

## Product datasheet for **TL317131V**

### SETD2 Human shRNA Lentiviral Particle (Locus ID 29072)

#### Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	SETD2 Human shRNA Lentiviral Particle (Locus ID 29072)
Locus ID:	29072
Synonyms:	HIF1, FLJ16420, FLJ22472, FLJ45883, KIAA1732
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	HYPB - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, >10 <sup>7</sup> TU/ml.
RefSeq:	<a href="#">NM_012271</a> , <a href="#">NM_014159</a> , <a href="#">NM_001349370</a> , <a href="#">NR_146158</a> , <a href="#">NM_014159.1</a> , <a href="#">NM_014159.2</a> , <a href="#">NM_014159.3</a> , <a href="#">NM_014159.4</a> , <a href="#">NM_014159.6</a> , <a href="#">BC072440</a> , <a href="#">BC090954</a> , <a href="#">BC117162</a> , <a href="#">BC117164</a> , <a href="#">NM_014159.7</a>
UniProt ID:	<a href="#">Q9BYW2</a>
Summary:	Huntington's disease (HD), a neurodegenerative disorder characterized by loss of striatal neurons, is caused by an expansion of a polyglutamine tract in the HD protein huntingtin. This gene encodes a protein belonging to a class of huntingtin interacting proteins characterized by WW motifs. This protein is a histone methyltransferase that is specific for lysine-36 of histone H3, and methylation of this residue is associated with active chromatin. This protein also contains a novel transcriptional activation domain and has been found associated with hyperphosphorylated RNA polymerase II. [provided by RefSeq, Aug 2008]
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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a> . If you need a special design or shRNA sequence, please utilize our <a href="#">custom shRNA service</a> .



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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](mailto: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).