

## Product datasheet for **TL309714**

### RPS3 Human shRNA Plasmid Kit (Locus ID 6188)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	RPS3 Human shRNA Plasmid Kit (Locus ID 6188)
Locus ID:	6188
Synonyms:	S3
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	RPS3 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 6188). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_001005</a> , <a href="#">NM_001256802</a> , <a href="#">NM_001260506</a> , <a href="#">NM_001260507</a> , <a href="#">NM_001005.1</a> , <a href="#">NM_001005.2</a> , <a href="#">NM_001005.3</a> , <a href="#">NM_001005.4</a> , <a href="#">NM_001260507.1</a> , <a href="#">NM_001256802.1</a> , <a href="#">NM_001260506.1</a> , <a href="#">BC034149</a> , <a href="#">BC034149.1</a> , <a href="#">BC003137</a> , <a href="#">BC003577</a> , <a href="#">BC013196</a> , <a href="#">BC013231</a> , <a href="#">BC029981</a> , <a href="#">BC071669</a> , <a href="#">BC071917</a> , <a href="#">BC100284</a> , <a href="#">BC150501</a> , <a href="#">BM831460</a> , <a href="#">BM928330</a> , <a href="#">NM_001005.5</a> , <a href="#">NM_001260506.2</a> , <a href="#">NM_001256802.2</a>
UniProt ID:	<a href="#">P23396</a>



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<b>Summary:</b>	<p>Ribosomes, the organelles that catalyze protein synthesis, consist of a small 40S subunit and a large 60S subunit. Together these subunits are composed of 4 RNA species and approximately 80 structurally distinct proteins. This gene encodes a ribosomal protein that is a component of the 40S subunit, where it forms part of the domain where translation is initiated. The protein belongs to the S3P family of ribosomal proteins. Studies of the mouse and rat proteins have demonstrated that the protein has an extraribosomal role as an endonuclease involved in the repair of UV-induced DNA damage. The protein appears to be located in both the cytoplasm and nucleus but not in the nucleolus. Higher levels of expression of this gene in colon adenocarcinomas and adenomatous polyps compared to adjacent normal colonic mucosa have been observed. This gene is co-transcribed with the small nucleolar RNA genes U15A and U15B, which are located in its first and fifth introns, respectively. As is typical for genes encoding ribosomal proteins, there are multiple processed pseudogenes of this gene dispersed through the genome. Multiple alternatively spliced transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, May 2012]</p>
<b>shRNA Design:</b>	<p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. If you need a special design or shRNA sequence, please utilize our <a href="#">custom shRNA service</a>.</p>
<b>Performance Guaranteed:</b>	<p>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.</p> <p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).</p>