

# Product datasheet for TL314143V

## CCM2 Human shRNA Lentiviral Particle (Locus ID 83605)

### **Product data:**

Product Type:	shRNA Lentiviral Particles
Product Name:	CCM2 Human shRNA Lentiviral Particle (Locus ID 83605)
Locus ID:	83605
Synonyms:	C7orf22; OSM; PP10187
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	CCM2 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, >10^7 TU/ml.
RefSeq:	<u>NM 001029835</u> , <u>NM 001167934</u> , <u>NM 001167935</u> , <u>NM 031443</u> , <u>NR 030770</u> , <u>NM 031443.1</u> , <u>NM 031443.2</u> , <u>NM 031443.3</u> , <u>NM 001029835.1</u> , <u>NM 001029835.2</u> , <u>NM 001167935.1</u> , <u>NM 001167934.1</u> , <u>BC004903</u> , <u>BC008859</u> , <u>BC016832</u> , <u>BC025958</u> , <u>BC063663</u> , <u>NM 001363458</u> , <u>NM 001363459</u> , <u>NM 031443.4</u>
UniProt ID:	Q9BSQ5
Summary:	This gene encodes a scaffold protein that functions in the stress-activated p38 Mitogen- activated protein kinase (MAPK) signaling cascade. The protein interacts with SMAD specific E3 ubiquitin protein ligase 1 (also known as SMURF1) via a phosphotyrosine binding domain to promote RhoA degradation. The protein is required for normal cytoskeletal structure, cell- cell interactions, and lumen formation in endothelial cells. Mutations in this gene result in cerebral cavernous malformations. Multiple transcript variants encoding different isoforms have been found for this gene.[provided by RefSeq, Nov 2009]
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> .



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#### **GRIGENE** CCM2 Human shRNA Lentiviral Particle (Locus ID 83605) – TL314143V

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).

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