

Product datasheet for RC214351L1V

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

ADCY7 (NM_001114) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: ADCY7 (NM_001114) Human Tagged ORF Clone Lentiviral Particle

Symbol: ADCY7

Synonyms: AC7

Mammalian Cell None

Selection:

Vector: pLenti-C-Myc-DDK (PS100064)

Tag: Myc-DDK
ACCN: NM 001114

ORF Size: 3240 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC214351).

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.

RefSeg: NM 001114.2

RefSeq Size:6157 bpRefSeq ORF:3243 bp

Locus ID: 113

UniProt ID: P51828
Cytogenetics: 16q12.1
Domains: CYCc

Protein Families: Druggable Genome, Transmembrane





ADCY7 (NM_001114) Human Tagged ORF Clone Lentiviral Particle - RC214351L1V

Protein Pathways: Calcium signaling pathway, Chemokine signaling pathway, Dilated cardiomyopathy, Gap

junction, GnRH signaling pathway, Melanogenesis, Oocyte meiosis, Progesterone-mediated

oocyte maturation, Purine metabolism, Vascular smooth muscle contraction

MW: 120.1 kDa

Gene Summary: This gene encodes a membrane-bound adenylate cyclase that catalyses the formation of

cyclic AMP from ATP and is inhibitable by calcium. The product of this gene is a member of the adenylyl cyclase class-4/guanylyl cyclase enzyme family that is characterized by the presence of twelve membrane-spanning domains in its sequences. Several transcript variants

have been observed for this gene, but the full-length natures of only two have been

determined so far. [provided by RefSeq, Oct 2013]