

Product datasheet for TL319098V

OriGene Technologies, Inc.

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MRPL52 Human shRNA Lentiviral Particle (Locus ID 122704)

Product data:

Product Type: shRNA Lentiviral Particles

Locus ID: 122704

Vector: pGFP-C-shLenti (TR30023)

Format: Lentiviral particles

Components: MRPL52 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble

control), 0.5 ml each, >10^7 TU/ml.

RefSeq: NM_001318870, NM_178336, NM_180981, NM_180982, NM_181304, NM_181305, NM_181306,

NM_181307, NM_178336.1, NM_178336.2, NM_180982.1, NM_180982.2, NM_181307.1, NM_181307.2, NM_181306.1, NM_181306.2, NM_181304.1, NM_181304.2, NM_181305.1, NM_181305.2, NM_180981.1, BC068070, BC068070.1, BC055381, NM_180982.3, NM_181304.3, NM_181307.3, NM_181305.3,

NM_181306.3, NM_178336.3

UniProt ID: Q86TS9

Summary: Mammalian mitochondrial ribosomal proteins are encoded by nuclear genes and help in

protein synthesis within the mitochondrion. Mitochondrial ribosomes (mitoribosomes) consist of a small 28S subunit and a large 39S subunit. They have an estimated 75% protein to rRNA

composition compared to prokaryotic ribosomes, where this ratio is reversed. Another

difference between mammalian mitoribosomes and prokaryotic ribosomes is that the latter contain a 5S rRNA. Among different species, the proteins comprising the mitoribosome differ

greatly in sequence, and sometimes in biochemical properties, which prevents easy recognition by sequence homology. This gene encodes a 39S subunit protein which has no

bacterial homolog. Multiple transcript variants encoding different protein isoforms were

identified through sequence analysis. [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 $\underline{\mathsf{techsupport}} \underline{\mathsf{oorigene.com}}.$

If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u>.





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).