

## Product datasheet for RC227032L4V

## OriGene Technologies, Inc.

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## LITAF (NM\_001136473) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** LITAF (NM\_001136473) Human Tagged ORF Clone Lentiviral Particle

Symbol: LITAF

**Synonyms:** PIG7; SIMPLE; TP53I7

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

**ACCN:** NM\_001136473

ORF Size: 456 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC227032).

Sequence:

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:** NM 001136473.1, NP 001129945.1

 RefSeq ORF:
 459 bp

 Locus ID:
 9516

 UniProt ID:
 Q99732

Cytogenetics: 16p13.13

**Protein Families:** Druggable Genome, Transcription Factors

**MW:** 15.6 kDa







## **Gene Summary:**

Lipopolysaccharide is a potent stimulator of monocytes and macrophages, causing secretion of tumor necrosis factor-alpha (TNF-alpha) and other inflammatory mediators. This gene encodes lipopolysaccharide-induced TNF-alpha factor, which is a DNA-binding protein and can mediate the TNF-alpha expression by direct binding to the promoter region of the TNF-alpha gene. The transcription of this gene is induced by tumor suppressor p53 and has been implicated in the p53-induced apoptotic pathway. Mutations in this gene cause Charcot-Marie-Tooth disease type 1C (CMT1C) and may be involved in the carcinogenesis of extramammary Paget's disease (EMPD). Multiple alternatively spliced transcript variants have been found for this gene. [provided by RefSeq, Dec 2014]