

## Product datasheet for SC206995

## EIF4A1 (NM 001416) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** EIF4A1 (NM\_001416) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: EIF4A1

Synonyms: DDX2A; EIF-4A; eIF-4A-I; EIF4A; eIF4A-I

ACCN: NM 001416

Insert Size: 550 bp

>SC206995 3'UTR clone of NM\_001416 **Insert Sequence:** 

The sequence shown below is from the reference sequence of NM\_001416. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCTGGGGGCTGAGGAGCAGCAGGAGGGGAGGGAAGGGAGCCAAGGGATGGACATCTTGTCATTTTTT TTCTTTGAATAAATGTCACTTTTTGAGGCAAAAGAAGGAACCGTGAACATTTTAGACACCCTTTTCTTT TAACCTAGTAACCTCCAGATCCCAGAGGCTCTCCTCACCTCAGCTGAGCTCCTTTGAAAGTGATTCAAG GTTGCCCAGGGGGTTGTCCCCAGGTGGGGGGAAGCAGGGGAGAAAATGGTAGCCATTTTTACATTGT TTTGTATAGTATTTATTGATTCAGGAAACAAACACAAAATTCTGAATAAAATGACTTGGAAACTGCC 

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul

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

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: NM 001416.4



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## EIF4A1 (NM\_001416) Human 3' UTR Clone - SC206995

Summary: ATP-dependent RNA helicase which is a subunit of the eIF4F complex involved in cap

recognition and is required for mRNA binding to ribosome. In the current model of

translation initiation, eIF4A unwinds RNA secondary structures in the 5'-UTR of mRNAs which

is necessary to allow efficient binding of the small ribosomal subunit, and subsequent

scanning for the initiator codon.[UniProtKB/Swiss-Prot Function]

**Locus ID:** 1973

**MW:** 19.9