

Product datasheet for SC207653

OriGene Technologies, Inc.

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DEPDC5 (NM_001007188) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: DEPDC5 (NM_001007188) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: DEPDC5

Synonyms: DEP.5; FFEVF; FFEVF1

ACCN: NM_001007188

Insert Size: 592 bp

Insert Sequence: >SC207653 3'UTR clone of NM_001007188

The sequence shown below is from the reference sequence of NM_001007188. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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.





MW:

DEPDC5 (NM_001007188) Human 3' UTR Clone - SC207653

RefSeq: <u>NM 001007188.4</u>

Summary: This gene encodes a member of the IML1 family of proteins involved in G-protein signaling

pathways. The mechanistic target of rapamycin complex 1 (mTORC1) pathway regulates cell

growth by sensing the availability of nutrients. The protein encoded by this gene is a

component of the GATOR1 (GAP activity toward Rags) complex which inhibits the amino acid-

sensing branch of the mTORC1 pathway. Mutations in this gene are associated with autosomal dominant familial focal epilepsy with variable foci. A single nucleotide polymorphism in an intron of this gene has been associated with an increased risk of hepatocellular carcinoma in individuals with chronic hepatitis C virus infection. Alternative

splicing results in multiple transcript variants. [provided by RefSeq, Mar 2014]

Locus ID: 9681

22.3

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