

## **Product datasheet for SC203970**

## SGK196 (POMK) (NM 032237) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: SGK196 (POMK) (NM\_032237) Human 3' UTR Clone

Symbol: SGK196

Synonyms: MDDGA12; MDDGC12; SGK196

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_032237

**Insert Size:** 587 bp

Insert Sequence: >SC203970 3'UTR clone of NM\_032237

The sequence shown below is from the reference sequence of NM\_032237. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCTTCCATTGTGGTCTGAGAAGGTACTTGATATGA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).



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## SGK196 (POMK) (NM\_032237) Human 3' UTR Clone - SC203970

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**RefSeq:** <u>NM 032237.5</u>

**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]

**Locus ID:** 84197 **MW:** 21.7