

Product datasheet for **TL303519**

LIPT1 Human shRNA Plasmid Kit (Locus ID 51601)

Product data:

Product Type:	shRNA Plasmids
Product Name:	LIPT1 Human shRNA Plasmid Kit (Locus ID 51601)
Locus ID:	51601
Synonyms:	LIPT1D
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	LIPT1 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 51601). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	NM_001204830 , NM_015929 , NM_145196 , NM_145197 , NM_145198 , NM_145199 , NR_037935 , NR_037936 , NM_145198.1 , NM_145198.2 , NM_015929.1 , NM_015929.2 , NM_015929.3 , NM_145199.1 , NM_145199.2 , NM_145197.1 , NM_145197.2 , NM_001204830.1 , NM_145196.1 , NM_145196.2 , BC070145 , BC070145.1 , BC007001 , BC009772 , NM_145199.3 , NM_145197.3 , NM_015929.4 , NM_145198.3
UniProt ID:	Q9Y234
Summary:	The process of transferring lipoic acid to proteins is a two-step process. The first step is the activation of lipoic acid by lipoate-activating enzyme to form lipoyl-AMP. For the second step, the protein encoded by this gene transfers the lipoyl moiety to apoproteins. Alternative splicing results in multiple transcript variants. A related pseudogene has been identified on chromosome 13. Read-through transcription also exists between this gene and the neighboring downstream mitochondrial ribosomal protein L30 (MRPL30) gene. [provided by RefSeq, Mar 2011]
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).