

Product datasheet for SC201967

FIG4 (NM 014845) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: FIG4 (NM_014845) Human 3' UTR Clone

Symbol: FIG4

Synonyms: ALS11; BTOP; CMT4J; dJ249I4.1; KIAA0274; SAC3; YVS

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_014845

Insert Size: 191 bp

Insert Sequence: >SC201967 3'UTR clone of NM_014845

The sequence shown below is from the reference sequence of NM_014845. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CTTTTCTGTCACTTGCAAATTCCAAATTATAGCTAATAAAGATGACTAGATAA

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).

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 014845.6</u>



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ORIGENE

Summary: The protein encoded by this gene belongs to the SAC domain-containing protein gene family.

The SAC domain, approximately 400 amino acids in length and consisting of seven conserved motifs, has been shown to possess phosphoinositide phosphatase activity. The yeast homolog, Sac1p, is involved in the regulation of various phosphoinositides, and affects diverse cellular functions such as actin cytoskeleton organization, Golgi function, and maintenance of vacuole morphology. Membrane-bound phosphoinositides function as signaling molecules and play a key role in vesicle trafficking in eukaryotic cells. Mutations in this gene have been associated with Charcot-Marie-Tooth disease, type 4J. [provided by

RefSeq, Jul 2008]

Locus ID: 9896

MW: 7.8