

## Product datasheet for RC218841L1V

## OriGene Technologies, Inc.

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## TAZ (NM\_000116) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

Product Type: Lentiviral Particles

**Product Name:** TAZ (NM\_000116) Human Tagged ORF Clone Lentiviral Particle

Symbol: TAZ

Synonyms: BTHS; CMD3A; EFE; EFE2; G4.5; LVNCX; TAZ; Taz1

Mammalian Cell

Selection:

None

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

 Tag:
 Myc-DDK

 ACCN:
 NM\_000116

ORF Size: 876 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC218841).

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 000116.2

 RefSeq Size:
 1904 bp

 RefSeq ORF:
 879 bp

 Locus ID:
 6901

 UniProt ID:
 Q16635

 Cytogenetics:
 Xq28

**Domains:** Acyltransferase

**Protein Families:** ES Cell Differentiation/IPS, Transmembrane





ORÏGENE

MW: 33.3 kDa

**Gene Summary:** 

This gene encodes a protein that is expressed at high levels in cardiac and skeletal muscle. Mutations in this gene have been associated with a number of clinical disorders including Barth syndrome, dilated cardiomyopathy (DCM), hypertrophic DCM, endocardial fibroelastosis, and left ventricular noncompaction (LVNC). Multiple transcript variants encoding different isoforms have been described. A long form and a short form of each of these isoforms is produced; the short form lacks a hydrophobic leader sequence and may exist as a cytoplasmic protein rather than being membrane-bound. Other alternatively spliced transcripts have been described but the full-length nature of all these transcripts is not known. [provided by RefSeq, Jul 2008]