

## Product datasheet for **TL504245**

### Ctc1 Mouse shRNA Plasmid (Locus ID 68964)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Ctc1 Mouse shRNA Plasmid (Locus ID 68964)
Locus ID:	68964
Synonyms:	1500010J02Rik; AAF-132; AAF132
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	Ctc1 - Mouse, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 68964). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">BC060629</a> , <a href="#">NM_001013256</a> , <a href="#">NM_001143790</a> , <a href="#">NM_001281464</a> , <a href="#">NM_001281465</a> , <a href="#">NM_026889</a> , <a href="#">NR_026565</a> , <a href="#">NM_001013256.1</a> , <a href="#">NM_001013256.2</a> , <a href="#">NM_001143790.1</a> , <a href="#">NM_001281464.1</a> , <a href="#">NM_001281465.1</a> , <a href="#">BC060629.1</a> , <a href="#">BC147495</a> , <a href="#">BC014687</a> , <a href="#">BC066078</a> , <a href="#">BC072625</a> , <a href="#">BC147497</a>
UniProt ID:	<a href="#">Q5SUQ9</a>



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<b>Summary:</b>	<p>Component of the CST complex proposed to act as a specialized replication factor promoting DNA replication under conditions of replication stress or natural replication barriers such as the telomere duplex. The CST complex binds single-stranded DNA with high affinity in a sequence-independent manner, while isolated subunits bind DNA with low affinity by themselves. Initially the CST complex has been proposed to protect telomeres from DNA degradation (PubMed:19854130). However, the CST complex has been shown to be involved in several aspects of telomere replication. The CST complex inhibits telomerase and is involved in telomere length homeostasis; it is proposed to bind to newly telomerase-synthesized 3' overhangs and to terminate telomerase action implicating the association with the ACD:POT1 complex thus interfering with its telomerase stimulation activity. The CST complex is also proposed to be involved in fill-in synthesis of the telomeric C-strand probably implicating recruitment and activation of DNA polymerase alpha. The CST complex facilitates recovery from many forms of exogenous DNA damage; seems to be involved in the re-initiation of DNA replication at repaired forks and/or dormant origins. Involved in telomere maintenance. Involved in genome stability (By similarity). May be involved in telomeric C-strand fill-in during late S/G2 phase (PubMed:22748632).[UniProtKB/Swiss-Prot Function]</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>