

## Product datasheet for RC214918L2V

## OriGene Technologies, Inc.

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## KDM5B (NM\_006618) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

Product Type: Lentiviral Particles

**Product Name:** KDM5B (NM\_006618) Human Tagged ORF Clone Lentiviral Particle

Symbol: KDM5B

Synonyms: CT31; JARID1B; MRT65; PLU-1; PLU1; PPP1R98; PUT1; RBBP2H1A; RBP2-H1

**Mammalian Cell** 

Selection:

None

**Vector:** pLenti-C-mGFP (PS100071)

Tag: mGFP

**ACCN:** NM\_006618 **ORF Size:** 4632 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC214918).

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 006618.3

 RefSeq Size:
 6393 bp

 RefSeq ORF:
 4635 bp

 Locus ID:
 10765

 UniProt ID:
 Q9UGL1

 Cytogenetics:
 1q32.1

**Domains:** ARID, PHD, JmjC, JmjN, zf-C5HC2

**Protein Families:** Druggable Genome, Transcription Factors





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**MW:** 175.5 kDa

**Gene Summary:** This gene encodes a lysine-specific histone demethylase that belongs to the jumonji/ARID

domain-containing family of histone demethylases. The encoded protein is capable of demethylating tri-, di- and monomethylated lysine 4 of histone H3. This protein plays a role in the transcriptional repression or certain tumor suppressor genes and is upregulated in certain cancer cells. This protein may also play a role in genome stability and DNA repair. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Nov 2016]