

Product datasheet for SC334524

OriGene Technologies, Inc.

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RNA terminal phosphate cyclase like 1 (RCL1) (NM 001286699) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: RNA terminal phosphate cyclase like 1 (RCL1) (NM 001286699) Human Untagged Clone

Tag: Tag Free

Symbol: RNA terminal phosphate cyclase like 1

Synonyms: RNAC; RPCL1

Mammalian Cell Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001286699, the custom clone sequence may differ by one or

more nucleotides

AGCAAGACCCTCAAGTGA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001286699

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).

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).





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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: <u>NM 001286699.1</u>, <u>NP 001273628.1</u>

RefSeq Size: 1950 bp
RefSeq ORF: 648 bp
Locus ID: 10171
UniProt ID: Q9Y2P8
Cytogenetics: 9p24.1

Gene Summary: Does not have cyclase activity. Plays a role in 40S-ribosomal-subunit biogenesis in the early

pre-rRNA processing steps at sites A0, A1 and A2 that are required for proper maturation of

the 18S RNA (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) lacks two alternate exons in the 5' region and it thus differs in its 5' UTR and initiates translation at a downstream in-frame start codon, compared to variant 1. The encoded isoform (b) is shorter at the N-terminus, compared to isoform a. Both

variants 2 and 3 encode isoform b.