

## Product datasheet for RC201016L3V

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## Syntaxin 18 (STX18) (NM\_016930) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** Syntaxin 18 (STX18) (NM\_016930) Human Tagged ORF Clone Lentiviral Particle

Symbol: Syntaxin 18

Synonyms: Ufe1

Mammalian Cell P

Selection:

Puromycin

**Vector:** pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK
ACCN: NM 016930

ORF Size: 1005 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC201016).

Sequence:

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through

naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**RefSeg:** NM 016930.2

 RefSeq Size:
 2162 bp

 RefSeq ORF:
 1008 bp

 Locus ID:
 53407

 UniProt ID:
 Q9P2W9

Cytogenetics: 4p16.3-p16.2

**Protein Families:** Transmembrane

**Protein Pathways:** SNARE interactions in vesicular transport





**MW:** 38.7 kDa

**Gene Summary:** 

This gene encodes a member of the syntaxin family of soluble N-ethylmaleimide-sensitive factor attachment protein receptors (SNAREs) which is part of a membrane tethering complex that includes other SNAREs and several peripheral membrane proteins, and is involved in vesicular transport between the endoplasmic reticulum (ER) and the Golgi complex. The encoded protein is important for the organization of the smooth, rough, and exit site ER subdomains. A pseudogene of this gene has been identified. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Oct 2016]