

Product datasheet for **SC207618**

SRRM2 (NM_016333) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	SRRM2
Synonyms:	300-KD; Cwc2l; CWF2l; HSPC075; SRL300; SRm300
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PSI00062)
ACCN:	NM_016333
Insert Size:	575 bp
Insert Sequence:	>SC207618 3'UTR clone of NM_016333 The sequence shown below is from the reference sequence of NM_016333. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC ATGAGACACCGCTCCTCCAGGTCTCCAATAATTGTCTTTGGGGGATTCCACCACACCCAATGCTCTGGA GCCACAAGGAGTGTCCCTTCTCCCCAGCAGAGCCGTGGGAGGGTCCTTGTCTGCTCTCCTTTGAACCT TGGCAGCCCTTGGATGGAGGGCTCCCTTTCCCTCCCTTTTTTTTTTTCTTTGTTCTGTGAAATGTTAA TCTCCGTGAGTTCTTCTGTTTCATGTGTTCTGGGGGGTTTGGGGTGGGAGGGAATGCAGATGGGAGTT GGGGGAGGGGAGGATACAGTTCAAGATACCCAGCCTGGAGTCAGGGCCAGGGAGGCATGGCCCCACTT GTATCCAGAAGTTCCAGGGGTGATTGTGATGGTGGTGGGACTGGAGTTGTATAAGGTGTTCTTGGA AGGAAGGGGCAGGAGTTGGAATTAGTTGGTCCCTACTGTCCCCCATGAGGTTGTGAACCCCTCCCCCA ACTTTTCATGTTTCTTAAAGGCATTTTGGTTTTTAAATCTGTACAGCAAGAGCAACTTTTCTGTCA AATAAAATGAGAAATGCAGGAA ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG </pre>
Restriction Sites:	SgfI-MluI
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).



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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_016333.4
Summary:	Required for pre-mRNA splicing as component of the spliceosome.[UniProtKB/Swiss-Prot Function]
Locus ID:	23524
MW:	20.8