

Product datasheet for **TL309754V**

RPL15 Human shRNA Lentiviral Particle (Locus ID 6138)

Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	RPL15 Human shRNA Lentiviral Particle (Locus ID 6138)
Locus ID:	6138
Synonyms:	DBA12; EC45; L15; RPL10; RPLY10; RPYL10
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	RPL15 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, >10 ⁷ TU/ml.
RefSeq:	NM_001253379 , NM_001253380 , NM_001253382 , NM_001253383 , NM_001253384 , NM_002948 , NM_002948.1 , NM_002948.2 , NM_002948.3 , NM_002948.4 , NM_001253384.1 , NM_001253384.2 , NM_001253379.1 , NM_001253379.2 , NM_001253380.1 , NM_001253380.2 , NM_001253382.1 , NM_001253382.2 , NM_001253383.1 , NM_001253383.2 , BC014837 , BC014837.1 , BC068198 , BC068198.1 , BC070328 , BC071672 , BC081565 , NM_001253383.3 , NM_002948.5
UniProt ID:	P61313
Summary:	Ribosomes, the organelles that catalyze protein synthesis, consist of a small 40S subunit and a large 60S subunit. Together these subunits are composed of four RNA species and approximately 80 structurally distinct proteins. This gene encodes a member of the L15E family of ribosomal proteins and a component of the 60S subunit. This gene shares sequence similarity with the yeast ribosomal protein YL10 gene. Elevated expression of this gene has been observed in esophageal tumors and gastric cancer tissues, and deletion of this gene has been observed in a Diamond-Blackfan anemia (DBA) patient. As is typical for genes encoding ribosomal proteins, there are multiple processed pseudogenes of this gene dispersed through the genome. [provided by RefSeq, Mar 2017]
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).