGAGTAGTGATAAGGTCTATGCCAC CAGATGGTCGCAGCAGCACGCCCAGGCCATTGTCACAGAGGAGTAGGCATAGTCGCAGCCTGGAGAGGCT GGCCAGGCTAGGCAGCCAAGGCCATTGTCACAGAGGGATAGGCATTGCTGCAGCCTGGAAGTGGCT GCCAAAAATCAGAGCTTGGAGGGGGATACAACAAAGGGGACTTCGGAAGATGTCGAGGCTGGAGAATG GGGAGGCTAGCTTGCCAGGAAGGAAGTGCTTGAACTCCCAGGCGAAAGGACATG GGGAGGATGATTCCCGAAAGGAAATGCCTGGAGCTTGGACATGGGCCTTGCC CGGAAGATGATTCCCGAAAGGAAATGCCGGCGGGGAATTCCTCAGGGAGAATG GTGGAAGATGGTTTCCCAGGAAGGAATGCTGGGACGTTATGGAGAAGGACTGTTGCA CAGCATCATAACCACCCCTTCGTGGGCCTGTGTGAATCCTCCAGGGACATGAGGTTCTC CGGGAGATGTTGCCATAACCACCCCTTGGTGGACTTGGAGAGAGCTGGGCCACGCA CAGCATCATAACCACACCTCTTGCTGAGGCGTGTGTAATCCTCAGGAGGACAGAGGACGA CAGCATCAAACCAAGTTATACCTTCCAAGAAGGAAGGTGGCTGGTGACAAGGAGGATG CTTGCAAAGAAGGACTTGCTGAATACATTGTGAGGTGTCTCCAGGAAGGA</pre>
Restriction Sites:	Please inquire
ACCN:	NM_001167618
OTI Disclaimer:	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).
OTI Annotation:	This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.
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).

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DRIGENE MLH1 (NM_001167618) Human Untagged Clone – SC328850

Reconstitution Method:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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:	<u>NM 001167618.1, NP 001161090.1</u>
RefSeq Size:	2473 bp
RefSeq ORF:	1548 bp
Locus ID:	4292
UniProt ID:	<u>P40692</u>
Cytogenetics:	3p22.2
Protein Families:	Druggable Genome
Protein Pathways:	Colorectal cancer, Endometrial cancer, Mismatch repair, Pathways in cancer
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] Transcript Variant: This variant (3) contains a distinct 5'-terminal exon and uses an alternate splice site in a 5' exon and therefore lacks an in-frame portion of the 5' coding region compared to variant 1. The resulting isoform (3) has a shorter N-terminus compared to

isoform 1. Variants 3, 4, and 6-12 all encode the same isoform (3). Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.

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