

Product datasheet for **TR301345**

STEAP2 Human shRNA Plasmid Kit (Locus ID 261729)

Product data:

Product Type:	shRNA Plasmids
Product Name:	STEAP2 Human shRNA Plasmid Kit (Locus ID 261729)
Locus ID:	261729
Synonyms:	IPCA1; PCANAP1; PUMPCn; STAMP1; STMP
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	STEAP2 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 261729). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	NM_001040665 , NM_001040666 , NM_001244944 , NM_001244945 , NM_001244946 , NM_152999 , NM_001040665.1 , NM_152999.1 , NM_152999.2 , NM_152999.3 , NM_001040666.1 , NM_001244946.1 , NM_001244945.1 , NM_001244944.1 , BC146341 , BC148824 , NM_001244944.2 , NM_001244945.2 , NM_001040665.2 , NM_001244946.2 , NM_152999.4
UniProt ID:	Q8NFT2
Summary:	This gene is a member of the STEAP family and encodes a multi-pass membrane protein that localizes to the Golgi complex, the plasma membrane, and the vesicular tubular structures in the cytosol. A highly similar protein in mouse has both ferrireductase and cupric reductase activity, and stimulates the cellular uptake of both iron and copper in vitro. Increased transcriptional expression of the human gene is associated with prostate cancer progression. Alternate transcriptional splice variants, encoding different isoforms, have been characterized. [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).