

### Product datasheet for RC228921L4V

#### OriGene Technologies, Inc.

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## Calcitonin receptor (CALCR) (NM 001164738) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** Calcitonin receptor (CALCR) (NM\_001164738) Human Tagged ORF Clone Lentiviral Particle

Symbol: Calcitonin receptor
Synonyms: CRT; CT-R; CTR1

**Mammalian Cell** 

Selection:

Puromycin

**Vector:** pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

**ACCN:** NM\_001164738

ORF Size: 1422 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC228921).

Sequence:
OTI Disclaimer:

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**RefSeq:** NM 001164738.1, NP 001158210.1

 RefSeq ORF:
 1425 bp

 Locus ID:
 799

 UniProt ID:
 P30988

 Cytogenetics:
 7q21.3

**Protein Families:** Druggable Genome, GPCR, Transmembrane

**Protein Pathways:** Neuroactive ligand-receptor interaction

MW: 55.34 kDa





# Calcitonin receptor (CALCR) (NM\_001164738) Human Tagged ORF Clone Lentiviral Particle – RC228921L4V

#### **Gene Summary:**

This gene encodes a high affinity receptor for the peptide hormone calcitonin and belongs to a subfamily of seven transmembrane-spanning G protein-coupled receptors. The encoded protein is involved in maintaining calcium homeostasis and in regulating osteoclast-mediated bone resorption. Polymorphisms in this gene have been associated with variations in bone mineral density and onset of osteoporosis. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Sep 2009]