

## Product datasheet for RC208258L4V

## OriGene Technologies, Inc.

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## **ELL (NM 006532) Human Tagged ORF Clone Lentiviral Particle**

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** ELL (NM\_006532) Human Tagged ORF Clone Lentiviral Particle

Symbol:

C19orf17; ELL1; MEN; PPP1R68 Synonyms:

**Mammalian Cell** 

Puromycin

Selection:

Vector:

pLenti-C-mGFP-P2A-Puro (PS100093)

mGFP Tag:

NM 006532 ACCN: **ORF Size:** 1863 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC208258).

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 006532.2

RefSeq Size: 4077 bp RefSeq ORF: 1866 bp Locus ID: 8178 **UniProt ID:** P55199 Cytogenetics: 19p13.11

**Protein Families: Transcription Factors** 

MW: 68.3 kDa





## **Gene Summary:**

Elongation factor component of the super elongation complex (SEC), a complex required to increase the catalytic rate of RNA polymerase II transcription by suppressing transient pausing by the polymerase at multiple sites along the DNA. Elongation factor component of the little elongation complex (LEC), a complex required to regulate small nuclear RNA (snRNA) gene transcription by RNA polymerase II and III (PubMed:22195968, PubMed:23932780). Specifically required for stimulating the elongation step of RNA polymerase II- and III-dependent snRNA gene transcription (PubMed:23932780). ELL also plays an early role before its assembly into in the SEC complex by stabilizing RNA polymerase II recruitment/initiation and entry into the pause site. Required to stabilize the pre-initiation complex and early elongation.[UniProtKB/Swiss-Prot Function]