

Product datasheet for TL509666

Siah1a Mouse shRNA Plasmid (Locus ID 20437)

Product data:

OriGene Technologies, Inc.

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Product Type:	shRNA Plasmids
Product Name:	Siah1a Mouse shRNA Plasmid (Locus ID 20437)
Locus ID:	20437
Synonyms:	AA982064; Al853500; Sinh1a
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	Siah1a - Mouse, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 20437). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<u>BC046317, NM 009172, NM 009172.1, NM 009172.2</u>
UniProt ID:	<u>P61092</u>



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GRIGENE Siah1a Mouse shRNA Plasmid (Locus ID 20437) – TL509666

Summary:	E3 ubiquitin-protein ligase that mediates ubiquitination and subsequent proteasomal
	degradation of target proteins. E3 ubiquitin ligases accept ubiquitin from an E2 ubiquitin-
	conjugating enzyme in the form of a thioester and then directly transfers the ubiquitin to
	targeted substrates. Mediates E3 ubiquitin ligase activity either through direct binding to
	substrates or by functioning as the essential RING domain subunit of larger E3 complexes.
	Triggers the ubiquitin-mediated degradation of many substrates, including proteins involved
	in transcription regulation (ELL2, MYB, POU2AF1, PML and RBBP8), a cell surface receptor
	(DCC), the cell-surface receptor-type tyrosine kinase FLT3, the cytoplasmic signal transduction
	molecules (KLF10/TIEG1 and NUMB), an antiapoptotic protein (BAG1), a microtubule motor
	protein (KIF22), a protein involved in synaptic vesicle function in neurons (SYP), a structural
	protein (CTNNB1) and SNCAIP. Confers constitutive instability to HIPK2 through proteasomal
	degradation. It is thereby involved in many cellular processes such as apoptosis, tumor
	suppression, cell cycle, axon guidance, transcription, spermatogenesis and TNF-alpha
	signaling. Has some overlapping function with SIAH2. Required for completion of meiosis I in
	males. Induces apoptosis in cooperation with PEG3. Upon nitric oxid (NO) generation that
	follows apoptotic stimulation, interacts with S-nitrosylated GAPDH, mediating the
	translocation of GAPDH to the nucleus. GAPDH acts as a stabilizer of SIAH1, facilitating the
	degradation of nuclear proteins.[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 <u>techsupport@origene.com</u> .
	If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u> .
Performance	OriGene guarantees that the sequences in the shRNA expression cassettes are verified to
Guaranteed:	correspond to the target gene with 100% identity. One of the four constructs at minimum are

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).

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