

## Product datasheet for RC222797L3V

## OriGene Technologies, Inc.

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## SGK196 (POMK) (NM\_032237) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** SGK196 (POMK) (NM\_032237) Human Tagged ORF Clone Lentiviral Particle

Symbol: SGK196

Synonyms: MDDGA12; MDDGC12; SGK196

Mammalian Cell

Selection:

Puromycin

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

Tag:Myc-DDKACCN:NM\_032237

ORF Size: 1050 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC222797).

OTI Disclaimer:

Sequence:

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 032237.2

 RefSeq Size:
 1623 bp

 RefSeq ORF:
 1053 bp

 Locus ID:
 84197

 UniProt ID:
 Q9H5K3

 Cytogenetics:
 8p11.21

**Protein Families:** Druggable Genome, Protein Kinase, Transmembrane

**MW:** 40 kDa





## **Gene Summary:**

This gene encodes a protein that may be involved in the presentation of the laminin-binding O-linked carbohydrate chain of alpha-dystroglycan (a-DG), which forms transmembrane linkages between the extracellular matrix and the exoskeleton. Some pathogens use this O-linked carbohydrate unit for host entry. Loss of function compound heterozygous mutations in this gene were found in a human patient affected by the Walker-Warburg syndrome (WWS) phenotype. Mice lacking this gene contain misplaced neurons (heterotopia) in some regions of the brain, possibly from defects in neuronal migration. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, May 2013]