

## **Product datasheet for TL317024**

## OriGene Technologies, Inc.

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## **SNAPIN Human shRNA Plasmid Kit (Locus ID 23557)**

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** SNAPIN Human shRNA Plasmid Kit (Locus ID 23557)

**Locus ID:** 23557

Synonyms: BLOC1S7; BLOS7; BORCS3; SNAPAP

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

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

**Mammalian Cell** 

Selection:

Puromycin

Format: Lentiviral plasmids

**Components:** SNAPIN - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 23557).

5µg purified plasmid DNA per construct

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

RefSeq: NM 012437, NR 052019, NR 052020, NM 012437.1, NM 012437.3, NM 012437.5, BC004494,

BC000761, BM051828, NM 012437.6

UniProt ID: 095295

**Summary:** The protein encoded by this gene is a coiled-coil-forming protein that associates with the

SNARE (soluble N-ethylmaleimide-sensitive fusion protein attachment protein receptor) complex of proteins and the BLOC-1 (biogenesis of lysosome-related organelles) complex. Biochemical studies have identified additional binding partners. As part of the SNARE

complex, it is required for vesicle docking and fusion and regulates neurotransmitter release.

The BLOC-1 complex is required for the biogenesis of specialized organelles such as

melanosomes and platelet dense granules. Mutations in gene products that form the BLOC-1

complex have been identified in mouse strains that are models of Hermansky-Pudlak

syndrome. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jun

2012]

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).