

Product datasheet for **TR301355**

AMSH (STAMPB) Human shRNA Plasmid Kit (Locus ID 10617)

Product data:

Product Type:	shRNA Plasmids
Product Name:	AMSH (STAMPB) Human shRNA Plasmid Kit (Locus ID 10617)
Locus ID:	10617
Synonyms:	AMSH; MICCAP
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	STAMPB - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 10617). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	NM_006463 , NM_201647 , NM_213622 , NM_001353967 , NM_001353968 , NM_001353969 , NM_001353970 , NM_001353971 , NM_001353972 , NM_001353973 , NM_001353974 , NM_001353975 , NM_001353976 , NR_148668 , NR_148669 , NR_148670 , NR_148671 , NM_201647.1 , NM_201647.2 , NM_006463.1 , NM_006463.2 , NM_006463.3 , NM_006463.4 , NM_213622.1 , BC007682 , BC065574 , BC101467 , BC101469 , BM478027 , NM_006463.5 , NM_213622.4 , NM_201647.3
UniProt ID:	O95630
Summary:	Cytokine-mediated signal transduction in the JAK-STAT cascade requires the involvement of adaptor molecules. One such signal-transducing adaptor molecule contains an SH3 domain that is required for induction of MYC and cell growth. The protein encoded by this gene binds to the SH3 domain of the signal-transducing adaptor molecule, and plays a critical role in cytokine-mediated signaling for MYC induction and cell cycle progression. Multiple alternatively spliced transcript variants encoding the same protein isoform have been found for this gene. [provided by RefSeq, Jul 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 techsupport@origene.com . If you need a special design or shRNA sequence, please utilize our custom shRNA service .



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**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).