

Product datasheet for SC201821

SF3B2 (NM 006842) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: SF3B2 (NM_006842) Human 3' UTR Clone

Symbol: SF3B2

Synonyms: Cus1; SAP145; SF3b1; SF3B145; SF3b150

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_006842

Insert Size: 593 bp

Insert Sequence: >SC201821 3'UTR clone of NM_006842

The sequence shown below is from the reference sequence of NM_006842. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TGTGCGACAACATAGCTCAAAAGCTAAATTGTCTTGAGTCC

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).



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



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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 006842.3</u>

Summary: This gene encodes subunit 2 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. Subunit 2 associates with pre-mRNA upstream of the branch site at the anchoring site. Subunit 2 also interacts directly with subunit 4 of the splicing factor 3b complex. Subunit 2 is a highly hydrophilic protein with a proline-rich N-terminus and a

glutamate-rich stretch in the C-terminus. [provided by RefSeq, Jul 2008]

Locus ID: 10992 MW: 22.3