

## Product datasheet for **TL514884**

### Anp32e Mouse shRNA Plasmid (Locus ID 66471)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Anp32e Mouse shRNA Plasmid (Locus ID 66471)
Locus ID:	66471
Synonyms:	2810018A15Rik; AI047746; AI326868; CPD1; LANP-L; LANPL; mLANP-L
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	Anp32e - Mouse, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 66471). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">BC005690</a> , <a href="#">BC080684</a> , <a href="#">NM_001253757</a> , <a href="#">NM_001253758</a> , <a href="#">NM_023210</a> , <a href="#">NM_023210.1</a> , <a href="#">NM_023210.2</a> , <a href="#">NM_023210.3</a> , <a href="#">NM_023210.4</a> , <a href="#">NM_001253758.1</a> , <a href="#">NM_001253757.1</a>
UniProt ID:	<a href="#">P97822</a>
Summary:	Histone chaperone that specifically mediates the genome-wide removal of histone H2A.Z/H2AFZ from the nucleosome: removes H2A.Z/H2AFZ from its normal sites of deposition, especially from enhancer and insulator regions. Not involved in deposition of H2A.Z/H2AFZ in the nucleosome. May stabilize the evicted H2A.Z/H2AFZ-H2B dimer, thus shifting the equilibrium towards dissociation and the off-chromatin state (PubMed:24463511). Inhibits activity of protein phosphatase 2A (PP2A). Does not inhibit protein phosphatase 1. May play a role in cerebellar development and synaptogenesis.[UniProtKB/Swiss-Prot Function]
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 <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> .



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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](mailto: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).