

Product datasheet for TL303193

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

VAM1 (MPP6) Human shRNA Plasmid Kit (Locus ID 51678)

Product data:

Product Type: shRNA Plasmids

Product Name: VAM1 (MPP6) Human shRNA Plasmid Kit (Locus ID 51678)

pGFP-C-shLenti (TR30023)

Locus ID: 51678

Synonyms: p55T; PALS2; VAM-1; VAM1

E. coli Selection: Chloramphenicol (34 ug/ml)

Mammalian Cell

Selection:

Vector:

Puromycin

Format: Lentiviral plasmids

Components: MPP6 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 51678).

5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.

RefSeq: NM 001303037, NM 016447, NM 016447.1, NM 016447.2, NM 016447.3, NM 001303037.1,

BC023638, BC023638.2, BC012034, BC013077, BC047330, BM685172, NM 001303037.2

UniProt ID: O9NZW5

Summary: Members of the peripheral membrane-associated guanylate kinase (MAGUK) family function

in tumor suppression and receptor clustering by forming multiprotein complexes containing distinct sets of transmembrane, cytoskeletal, and cytoplasmic signaling proteins. All MAGUKs contain a PDZ-SH3-GUK core and are divided into 4 subfamilies, DLG-like (see DLG1; MIM 601014), ZO1-like (see TJP1; MIM 601009), p55-like (see MPP1; MIM 305360), and LIN2-like (see CASK; MIM 300172), based on their size and the presence of additional domains. MPP6 is a member of the p55-like MAGUK subfamily (Tseng et al., 2001 [PubMed 11311936]).[supplied

by OMIM, Mar 2008]

shRNA Design: These shRNA constructs were designed against multiple splice variants at this gene locus. To

be certain that your variant of interest is targeted, please contact <u>techsupport@origene.com</u>. If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u>.







Performance Guaranteed:

OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).