

Product datasheet for RC200639L2V

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

Lipoamide Dehydrogenase (DLD) (NM_000108) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Lipoamide Dehydrogenase (DLD) (NM_000108) Human Tagged ORF Clone Lentiviral Particle

Symbol: DLD

Synonyms: DLDD; DLDH; E3; GCSL; LAD; OGDC-E3; PHE3

Mammalian Cell

Selection:

None

Vector: pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_000108 **ORF Size:** 1527 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC200639).

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 000108.3

 RefSeq Size:
 3613 bp

 RefSeq ORF:
 1530 bp

 Locus ID:
 1738

 UniProt ID:
 P09622

 Cytogenetics:
 7q31.1

Domains: pyr_redox, pyr_redox_dim

Protein Families: Druggable Genome





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Protein Pathways: Citrate cycle (TCA cycle), Glycine, serine and threonine metabolism, Glycolysis /

Gluconeogenesis, Metabolic pathways, Pyruvate metabolism, Valine, leucine and isoleucine

degradation

MW: 54.2 kDa

Gene Summary: This gene encodes a member of the class-I pyridine nucleotide-disulfide oxidoreductase

family. The encoded protein has been identified as a moonlighting protein based on its ability to perform mechanistically distinct functions. In homodimeric form, the encoded protein functions as a dehydrogenase and is found in several multi-enzyme complexes that regulate energy metabolism. However, as a monomer, this protein can function as a protease.

Mutations in this gene have been identified in patients with E3-deficient maple syrup urine disease and lipoamide dehydrogenase deficiency. Alternative splicing results in multiple

transcript variants. [provided by RefSeq, Jan 2014]