

## Product datasheet for RC214969L3V

## OriGene Technologies, Inc.

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## COL13A1 (NM\_080800) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** COL13A1 (NM\_080800) Human Tagged ORF Clone Lentiviral Particle

Symbol: COL13A1

Synonyms: CMS19; COLXIIIA1

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK
ACCN: NM 080800

ORF Size: 2058 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC214969).

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 080800.2

RefSeq Size: 3079 bp
RefSeq ORF: 2061 bp
Locus ID: 1305
UniProt ID: Q5TAT6

Cytogenetics: 10q22.1

MW: 66.7 kDa







## **Gene Summary:**

This gene encodes the alpha chain of one of the nonfibrillar collagens. The function of this gene product is not known, however, it has been detected at low levels in all connective tissue-producing cells so it may serve a general function in connective tissues. Unlike most of the collagens, which are secreted into the extracellular matrix, collagen XIII contains a transmembrane domain and the protein has been localized to the plasma membrane. The transcripts for this gene undergo complex and extensive splicing involving at least eight exons. Like other collagens, collagen XIII is a trimer; it is not known whether this trimer is composed of one or more than one alpha chain isomer. A number of alternatively spliced transcript variants have been described, but the full length nature of some of them has not been determined. [provided by RefSeq, Jul 2008]