

## Product datasheet for RC229337L3V

## OriGene Technologies, Inc.

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## **SETMAR (NM\_006515) Human Tagged ORF Clone Lentiviral Particle**

**Product data:** 

**Product Type:** Lentiviral Particles

Product Name: SETMAR (NM 006515) Human Tagged ORF Clone Lentiviral Particle

Symbol: SETMAR

**Synonyms:** Mar1; METNASE

**Mammalian Cell** 

Selection:

Puromycin

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

Tag: Myc-DDK
ACCN: NM 006515

ORF Size: 2052 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC229337).

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 006515.3, NP 006506.3

 RefSeq ORF:
 2055 bp

 Locus ID:
 6419

 UniProt ID:
 Q53H47

Cytogenetics: Q53H47

Q53H47

**Domains:** SET, Transposase\_1, PreSET, Pre-SET

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

**Protein Pathways:** Lysine degradation





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**MW:** 77.9 kDa

**Gene Summary:** 

This gene encodes a fusion protein that contains an N-terminal histone-lysine N-methyltransferase domain and a C-terminal mariner transposase domain. The encoded protein binds DNA and functions in DNA repair activities including non-homologous end joining and double strand break repair. The SET domain portion of this protein specifically methylates histone H3 lysines 4 and 36. This gene exists as a fusion gene only in anthropoid primates, other organisms lack mariner transposase domain. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Jan 2013]