

Product datasheet for SC208847

MCM2 (NM_004526) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: MCM2 (NM_004526) Human 3' UTR Clone

Symbol: MCM2

Synonyms: BM28; CCNL1; cdc19; CDCL1; D3S3194; DFNA70; MITOTIN

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_004526

Insert Size: 693 bp

Insert Sequence: >SC208847 3'UTR clone of NM_004526

The sequence shown below is from the reference sequence of NM_004526. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).



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Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 004526.4</u>

Summary: The protein encoded by this gene is one of the highly conserved mini-chromosome

maintenance proteins (MCM) that are involved in the initiation of eukaryotic genome

replication. The hexameric protein complex formed by MCM proteins is a key component of the pre-replication complex (pre_RC) and may be involved in the formation of replication forks and in the recruitment of other DNA replication related proteins. This protein forms a complex with MCM4, 6, and 7, and has been shown to regulate the helicase activity of the complex. This protein is phosphorylated, and thus regulated by, protein kinases CDC2 and CDC7. Multiple alternatively spliced transcript variants have been found, but the full-length

nature of some variants has not been defined. [provided by RefSeq, Oct 2012]

Locus ID: 4171

MW: 25.1