

# **Product datasheet for SC324797**

### OriGene Technologies, Inc.

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## LANPL (ANP32E) (NM\_001136479) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

Product Name: LANPL (ANP32E) (NM\_001136479) Human Untagged Clone

Tag: Tag Free Symbol: ANP32E

Synonyms: LANP-L; LANPL
Vector: pCMV6 series

Fully Sequenced ORF: >NCBI ORF sequence for NM\_001136479, the custom clone sequence may differ by one or

more nucleotides

**Restriction Sites:** Please inquire **ACCN:** NM 001136479

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).





#### **Reconstitution Method:**

- 1. Centrifuge at 5,000xg for 5min.
- 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
- 3. Close the tube and incubate for 10 minutes at room temperature.
- 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

**RefSeq:** NM 001136479.1, NP 001129951.1

RefSeq Size: 3146 bp
RefSeq ORF: 663 bp
Locus ID: 81611
UniProt ID: Q9BTT0
Cytogenetics: 1q21.2

**Protein Families:** Druggable Genome

**Gene 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

Function1

Transcript Variant: This variant (3) differs in the 5' UTR and initiates translation at a

downstream start codon compared to variant 1. The resulting isoform (3) is shorter at the N-

terminus compared to isoform 1.