## Product datasheet for RC204079L4V

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200
Rockville, MD 20850, US
Phone: +1-888-267-4436
https://www.origene.com techsupport@origene.com
EU: info-de@origene.com
CN: techsupport@origene.cn

## AIPL1 (NM_014336) Human Tagged ORF Clone Lentiviral Particle

## Product data:

Product Type:
Product Name:
Symbol:
Synonyms:
Mammalian Cell
Selection:
Vector:
Tag:
ACCN:
ORF Size:
ORF Nucleotide
Sequence:
OTI Disclaimer:

OTI Annotation:

RefSeq:
RefSeq Size:
RefSeq ORF:
Locus ID:
UniProt ID:
Cytogenetics:
Protein Families:
MW:

Lentiviral Particles
AIPL1 (NM_014336) Human Tagged ORF Clone Lentiviral Particle
AIPL1
AIPL2; LCA4
Puromycin
pLenti-C-mGFP-P2A-Puro (PS100093)
mGFP
NM_014336
1152 bp
The ORF insert of this clone is exactly the same as(RC204079).

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
This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

## NM 014336.3

2990 bp
1155 bp
23746
Q9NZN9
17p13.2
Druggable Genome
43.9 kDa

Leber congenital amaurosis (LCA) is the most severe inherited retinopathy with the earliest age of onset and accounts for at least 5\% of all inherited retinal diseases. Affected individuals are diagnosed at birth or in the first few months of life with nystagmus, severely impaired vision or blindness and an abnormal or flat electroretinogram. The photoreceptor/pinealexpressed gene, AIPL1, encoding aryl-hydrocarbon interacting protein-like 1, is located within the LCA4 candidate region. The encoded protein contains three tetratricopeptide motifs, consistent with chaperone or nuclear transport activity. Mutations in this gene may cause approximately 20\% of recessive LCA. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2014]

