

Product datasheet for RC216737L3

LIPT1 (NM_015929) Human Tagged Lenti ORF Clone

Product data:

Product Type: Expression Plasmids

Product Name: LIPT1 (NM_015929) Human Tagged Lenti ORF Clone

Tag: Myc-DDK

Symbol: LIPT1

Synonyms: LIPT1D

Mammalian Cell Puromycin

Selection:

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

E. coli Selection: Chloramphenicol (34 ug/mL)

ORF Nucleotide The ORF insert of this clone is exactly the same as(RC216737).

Sequence:

Restriction Sites: Sgfl-Mlul

Cloning Scheme:





^{*} The last codon before the Stop codon of the ORF.

ACCN: NM_015929

ORF Size: 1119 bp



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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.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: NM 015929.2

 RefSeq Size:
 1456 bp

 RefSeq ORF:
 1122 bp

 Locus ID:
 51601

 UniProt ID:
 Q9Y234

Cytogenetics: 2q11.2

Domains: BPL LipA LipB

Protein Pathways: Lipoic acid metabolism, Metabolic pathways

MW: 42.5 kDa

Gene Summary: The process of transferring lipoic acid to proteins is a two-step process. The first step is the

activation of lipoic acid by lipoate-activating enzyme to form lipoyl-AMP. For the second step, the protein encoded by this gene transfers the lipoyl moiety to apoproteins. Alternative splicing results in multiple transcript variants. A related pseudogene has been identified on

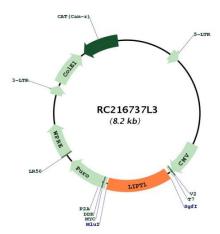
chromosome 13. Read-through transcription also exists between this gene and the

neighboring downstream mitochondrial ribosomal protein L30 (MRPL30) gene. [provided by

RefSeq, Mar 2011]



Product images:



Circular map for RC216737L3