

## Product datasheet for RC201757L1V

## 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

## Aspartate Aminotransferase (GOT1) (NM\_002079) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** Aspartate Aminotransferase (GOT1) (NM\_002079) Human Tagged ORF Clone Lentiviral

**Particle** 

**Symbol:** Aspartate Aminotransferase

Synonyms: AST1; ASTQTL1; cAspAT; cCAT; GIG18

Mammalian Cell

Selection:

None

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

 Tag:
 Myc-DDK

 ACCN:
 NM\_002079

ORF Size: 1239 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC201757).

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.

**RefSeq:** <u>NM 002079.1</u>

 RefSeq Size:
 2140 bp

 RefSeq ORF:
 1242 bp

 Locus ID:
 2805

 UniProt ID:
 P17174

 Cytogenetics:
 10q24.2

**Domains:** aminotran\_1\_2





## Aspartate Aminotransferase (GOT1) (NM\_002079) Human Tagged ORF Clone Lentiviral Particle – RC201757L1V

Protein Pathways: Alanine, aspartate and glutamate metabolism, Arginine and proline metabolism, Cysteine and

methionine metabolism, Metabolic pathways, Phenylalanine, tyrosine and tryptophan

biosynthesis, Phenylalanine metabolism, Tyrosine metabolism

MW: 46.2 kDa

**Gene Summary:** Glutamic-oxaloacetic transaminase is a pyridoxal phosphate-dependent enzyme which exists

in cytoplasmic and mitochondrial forms, GOT1 and GOT2, respectively. GOT plays a role in amino acid metabolism and the urea and tricarboxylic acid cycles. The two enzymes are

homodimeric and show close homology. [provided by RefSeq, Jul 2008]