

Product datasheet for RC204177L3V

OriGene Technologies, Inc.

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Solute carrier family 22 member 5 (SLC22A5) (NM_003060) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Solute carrier family 22 member 5 (SLC22A5) (NM_003060) Human Tagged ORF Clone

Lentiviral Particle

Symbol: Solute carrier family 22 member 5

Synonyms: CDSP; OCTN2

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK
ACCN: NM_003060

ORF Size: 1671 bp

ORF Nucleotide

Sequence:

The ORF insert of this clone is exactly the same as(RC204177).

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.

RefSeq: <u>NM 003060.2</u>

 RefSeq Size:
 3295 bp

 RefSeq ORF:
 1674 bp

 Locus ID:
 6584

 UniProt ID:
 076082

 Cytogenetics:
 5q31.1

Domains: sugar_tr





Protein Families: Transmembrane

MW: 62.7 kDa

Gene Summary: Polyspecific organic cation transporters in the liver, kidney, intestine, and other organs are

critical for elimination of many endogenous small organic cations as well as a wide array of drugs and environmental toxins. The encoded protein is a plasma integral membrane protein which functions both as an organic cation transporter and as a sodium-dependent high affinity carnitine transporter. The encoded protein is involved in the active cellular uptake of

carnitine. Mutations in this gene are the cause of systemic primary carnitine deficiency (CDSP), an autosomal recessive disorder manifested early in life by hypoketotic hypoglycemia

and acute metabolic decompensation, and later in life by skeletal myopathy or

cardiomyopathy. Alternative splicing of this gene results in multiple transcript variants.

[provided by RefSeq, Apr 2015]