

Product datasheet for **TL303337V**

MASP2 Human shRNA Lentiviral Particle (Locus ID 10747)

Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	MASP2 Human shRNA Lentiviral Particle (Locus ID 10747)
Locus ID:	10747
Synonyms:	MAP-2; MAP19; MASP-2; MASP1P1; sMAP
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	MASP2 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, $>10^7$ TU/ml.
RefSeq:	<u>NM_006610</u> , <u>NM_139208</u> , <u>NM_006610.1</u> , <u>NM_006610.2</u> , <u>NM_006610.3</u> , <u>NM_139208.1</u> , <u>NM_139208.2</u> , <u>BC052299</u> , <u>BC067359</u> , <u>BC080556</u> , <u>BC156086</u> , <u>BC156886</u> , <u>NM_139208.3</u> , <u>NM_006610.4</u>
UniProt ID:	<u>O00187</u>
Summary:	This gene encodes a member of the peptidase S1 family of serine proteases. The encoded preproprotein is proteolytically processed to generate A and B chains that heterodimerize to form the mature protease. This protease cleaves complement components C2 and C4 in order to generate C3 convertase in the lectin pathway of the complement system. The encoded protease also plays a role in the coagulation cascade through cleavage of prothrombin to form thrombin. Myocardial infarction and acute stroke patients exhibit reduced serum concentrations of the encoded protein. Alternative splicing results in multiple transcript variants, at least one of which encodes an isoform that is proteolytically processed. [provided by RefSeq, Feb 2016]
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 techsupport@origene.com . If you need a special design or shRNA sequence, please utilize our custom shRNA service .



[View online »](#)

**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).