

Product datasheet for **SC201636**

SAP155 (SF3B1) (NM_001005526) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	SAP155 (SF3B1) (NM_001005526) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	SF3B1
Synonyms:	Hsh155; MDS; PRP10; PRPF10; SAP155; SF3b155
ACCN:	NM_001005526
Insert Size:	166 bp
Insert Sequence:	>SC201636 3'UTR clone of NM_001005526 The sequence shown below is from the reference sequence of NM_001005526. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC TTTGCAGATGGCTTCTATTCTGCTGCTTGAAGTCAGAACTGCTGATGGAGACAAAGGCACGAAAGTGTA CGTATTCCGATTAGCAACCCAGGAACCCATCACTTCTGAAGACTCTAACTGTGCTGTCAATTTGTTT TTATATGCATTAATAATTTGTTTTAAA ACGCGT AAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
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).
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_001005526.2</u>



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Summary:

This gene encodes subunit 1 of the splicing factor 3b protein complex. Splicing factor 3b, together with splicing factor 3a and a 12S RNA unit, forms the U2 small nuclear ribonucleoproteins complex (U2 snRNP). The splicing factor 3b/3a complex binds pre-mRNA upstream of the intron's branch site in a sequence independent manner and may anchor the U2 snRNP to the pre-mRNA. Splicing factor 3b is also a component of the minor U12-type spliceosome. The carboxy-terminal two-thirds of subunit 1 have 22 non-identical, tandem HEAT repeats that form rod-like, helical structures. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]

Locus ID:

23451

MW:

6.3