

Product datasheet for **SC309011**

Spt6 (SUPT6H) (NM_003170) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Spt6 (SUPT6H) (NM_003170) Human Untagged Clone
Tag:	Tag Free
Symbol:	Spt6
Synonyms:	emb-5; SPT6; SPT6H
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL6</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>NCBI ORF sequence for NM_003170, the custom clone sequence may differ by one or more nucleotides

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ATGTCTGATTTTGTGAAAGCGAGGCTGAGGAGTCAGAGGAAGAATACAATGATGAAGGC
GAGGTGGTACCCCGAGTCACCAAGAAATTTGTGGAAGAGGAGGATGATGATGAGGAGGAG
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GAAGGTGACGAGGCAGAAGATGAGGAGCAGAGGGGCCTGAGCTCAAGCAAGCCTCTCGC
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 TCCTTTGCCCGGACCTTCTGAATCACAAGTATTATCAGGACTGCAGCGGTGGGACCCG
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 TGGCTGCAGGAAAAGGAGGCAGAACGGCGGAAACAGAAGCAGCGGCTGACACCTCGGCC
 TCCCCAGCCCCATGATCGAAAGCACCCCATGTCCATTGCTGGCGATGCCACCCCACTC
 CTGGACGAGATGGATCGGTAG

Restriction Sites: Please inquire

ACCN: NM_003170

OTI Disclaimer: Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.

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 TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_003170.3](#), [NP_003161.2](#)

RefSeq Size: 5896 bp

RefSeq ORF: 5181 bp

Locus ID: 6830

UniProt ID: [Q7KZ85](#)

Cytogenetics: 17q11.2

Domains: SH2, S1, YqgFc

Protein Families: Transcription Factors

Gene Summary: Transcription elongation factor which binds histone H3 and plays a key role in the regulation of transcription elongation and mRNA processing. Enhances the transcription elongation by RNA polymerase II (RNAPII) and is also required for the efficient activation of transcriptional elongation by the HIV-1 nuclear transcriptional activator, Tat. Besides chaperoning histones in transcription, acts to transport and splice mRNA by forming a complex with IWS1 and the C-terminal domain (CTD) of the RNAPII subunit RPB1 (POLR2A). The SUPT6H:IWS1:CTD complex recruits mRNA export factors (ALYREF/THOC4, EXOSC10) as well as histone modifying enzymes (such as SETD2), to ensure proper mRNA splicing, efficient mRNA export and elongation-coupled H3K36 methylation, a signature chromatin mark of active transcription. SUPT6H via its association with SETD1A, regulates both class-switch recombination and somatic hypermutation through formation of H3K4me3 epigenetic marks on activation-induced cytidine deaminase (AICDA) target loci. Promotes the activation of the myogenic gene program by entailing erasure of the repressive H3K27me3 epigenetic mark through stabilization of the chromatin interaction of the H3K27 demethylase KDM6A.
[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (2) differs in the 5' UTR compared to variant 1. Both variants 1 and 2 encode the same protein.