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Product datasheet for TR309207

NCOA62 (SNW1) Human shRNA Plasmid Kit (Locus ID 22938)

Product data:

Product Type:	shRNA Plasmids
Product Name:	NCOA62 (SNW1) Human shRNA Plasmid Kit (Locus ID 22938)
Locus ID:	22938
Synonyms:	Bx42; FUN20; NCOA-62; Prp45; PRPF45; SKIIP; SKIP; SKIP1
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	SNW1 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 22938). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<u>NM_001318844</u> , <u>NM_012245</u> , <u>NM_012245.1</u> , <u>NM_012245.2</u> , <u>BC032377</u> , <u>BC040112</u> , <u>BC046105</u> , <u>BC065286</u> , <u>BC105585</u> , <u>BC108902</u> , <u>BC108903</u> , <u>NM_012245.3</u>
UniProt ID:	<u>Q13573</u>
Summary:	This gene, a member of the SNW gene family, encodes a coactivator that enhances transcription from some Pol II promoters. This coactivator can bind to the ligand-binding domain of the vitamin D receptor and to retinoid receptors to enhance vitamin D-, retinoic acid-, estrogen-, and glucocorticoid-mediated gene expression. It can also function as a splicing factor by interacting with poly(A)-binding protein 2 to directly control the expression of muscle-specific genes at the transcriptional level. Finally, the protein may be involved in oncogenesis since it interacts with a region of SKI oncoproteins that is required for transforming activity. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2016]
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> .



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GRIGENE NCOA62 (SNW1) Human shRNA Plasmid Kit (Locus ID 22938) – TR309207

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).

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