

## Product datasheet for **TL310456V**

### PHF8 Human shRNA Lentiviral Particle (Locus ID 23133)

#### Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	PHF8 Human shRNA Lentiviral Particle (Locus ID 23133)
Locus ID:	23133
Synonyms:	JHDM1F; KDM7B; MRXSSD; ZNF422
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	PHF8 - 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_001184896</a> , <a href="#">NM_001184897</a> , <a href="#">NM_001184898</a> , <a href="#">NM_015107</a> , <a href="#">NM_015107.1</a> , <a href="#">NM_015107.2</a> , <a href="#">NM_001184898.1</a> , <a href="#">NM_001184897.1</a> , <a href="#">NM_001184896.1</a> , <a href="#">BC017720</a> , <a href="#">BC042108</a> , <a href="#">BC053861</a> , <a href="#">NM_015107.3</a> , <a href="#">NM_001184898.2</a>
UniProt ID:	<a href="#">Q9UPP1</a>
Summary:	The protein encoded by this gene is a histone lysine demethylase that preferentially acts on histones in the monomethyl or dimethyl states. The encoded protein requires Fe(2+) ion, 2-oxoglutarate, and oxygen for its catalytic activity. The protein has an N-terminal PHD finger and a central Jumonji C domain. This gene is thought to function as a transcription activator. Defects in this gene are a cause of syndromic X-linked Siderius type intellectual disability (MRXSSD) and over-expression of this gene is associated with several forms of cancer. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2017]
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).