

## Product datasheet for **RC229002L4V**

### SETDB2 (NM\_001160308) Human Tagged ORF Clone Lentiviral Particle

#### Product data:

|                           |  |
|---------------------------|--|
| Product Type:             | Lentiviral Particles   |
| Product Name:             | SETDB2 (NM_001160308) Human Tagged ORF Clone Lentiviral Particle   |
| Symbol:                   | SETDB2   |
| Synonyms:                 | C13orf4; CLLD8; CLLL8; KMT1F   |
| Mammalian Cell Selection: | Puromycin  |
| Vector:                   | pLenti-C-mGFP-P2A-Puro (PS100093)  |
| Tag:                      | mGFP   |
| ACCN:                     | NM_001160308   |
| ORF Size:                 | 2121 bp  |
| ORF Nucleotide Sequence:  | The ORF insert of this clone is exactly the same as(RC229002).   |
| 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. <a href="#">More info</a> |
| 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:                   | <a href="#">NM_001160308.1</a>   |
| RefSeq ORF:               | 2124 bp  |
| Locus ID:                 | 83852  |
| UniProt ID:               | <a href="#">Q96T68</a>   |
| Cytogenetics:             | 13q14.2  |
| Protein Families:         | Druggable Genome   |
| Protein Pathways:         | Lysine degradation   |
| MW:                       | 80.5 kDa   |



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**Gene Summary:**

This gene encodes a member of a family of proteins that contain a methyl-CpG-binding domain (MBD) and a SET domain and function as histone methyltransferases. This protein is recruited to heterochromatin and plays a role in the regulation of chromosome segregation. This region is commonly deleted in chronic lymphocytic leukemia. Naturally-occurring readthrough transcription occurs from this gene to the downstream PHF11 (PHD finger protein 11) gene. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2016]