

Product datasheet for **SC205369**

SFRS15 (SCAF4) (NM_020706) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	SFRS15 (SCAF4) (NM_020706) Human 3' UTR Clone
Symbol:	SFRS15
Synonyms:	SFRS15; SRA4
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_020706
Insert Size:	429 bp
Insert Sequence:	<p>>SC205369 3'UTR clone of NM_020706</p> <p>The sequence shown below is from the reference sequence of NM_020706. The complete sequence of this clone may contain minor differences, such as SNPs.</p> <p>Blue=Stop Codon Red=Cloning site</p>

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GGCAAGTTGGACGCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAACGATCGC
TCTGGCTCAGCAGCAGAGGCTCCTCGTAGAGACTGGAATTTGTGAAAATGTGACAGTGACACTTCCTG
GAGTGTAGAGCTTGAGGTGTACAGATGCTGTATTATATCCGCTCCCGCTGTACTGCAGCCCCGCCAG
CTGGTGGGGAAGTGAAGCAATTTGATTGCTTCCCTTCTATTTAAAAATAGCCACAAAATAACAAAAA
TACTGAAAATATGAATAAATATTACCCTTTTTGCTGTAACTTTTTAAAGTTTTGACTTTAAAAAGTTT
ACAAATCGTAATTAGAAGTGCTCTATTTTTTTTTTTTTTTTAAATTTAAGACAAGGTAAACGGTAAA
GCTCCTCAAAACAATAGGGATGTTTTTAATAAACTCTATTTTCGTAACAACTTTAATGTGTGCTATTC
TTCCTACACTGCATT
ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
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Restriction Sites:	Sgfl-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.


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RefSeq: [NM_020706.2](#)

Summary: This gene likely encodes a member of the arginine/serine-rich splicing factor family. A similar protein in Rat appears to bind the large subunit of RNA polymerase II and provide a link between transcription and pre-mRNA splicing. Alternatively spliced transcript variants have been described. [provided by RefSeq, Feb 2009]

Locus ID: 57466

MW: 17.1