

Product datasheet for **SC205368**

SFRS15 (SCAF4) (NM_001145445) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: SFRS15

Synonyms: SFRS15; SRA4

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PSI00062)

ACCN: NM_001145445

Insert Size: 429 bp

Insert Sequence: >SC205368 3'UTR clone of NM_001145445
The sequence shown below is from the reference sequence of NM_001145445. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
TCTGGCTCAGCAGCAGAGGCTCCTCGTAGAGACTGGAATTTGTGAAATGTGACAGTGACACTTCCTG
GAGTGTAGAGCTTGAGGTGTACAGATGCTGTATTATATCCGCTCCCGCTGACTGCAGCCCCGCGCCAG
CTGGTGGGGAAGTGAAGCAATTTGATTGCTTCCCTTCTATTTAAAAATAGCCACAAATAACAAAAA
TACTGAAAATATGAATAAATATTACCTTTTTGCTGTAACTTTTAAAAAGTTTGAAGTTTAAAAAGTTT
ACAAATCGTAATTAGAAGTGCTCTCTATTTTTTTTTTTTTTTTAAATTTAAGACAAGGTAACGGTGAAA
GCTCCTCAAAACAATAGGGATGTTTTTAATAAACTCTATTTTCGTAACAACTTTAATGTGTGCTATTC
TTCCTACACTGCATT
ACGCGTAAGCGGCCGCGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG
```

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_001145445.1
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