

## Product datasheet for **TR309007**

### SUPT16H Human shRNA Plasmid Kit (Locus ID 11198)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	SUPT16H Human shRNA Plasmid Kit (Locus ID 11198)
Locus ID:	11198
Synonyms:	CDC68; FACTP140; SPT16; SPT16/CDC68
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	SUPT16H - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 11198). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<a href="#">NM_007192</a> , <a href="#">NM_007192.1</a> , <a href="#">NM_007192.2</a> , <a href="#">NM_007192.3</a> , <a href="#">BC000565</a> , <a href="#">BC012582</a> , <a href="#">BC014046</a> , <a href="#">BC064561</a> , <a href="#">BC073849</a> , <a href="#">BC111402</a>
UniProt ID:	<a href="#">Q9Y5B9</a>
Summary:	Transcription of protein-coding genes can be reconstituted on naked DNA with only the general transcription factors and RNA polymerase II. However, this minimal system cannot transcribe DNA packaged into chromatin, indicating that accessory factors may facilitate access to DNA. One such factor, FACT (facilitates chromatin transcription), interacts specifically with histones H2A/H2B to effect nucleosome disassembly and transcription elongation. FACT is composed of an 80 kDa subunit and a 140 kDa subunit; this gene encodes the 140 kDa subunit. [provided by RefSeq, Feb 2009]
shRNA Design:	These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact <a href="mailto:techsupport@origene.com">techsupport@origene.com</a> . If you need a special design or shRNA sequence, please utilize our <a href="#">custom shRNA service</a> .



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**Performance  
Guaranteed:**

OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at [techsupport@origene.com](mailto:techsupport@origene.com). Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).