

Product datasheet for RC221281L4V

OriGene Technologies, Inc.

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Claudin 10 (CLDN10) (NM 182848) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Claudin 10 (CLDN10) (NM_182848) Human Tagged ORF Clone Lentiviral Particle

Symbol: CLDN10

Synonyms: CPETRL3; HELIX; OSP-L; OSPL

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

ACCN: NM_182848

ORF Size: 678 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC221281).

OTI Disclaimer:

Sequence:

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 182848.3, NP 878268.1

 RefSeq Size:
 2658 bp

 RefSeq ORF:
 681 bp

 Locus ID:
 9071

 UniProt ID:
 P78369

 Cytogenetics:
 13q32.1

Protein Families: Transmembrane

Protein Pathways: Cell adhesion molecules (CAMs), Leukocyte transendothelial migration, Tight junction





MW: 24.3 kDa

Gene Summary:

This gene encodes a member of the claudin family. Claudins are integral membrane proteins and components of tight junction strands. Tight junction strands serve as a physical barrier to prevent solutes and water from passing freely through the paracellular space between epithelial or endothelial cell sheets, and also play critical roles in maintaining cell polarity and signal transductions. The expression level of this gene is associated with recurrence of primary hepatocellular carcinoma. Six alternatively spliced transcript variants encoding different isoforms have been reported, but the transcript sequences of some variants are not determined.[provided by RefSeq, Jun 2010]