

## **Product datasheet for TG503112**

**Becn1 Mouse shRNA Plasmid (Locus ID 56208)** 

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** Becn1 Mouse shRNA Plasmid (Locus ID 56208)

**Locus ID:** 56208

**Synonyms:** 4921513J16Rik; 5430417M23Rik; Atg6

**Vector:** pGFP-V-RS (TR30007)

E. coli Selection: Kanamycin

Mammalian Cell Puromycin

Selection:

Format: Retroviral plasmids

**Components:** Becn1 - Mouse, 4 unique 29mer shRNA constructs in retroviral GFP vector(Gene ID = 56208).

5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pGFP-V-RS Vector, TR30013, included for free.

**RefSeq:** BC005770, NM 019584, NM 019584.1, NM 019584.2, NM 019584.3, BC027738,

NM 001359819, NM 001359820, NM 001359821, NM 019584.4

**UniProt ID:** <u>O88597</u>

Summary: Plays a central role in autophagy (PubMed:10604474, PubMed:12372286, PubMed:19270693,

PubMed:28445460). Acts as core subunit of different PI3K complex forms that mediate formation of phosphatidylinositol 3-phosphate and are believed to play a role in multiple membrane trafficking pathways: PI3KC3-C1 is involved in initiation of autophagosomes and

PI3KC3-C2 in maturation of autophagosomes and endocytosis (PubMed:19270693,

PubMed:25275521). Involved in regulation of degradative endocytic trafficking and required for the abcission step in cytokinesis, probably in the context of PI3KC3-C2 (By similarity).

Essential for the formation of PI3KC3-C2 but not PI3KC3-C1 PI3K complex forms

(PubMed:25275521). Involved in endocytosis including endosome formation in neuronal cells (PubMed:25275521). May play a role in antiviral host defense (By similarity).[UniProtKB/Swiss-

Prot Function]

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>.



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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. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).