

Product datasheet for RC203949L4V

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

PHGDH (NM_006623) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: PHGDH (NM_006623) Human Tagged ORF Clone Lentiviral Particle

Symbol: PHGDH

Synonyms: 3-PGDH; 3PGDH; HEL-S-113; NLS; NLS1; PDG; PGDD; PGDH; PHGDHD; SERA

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

ACCN: NM_006623 **ORF Size:** 1599 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC203949).

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.

RefSeg: NM 006623.2, NP 006614.2

 RefSeq Size:
 2021 bp

 RefSeq ORF:
 1602 bp

 Locus ID:
 26227

 UniProt ID:
 043175

 Cytogenetics:
 1p12

Domains: 2-Hacid_DH, 2-Hacid_DH_C

Protein Families: Druggable Genome, Stem cell - Pluripotency





PHGDH (NM_006623) Human Tagged ORF Clone Lentiviral Particle - RC203949L4V

Protein Pathways: Glycine, serine and threonine metabolism, Metabolic pathways

MW: 56.7 kDa

Gene Summary: This gene encodes the enzyme which is involved in the early steps of L-serine synthesis in

animal cells. L-serine is required for D-serine and other amino acid synthesis. The enzyme requires NAD/NADH as a cofactor and forms homotetramers for activity. Mutations in this gene have been found in a family with congenital microcephaly, psychomotor retardation and other symptoms. Multiple alternatively spliced transcript variants have been found, however the full-length nature of most are not known. [provided by RefSeq, Aug 2011]