

Product datasheet for RC207333L4V

OriGene Technologies, Inc.

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EIF4E (NM_001968) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: EIF4E (NM_001968) Human Tagged ORF Clone Lentiviral Particle

Symbol: EIF4E

Synonyms: AUTS19; CBP; eIF-4E; EIF4E1; EIF4EL1; EIF4F

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

ACCN: NM_001968

ORF Size: 651 bp

ORF Nucleotide

tide The OPE

OTI Disclaimer:

Sequence:

The ORF insert of this clone is exactly the same as(RC207333).

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 001968.2

 RefSeq Size:
 4749 bp

 RefSeq ORF:
 654 bp

 Locus ID:
 1977

 UniProt ID:
 P06730

 Cytogenetics:
 4q23

Domains: IF4E

Protein Pathways: Insulin signaling pathway, mTOR signaling pathway







MW: 25.1 kDa

Gene Summary:

The protein encoded by this gene is a component of the eukaryotic translation initiation factor 4F complex, which recognizes the 7-methylguanosine cap structure at the 5' end of messenger RNAs. The encoded protein aids in translation initiation by recruiting ribosomes to the 5'-cap structure. Association of this protein with the 4F complex is the rate-limiting step in translation initiation. This gene acts as a proto-oncogene, and its expression and activation is associated with transformation and tumorigenesis. Several pseudogenes of this gene are found on other chromosomes. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Sep 2015]