

## **Product datasheet for TL312383**

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## **HNMT Human shRNA Plasmid Kit (Locus ID 3176)**

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** HNMT Human shRNA Plasmid Kit (Locus ID 3176)

**Locus ID:** 3176

Synonyms: HMT; HNMT-S1; HNMT-S2; MRT51

Vector: pGFP-C-shLenti (TR30023)

E. coli Selection: Chloramphenicol (34 ug/ml)

**Mammalian Cell** 

Selection:

Puromycin

Format: Lentiviral plasmids

**Components:** HNMT - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 3176).

5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.

RefSeq: NM 001024074, NM 001024075, NM 006895, NM 006895.1, NM 006895.2, NM 001024074.1,

NM 001024074.2, NM 001024075.1, BC020677, BC020677.1, BC005907, NM 001024074.3,

NM 001024075.2, NM 006895.3

UniProt ID: P50135

**Summary:** In mammals, histamine is metabolized by two major pathways: N(tau)-methylation via

histamine N-methyltransferase and oxidative deamination via diamine oxidase. This gene encodes the first enzyme which is found in the cytosol and uses S-adenosyl-L-methionine as the methyl donor. In the mammalian brain, the neurotransmitter activity of histamine is controlled by N(tau)-methylation as diamine oxidase is not found in the central nervous system. A common genetic polymorphism affects the activity levels of this gene product in red blood cells. Multiple alternatively spliced transcript variants that encode different

proteins have been found for this gene. [provided by RefSeq, Jul 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 <u>techsupport@origene.com</u>. If you need a special design or shRNA sequence, please utilize our custom shRNA service.







## 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).