

## Product datasheet for RC221294L4V

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## **CLCN6 (NM\_001286) Human Tagged ORF Clone Lentiviral Particle**

**Product data:** 

Product Type: Lentiviral Particles

**Product Name:** CLCN6 (NM\_001286) Human Tagged ORF Clone Lentiviral Particle

Symbol: CLCN6

Synonyms: CLC-6; CONRIBA

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

**ACCN:** NM\_001286 **ORF Size:** 2607 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC221294).

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.

**RefSeg:** NM 001286.3

 RefSeq Size:
 5713 bp

 RefSeq ORF:
 2610 bp

 Locus ID:
 1185

 UniProt ID:
 P51797

 Cytogenetics:
 1p36.22

**Domains:** CBS, voltage\_CLC

**Protein Families:** Druggable Genome, Ion Channels: Other, Transmembrane





## CLCN6 (NM\_001286) Human Tagged ORF Clone Lentiviral Particle - RC221294L4V

**MW:** 97.2 kDa

**Gene Summary:** This gene encodes a member of the voltage-dependent chloride channel protein family.

Members of this family can function as either chloride channels or antiporters. This protein is primarily localized to late endosomes and functions as a chloride/proton antiporter. Alternate splicing results in both coding and non-coding variants. Additional alternately spliced variants have been described but their full-length structure is unknown. [provided by

RefSeq, Mar 2012]