

## Product datasheet for RC207689L2V

### OriGene Technologies, Inc.

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# CST6 (NM\_001323) Human Tagged ORF Clone Lentiviral Particle

### **Product data:**

Product Type: Lentiviral Particles

**Product Name:** CST6 (NM\_001323) Human Tagged ORF Clone Lentiviral Particle

Symbol:CST6Synonyms:ECTD15

Mammalian Cell

Selection:

None

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

Tag: mGFP

**ACCN:** NM\_001323

ORF Size: 447 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC207689).

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 001323.2

 RefSeq Size:
 618 bp

 RefSeq ORF:
 450 bp

 Locus ID:
 1474

 UniProt ID:
 Q15828

 Cytogenetics:
 11q13.1

**Protein Families:** Secreted Protein

**MW:** 16.5 kDa







#### **Gene Summary:**

The cystatin superfamily encompasses proteins that contain multiple cystatin-like sequences. Some of the members are active cysteine protease inhibitors, while others have lost or perhaps never acquired this inhibitory activity. There are three inhibitory families in the superfamily, including the type 1 cystatins (stefins), type 2 cystatins and the kininogens. The type 2 cystatin proteins are a class of cysteine proteinase inhibitors found in a variety of human fluids and secretions, where they appear to provide protective functions. This gene encodes a cystatin from the type 2 family, which is down-regulated in metastatic breast tumor cells as compared to primary tumor cells. Loss of expression is likely associated with the progression of a primary tumor to a metastatic phenotype. [provided by RefSeq, Jul 2008]