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Product datasheet for TL519366V

Sat2 Mouse shRNA Lentiviral Particle (Locus ID 69215)

Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	Sat2 Mouse shRNA Lentiviral Particle (Locus ID 69215)
Locus ID:	69215
Synonyms:	2610016A03Rik; SSAT-2; SSAT2
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	Sat2 - Mouse shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, >10^7 TU/ml.
RefSeq:	BC061227, NM_026991, NM_026991.1, NM_026991.2, NM_001356468, NM_001356469, NM_026991.3
UniProt ID:	Q6P8J2
Summary:	Enzyme which catalyzes the acetylation of polyamines. Substrate specificity: norspermidine > spermidine = spermine >> N(1)acetylspermine = putrescine (By similarity).[UniProtKB/Swiss-Prot Function]
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).