

Product datasheet for **TL309034**

Syntaxin 16 (STX16) Human shRNA Plasmid Kit (Locus ID 8675)

Product data:

Product Type:	shRNA Plasmids
Product Name:	Syntaxin 16 (STX16) Human shRNA Plasmid Kit (Locus ID 8675)
Locus ID:	8675
Synonyms:	SYN16
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	STX16 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 8675). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	NM_001001433 , NM_001001434 , NM_001134772 , NM_001134773 , NM_001204868 , NM_003763 , NR_037941 , NR_037942 , NR_037943 , NM_003763.1 , NM_003763.3 , NM_003763.4 , NM_003763.5 , NM_001001433.1 , NM_001001433.2 , NM_001134773.1 , NM_001134773.2 , NM_001134772.1 , NM_001134772.2 , NM_001204868.1 , NM_001001434.1 , BC019042 , BC019042.2 , BC073876 , NM_001134772.3 , NM_001001433.3 , NM_001134773.3 , NM_001204868.2
UniProt ID:	O14662
Summary:	This gene encodes a protein that is a member of the syntaxin or t-SNARE (target-SNAP receptor) family. These proteins are found on cell membranes and serve as the targets for V-SNARES (vesicle-SNAP receptors) permitting specific synaptic vesicle docking and fusion. A microdeletion in the region of chromosome 20 where this gene is located has been associated with pseudohypoparathyroidism type Ib. Multiple transcript variants have been found for this gene. Read-through transcription also exists between this gene and the neighboring downstream aminopeptidase-like 1 (NPEPL1) gene. [provided by RefSeq, Mar 2011]
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 techsupport@origene.com . If you need a special design or shRNA sequence, please utilize our custom shRNA service .



[View online »](#)

**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).