

Product datasheet for RC205846L4V

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

splicing factor 1 (SF1) (NM 201998) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: splicing factor 1 (SF1) (NM_201998) Human Tagged ORF Clone Lentiviral Particle

Symbol: splicing factor 1

Synonyms: BBP; D11S636; MBBP; ZCCHC25; ZFM1; ZNF162

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_201998 **ORF Size:** 1644 bp

ORF Nucleotide

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Sequence:

The ORF insert of this clone is exactly the same as(RC205846).

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This

clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeq: <u>NM 201998.1</u>

RefSeq Size: 2949 bp
RefSeq ORF: 1647 bp
Locus ID: 7536

UniProt ID: Q15637
Cytogenetics: 11q13.1

Protein Families: Transcription Factors

MW: 59.7 kDa





Gene Summary:

This gene encodes a nuclear pre-mRNA splicing factor. The encoded protein specifically recognizes the intron branch point sequence at the 3' splice site, together with the large subunit of U2 auxiliary factor (U2AF), and is required for the early stages of spliceosome assembly. It also plays a role in nuclear pre-mRNA retention and transcriptional repression. The encoded protein contains an N-terminal U2AF ligand motif, a central hnRNP K homology motif and quaking 2 region which bind a key branch-site adenosine within the branch point sequence, a zinc knuckles domain, and a C-terminal proline-rich domain. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Oct 2016]