

Product datasheet for SC208016

VAMP1 (NM 016830) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: VAMP1 (NM_016830) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: VAMP1

Synonyms: CMS25; SPAX1; SYB1; VAMP-1

ACCN: NM_016830

Insert Size: 625 bp

Insert Sequence: >SC208016 3'UTR clone of NM_016830

The sequence shown below is from the reference sequence of NM $_$ 016830. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TAAA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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.



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



MW:

VAMP1 (NM_016830) Human 3' UTR Clone - SC208016

RefSeq: <u>NM 016830.4</u>

Summary: Synapotobrevins, syntaxins, and the synaptosomal-associated protein SNAP25 are the main

components of a protein complex involved in the docking and/or fusion of synaptic vesicles with the presynaptic membrane. The protein encoded by this gene is a member of the vesicle-associated membrane protein (VAMP)/synaptobrevin family. Mutations in this gene are associated with autosomal dominant spastic ataxia 1. Multiple alternative splice variants have been described, but the full-length nature of some variants has not been defined.

[provided by RefSeq, Jul 2014]

Locus ID: 6843

22.8