

Product datasheet for RC206858L4V

OriGene Technologies, Inc.

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Ornithine Decarboxylase (ODC1) (NM_002539) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Ornithine Decarboxylase (ODC1) (NM_002539) Human Tagged ORF Clone Lentiviral Particle

Symbol: Ornithine Decarboxylase

Synonyms: BABS; NEDBA; NEDBIA; ODC

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_002539 **ORF Size:** 1383 bp

ORF Nucleotide

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OTI Disclaimer:

Sequence:

The ORF insert of this clone is exactly the same as(RC206858).

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 002539.1

 RefSeq Size:
 2307 bp

 RefSeq ORF:
 1386 bp

 Locus ID:
 4953

 UniProt ID:
 P11926

 Cytogenetics:
 2p25.1

Domains: Orn_Arg_deC_N

Protein Families: Druggable Genome





Ornithine Decarboxylase (ODC1) (NM_002539) Human Tagged ORF Clone Lentiviral Particle – RC206858L4V

Protein Pathways: Arginine and proline metabolism, Glutathione metabolism, Metabolic pathways

MW: 51.1 kDa

Gene Summary: This gene encodes the rate-limiting enzyme of the polyamine biosynthesis pathway which

catalyzes ornithine to putrescine. The activity level for the enzyme varies in response to growth-promoting stimuli and exhibits a high turnover rate in comparison to other mammalian proteins. Originally localized to both chromosomes 2 and 7, the gene encoding this enzyme has been determined to be located on 2p25, with a pseudogene located on 7q31-qter. Multiple alternatively spliced transcript variants encoding distinct isoforms have

been identified. [provided by RefSeq, Dec 2013]