

Product datasheet for SC334695

OriGene Technologies, Inc.

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EIF4EL3 (EIF4E2) (NM_001282958) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: EIF4EL3 (EIF4E2) (NM_001282958) Human Untagged Clone

Tag: Tag Free Symbol: EIF4E2

Synonyms: 4E-LP; 4EHP; EIF4EL3; h4EHP; IF4e

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001282958, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

ACCN: NM 001282958

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).





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Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: <u>NM 001282958.1</u>, <u>NP 001269887.1</u>

 RefSeq Size:
 3630 bp

 RefSeq ORF:
 711 bp

 Locus ID:
 9470

 Cytogenetics:
 2q37.1

Protein Families: Transcription Factors

Protein Pathways: Insulin signaling pathway, mTOR signaling pathway

Gene Summary: Recognizes and binds the 7-methylguanosine-containing mRNA cap during an early step in

the initiation (PubMed:9582349, PubMed:17368478, PubMed:25624349). Acts as a repressor of translation initiation (PubMed:22751931). In contrast to EIF4E, it is unable to bind eIF4G (EIF4G1, EIF4G2 or EIF4G3), suggesting that it acts by competing with EIF4E and block

assembly of eIF4F at the cap (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) contains alternate 3' exon structure and it thus differs in the 3' coding region and 3' UTR, compared to variant 1. The encoded isoform (B) has a distinct

C-terminus and is shorter than isoform A.