

#### OriGene Technologies, Inc.

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# Product datasheet for RC228962L3V

## FATP2 (SLC27A2) (NM\_001159629) Human Tagged ORF Clone Lentiviral Particle

## **Product data:**

Product Type:	Lentiviral Particles
Product Name:	FATP2 (SLC27A2) (NM_001159629) Human Tagged ORF Clone Lentiviral Particle
Symbol:	SLC27A2
Synonyms:	ACSVL1; FACVL1; FATP2; hFACVL1; HsT17226; VLACS; VLCS
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
Tag:	Myc-DDK
ACCN:	NM_001159629
ORF Size:	1701 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC228962).
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. <u>More info</u>
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 001159629.1, NP 001153101.1</u>
RefSeq ORF:	1704 bp
Locus ID:	11001
UniProt ID:	<u>014975</u>
Cytogenetics:	15q21.2
Protein Families:	Transmembrane
Protein Pathways:	PPAR signaling pathway
MW:	64.4 kDa



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Gene Summary:The protein encoded by this gene is an isozyme of long-chain fatty-acid-coenzyme A ligase<br/>family. Although differing in substrate specificity, subcellular localization, and tissue<br/>distribution, all isozymes of this family convert free long-chain fatty acids into fatty acyl-CoA<br/>esters, and thereby play a key role in lipid biosynthesis and fatty acid degradation. This<br/>isozyme activates long-chain, branched-chain and very-long-chain fatty acids containing 22 or<br/>more carbons to their CoA derivatives. It is expressed primarily in liver and kidney, and is<br/>present in both endoplasmic reticulum and peroxisomes, but not in mitochondria. Its<br/>decreased peroxisomal enzyme activity is in part responsible for the biochemical pathology<br/>in X-linked adrenoleukodystrophy. Alternatively spliced transcript variants encoding different<br/>isoforms have been found for this gene. [provided by RefSeq, Apr 2009]

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