

## Product datasheet for RC223907L2V

## 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

## VE Cadherin (CDH5) (NM\_001795) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

Product Name: VE Cadherin (CDH5) (NM 001795) Human Tagged ORF Clone Lentiviral Particle

Symbol: CDH5

Synonyms: 7B4; CD144

Mammalian Cell

Selection:

None

**Vector:** pLenti-C-mGFP (PS100071)

Tag: mGFP

**ACCN:** NM\_001795 **ORF Size:** 2352 bp

**ORF Nucleotide** 

OTI Disclaimer:

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Sequence:

The ORF insert of this clone is exactly the same as(RC223907).

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 001795.3

 RefSeq Size:
 4134 bp

 RefSeq ORF:
 2355 bp

 Locus ID:
 1003

 UniProt ID:
 P33151

 Cytogenetics:
 16q21

**Domains:** Cadherin\_C\_term, CA

**Protein Families:** Druggable Genome, ES Cell Differentiation/IPS, Transmembrane





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

**MW:** 87.5 kDa

**Gene Summary:** This gene encodes a classical cadherin of the cadherin superfamily. The encoded

preproprotein is proteolytically processed to generate the mature glycoprotein. This calcium-dependent cell-cell adhesion molecule is comprised of five extracellular cadherin repeats, a transmembrane region and a highly conserved cytoplasmic tail. Functioning as a classical cadherin by imparting to cells the ability to adhere in a homophilic manner, this protein plays a role in endothelial adherens junction assembly and maintenance. This gene is located in a gene cluster in a region on the long arm of chromosome 16 that is involved in loss of heterozygosity events in breast and prostate cancer. [provided by RefSeq, Nov 2015]