

## **Product datasheet for SC332158**

### OriGene Technologies, Inc.

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# Calcipressin 3 (RCAN3) (NM\_001251980) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

Product Name: Calcipressin 3 (RCAN3) (NM 001251980) Human Untagged Clone

Tag: Tag Free

Symbol: Calcipressin 3

Synonyms: DSCR1L2; hRCN3; MCIP3; RCN3

**Vector:** pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC332158 representing NM\_001251980.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

**CTGTGA** 

**Restriction Sites:** Sgfl-Mlul

**ACCN:** NM 001251980

**Insert Size:** 696 bp

**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).





#### **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 001251980.1</u>

 RefSeq Size:
 2503 bp

 RefSeq ORF:
 696 bp

 Locus ID:
 11123

 UniProt ID:
 Q9UKA8

 Cytogenetics:
 1p36.11

 MW:
 26.4 kDa

Gene Summary: Inhibits calcineurin-dependent transcriptional responses by binding to the catalytic domain of

calcineurin A. Could play a role during central nervous system development (By similarity).

[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (5) lacks an in-frame segment in the coding region, compared to variant 1. The resulting isoform (2) lacks an internal segment, compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.