

## **Product datasheet for TL312474**

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## **HEXB Human shRNA Plasmid Kit (Locus ID 3074)**

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** HEXB Human shRNA Plasmid Kit (Locus ID 3074)

Locus ID: 3074

**Synonyms:** ENC-1AS; HEL-248; HEL-S-111

**Vector:** pGFP-C-shLenti (TR30023)

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

**Mammalian Cell** 

Selection:

Puromycin

Format: Lentiviral plasmids

**Components:** HEXB - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 3074).

5µg purified plasmid DNA per construct

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

RefSeq: NM 000521, NM 001292004, NM 000521.1, NM 000521.2, NM 000521.3, NM 001292004.1,

BC017378, BC017378.2

UniProt ID: P07686

**Summary:** Hexosaminidase B is the beta subunit of the lysosomal enzyme beta-hexosaminidase that,

together with the cofactor GM2 activator protein, catalyzes the degradation of the ganglioside GM2, and other molecules containing terminal N-acetyl hexosamines. Beta-hexosaminidase is composed of two subunits, alpha and beta, which are encoded by separate genes. Both beta-hexosaminidase alpha and beta subunits are members of family 20 of glycosyl hydrolases. Mutations in the alpha or beta subunit genes lead to an accumulation of GM2 ganglioside in neurons and neurodegenerative disorders termed the GM2 gangliosidoses.

Beta subunit gene mutations lead to Sandhoff disease (GM2-gangliosidosis type II).

Alternatively spliced transcript variants encoding different isoforms have been found for this

gene. [provided by RefSeq, May 2014]

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).