

## **Product datasheet for TR300527**

## 9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436

OriGene Technologies, Inc.

Phone: +1-888-267-4436
https://www.origene.com
techsupport@origene.com
EU: info-de@origene.com
CN: techsupport@origene.cn

## **WBSCR22 Human shRNA Plasmid Kit (Locus ID 114049)**

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** WBSCR22 Human shRNA Plasmid Kit (Locus ID 114049)

**Locus ID:** 114049

Synonyms: HASJ4442; HUSSY-3; MERM1; PP3381; WBMT; WBSCR22

Vector: pRS (TR20003)

E. coli Selection: Ampicillin

Mammalian Cell Puromycin

Selection:

Format: Retroviral plasmids

Components: WBSCR22 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID

= 114049). 5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.

RefSeq: NM 001202560, NM 017528, NR 037776, NR 045512, NM 017528.1, NM 017528.2,

NM 017528.3, NM 017528.4, NM 001202560.1, NM 001202560.2, BC011696, BC011696.2,

BC000169, BC001780, BM671506, BM794274, NM 017528.5, NM 001202560.3

UniProt ID: 043709

Summary: This gene encodes a protein containing a nuclear localization signal and an S-adenosyl-L-

methionine binding motif typical of methyltransferases, suggesting that the encoded protein may act on DNA methylation. This gene is deleted in Williams syndrome, a multisystem developmental disorder caused by the deletion of contiguous genes at 7q11.23. Alternatively

spliced transcript variants have been found. [provided by RefSeq, Feb 2011]

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 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).