

Product datasheet for RC200395L4V

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

Apolipoprotein E (APOE) (NM 000041) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Apolipoprotein E (APOE) (NM_000041) Human Tagged ORF Clone Lentiviral Particle

Symbol: APOE

Synonyms: AD2; APO-E; ApoE4; LDLCQ5; LPG

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

ACCN: NM_000041

ORF Size: 951 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC200395).

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

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.

RefSeg: NM 000041.2

RefSeq Size: 1223 bp
RefSeq ORF: 954 bp
Locus ID: 348

 UniProt ID:
 P02649

 Cytogenetics:
 19q13.32

Domains: Apolipoprotein

Protein Families: Adult stem cells, Druggable Genome, Secreted Protein, Stem cell - Pluripotency





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Protein Pathways: Alzheimer's disease

MW: 36.2 kDa

Gene Summary: The protein encoded by this gene is a major apoprotein of the chylomicron. It binds to a

specific liver and peripheral cell receptor, and is essential for the normal catabolism of triglyceride-rich lipoprotein constituents. This gene maps to chromosome 19 in a cluster with

the related apolipoprotein C1 and C2 genes. Mutations in this gene result in familial

dysbetalipoproteinemia, or type III hyperlipoproteinemia (HLP III), in which increased plasma cholesterol and triglycerides are the consequence of impaired clearance of chylomicron and

VLDL remnants. [provided by RefSeq, Jun 2016]