

## Product datasheet for **TR309560**

### Semaphorin 3c (SEMA3C) Human shRNA Plasmid Kit (Locus ID 10512)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Semaphorin 3c (SEMA3C) Human shRNA Plasmid Kit (Locus ID 10512)
Locus ID:	10512
Synonyms:	SEMAE; SemE
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	SEMA3C - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 10512). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<a href="#">NM_006379</a> , <a href="#">NM_001350120</a> , <a href="#">NM_001350121</a> , <a href="#">NM_006379.1</a> , <a href="#">NM_006379.2</a> , <a href="#">NM_006379.3</a> , <a href="#">BC030690</a> , <a href="#">BC030690.1</a> , <a href="#">BM473251</a> , <a href="#">NM_006379.5</a>
UniProt ID:	<a href="#">Q99985</a>
Summary:	This gene encodes a secreted glycoprotein that belongs to the semaphorin class 3 family of neuronal guidance cues. The encoded protein contains an N-terminal sema domain, integrin and immunoglobulin-like domains, and a C-terminal basic domain. Homodimerization and proteolytic cleavage of the C-terminal propeptide are necessary for the function of the encoded protein. It binds a neuropilin co-receptor before forming a heterotrimeric complex with an associated plexin. An increase in the expression of this gene correlates with an increase in cancer cell invasion and adhesion. Naturally occurring mutations in this gene are associated with Hirschsprung disease. [provided by RefSeq, May 2017]
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).