

## Product datasheet for **TL312141**

### INMT Human shRNA Plasmid Kit (Locus ID 11185)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	INMT Human shRNA Plasmid Kit (Locus ID 11185)
Locus ID:	11185
Synonyms:	TEMT
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	INMT - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 11185). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_001199219</a> , <a href="#">NM_006774</a> , <a href="#">NM_006774.2</a> , <a href="#">NM_006774.3</a> , <a href="#">NM_006774.4</a> , <a href="#">NM_001199219.1</a> , <a href="#">BC033813</a> , <a href="#">BC103712</a> , <a href="#">BC106902</a> , <a href="#">BC106903</a> , <a href="#">NM_001199219.2</a> , <a href="#">NM_006774.5</a>
UniProt ID:	<a href="#">O95050</a>
Summary:	N-methylation of endogenous and xenobiotic compounds is a major method by which they are degraded. This gene encodes an enzyme that N-methylates indoles such as tryptamine. Alternative splicing results in multiple transcript variants. Read-through transcription also exists between this gene and the downstream MINDY4 (aka FAM188B) gene. In rodents and other mammals such as cetartiodactyla this gene is in the opposite orientation compared to its orientation in human and other primates and this gene appears to have been lost in carnivora and chiroptera. [provided by RefSeq, Jul 2019]
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).