

## Product datasheet for RC227720L3V

## OriGene Technologies, Inc.

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## SIRT1 (NM\_001142498) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** SIRT1 (NM\_001142498) Human Tagged ORF Clone Lentiviral Particle

Symbol: SIRT1

Synonyms: SIR2; SIR2alpha; SIR2L1

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK

**ACCN:** NM\_001142498

ORF Size: 1356 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC227720).

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 001142498.1, NP 001135970.1

RefSeq ORF: 1359 bp
Locus ID: 23411
UniProt ID: Q96EB6

Cytogenetics: 10q21.3

**Protein Families:** Druggable Genome, Stem cell - Pluripotency, Transcription Factors

MW: 50.3 kDa







## **Gene Summary:**

This gene encodes a member of the sirtuin family of proteins, homologs to the yeast Sir2 protein. Members of the sirtuin family are characterized by a sirtuin core domain and grouped into four classes. The functions of human sirtuins have not yet been determined; however, yeast sirtuin proteins are known to regulate epigenetic gene silencing and suppress recombination of rDNA. Studies suggest that the human sirtuins may function as intracellular regulatory proteins with mono-ADP-ribosyltransferase activity. The protein encoded by this gene is included in class I of the sirtuin family. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Dec 2008]