

## Product datasheet for RC212469L3V

## OriGene Technologies, Inc.

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## SPAG11B (NM 058200) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** SPAG11B (NM\_058200) Human Tagged ORF Clone Lentiviral Particle

Symbol:

EDDM2B; EP2; EP2C; EP2D; HE2; HE2C; SPAG11; SPAG11A Synonyms:

**Mammalian Cell** 

Selection:

ACCN:

Puromycin

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

Myc-DDK Tag: NM 058200

**ORF Size:** 324 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC212469).

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 058200.2, NP 478107.2

RefSeq Size: 755 bp RefSeq ORF: 327 bp Locus ID: 10407 **UniProt ID:** Q08648 Cytogenetics: 8p23.1

**Protein Families:** Secreted Protein

MW: 12 kDa







## **Gene Summary:**

This gene encodes several androgen-dependent, epididymis-specific secretory proteins. The specific functions of these proteins have not been determined, but they are thought to be involved in sperm maturation. Some of the isoforms contain regions of similarity to beta-defensins, a family of antimicrobial peptides. The gene is located on chromosome 8p23 near the defensin gene cluster. Alternative splicing of this gene results in seven transcript variants encoding different isoforms. Two different N-terminal and five different C-terminal protein sequences are encoded by the splice variants. Two additional variants have been described, but their full length sequences have not been determined. [provided by RefSeq, Jul 2008]