

## Product datasheet for RC200657L1V

## OriGene Technologies, Inc.

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## Citrate transport protein (SLC25A1) (NM\_005984) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** Citrate transport protein (SLC25A1) (NM\_005984) Human Tagged ORF Clone Lentiviral Particle

Symbol: Citrate transport protein

**Synonyms:** CMS23; CTP; D2L2AD; SEA; SLC20A3

Mammalian Cell

Selection:

None

**Vector:** pLenti-C-Myc-DDK (PS100064)

Tag: Myc-DDK

**ACCN:** NM\_005984

ORF Size: 933 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC200657).

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.

**RefSeq:** <u>NM 005984.1</u>

 RefSeq Size:
 1619 bp

 RefSeq ORF:
 936 bp

 Locus ID:
 6576

 UniProt ID:
 P53007

 Cytogenetics:
 22q11.21

**Domains:** mito\_carr





## Citrate transport protein (SLC25A1) (NM\_005984) Human Tagged ORF Clone Lentiviral Particle – RC200657L1V

**Protein Families:** Druggable Genome

MW: 34.01 kDa

**Gene Summary:** This gene encodes a member of the mitochondrial carrier subfamily of solute carrier

proteins. Members of this family include nuclear-encoded transporters that translocate small metabolites across the mitochondrial membrane. This protein regulates the movement of citrate across the inner membranes of the mitochondria. Mutations in this gene have been associated with combined D-2- and L-2-hydroxyglutaric aciduria. Pseudogenes of this gene have been identified on chromosomes 7, 11, 16, and 19. Alternative splicing results in

multiple transcript variants. [provided by RefSeq, Dec 2013]