

## Product datasheet for RC200580L4V

## OriGene Technologies, Inc.

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## ADA2a (TADA2A) (NM\_133439) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

Product Name: ADA2a (TADA2A) (NM\_133439) Human Tagged ORF Clone Lentiviral Particle

Symbol: ADA2a

Synonyms: ADA2; ADA2A; hADA2; KL04P; TADA2L

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

**ACCN:** NM\_133439

ORF Size: 915 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC200580).

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 133439.2

 RefSeq Size:
 1327 bp

 RefSeq ORF:
 918 bp

 Locus ID:
 6871

 UniProt ID:
 075478

Cytogenetics: 17q12

**Protein Families:** Transcription Factors

MW: 36.1 kDa







## **Gene Summary:**

Many DNA-binding transcriptional activator proteins enhance the initiation rate of RNA polymerase II-mediated gene transcription by interacting functionally with the general transcription machinery bound at the basal promoter. Adaptor proteins are usually required for this activation, possibly to acetylate and destabilize nucleosomes, thereby relieving chromatin constraints at the promoter. The protein encoded by this gene is a transcriptional activator adaptor and has been found to be part of the PCAF histone acetylase complex. Several alternatively spliced transcript variants encoding different isoforms of this gene have been described, but the full-length nature of some of these variants has not been determined. [provided by RefSeq, Oct 2009]