

Product datasheet for RC201592L4V

OriGene Technologies, Inc.

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CYB5R3 (NM_000398) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: CYB5R3 (NM_000398) Human Tagged ORF Clone Lentiviral Particle

Symbol: CYB5R3

Synonyms: B5R; DIA1

Mammalian Cell Puromycin

Selection:

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

Tag: mGFP

ACCN: NM_000398

ORF Size: 903 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC201592).

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 000398.4

 RefSeq Size:
 2923 bp

 RefSeq ORF:
 906 bp

 Locus ID:
 1727

 UniProt ID:
 P00387

 Cytogenetics:
 22q13.2

Domains: NAD_binding_1, FAD_binding_6

Protein Families: Druggable Genome





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Protein Pathways: Amino sugar and nucleotide sugar metabolism

MW: 34.2 kDa

Gene Summary: This gene encodes cytochrome b5 reductase, which includes a membrane-bound form in

somatic cells (anchored in the endoplasmic reticulum, mitochondrial and other membranes) and a soluble form in erythrocytes. The membrane-bound form exists mainly on the cytoplasmic side of the endoplasmic reticulum and functions in desaturation and elongation of fatty acids, in cholesterol biosynthesis, and in drug metabolism. The erythrocyte form is located in a soluble fraction of circulating erythrocytes and is involved in methemoglobin reduction. The membrane-bound form has both membrane-binding and catalytic domains, while the soluble form has only the catalytic domain. Alternate splicing results in multiple transcript variants. Mutations in this gene cause methemoglobinemias. [provided by RefSeq,

Jan 2010]