

Product datasheet for **TR301309**

CSPS (SULT1A3) Human shRNA Plasmid Kit (Locus ID 6818)

Product data:

Product Type:	shRNA Plasmids
Product Name:	CSPS (SULT1A3) Human shRNA Plasmid Kit (Locus ID 6818)
Locus ID:	6818
Synonyms:	STM, HAST, HAST3, M-PST, ST1A5, TL-PST, SULT1A4, MGC117469
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	SULT1A3 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 6818). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	NM_001017387 , NM_003166 , NM_177552 , NM_003166.1 , NM_003166.2 , NM_003166.3 , NM_177552.1 , NM_177552.2 , NM_177552.3 , NM_001017387.1 , BC078144 , BC078144.1 , BC132914 , BC132920 , BC141504 , BC144354 , BC144355 , BC156739 , BC171772 , NM_177552.4
UniProt ID:	P50224
Summary:	Sulfotransferase enzymes catalyze the sulfate conjugation of many hormones, neurotransmitters, drugs, and xenobiotic compounds. These cytosolic enzymes are different in their tissue distributions and substrate specificities. The gene structure (number and length of exons) is similar among family members. This gene encodes a phenol sulfotransferase with thermolabile enzyme activity. Four sulfotransferase genes are located on the p arm of chromosome 16; this gene and SULT1A4 arose from a segmental duplication. This gene is the most centromeric of the four sulfotransferase genes. Read-through transcription exists between this gene and the upstream SLX1A (SLX1 structure-specific endonuclease subunit homolog A) gene that encodes a protein containing GIY-YIG domains. [provided by RefSeq, Nov 2010]
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 techsupport@origene.com . If you need a special design or shRNA sequence, please utilize our custom shRNA service .



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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. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).