

Product datasheet for RC216583L4V

OriGene Technologies, Inc.

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Deformed Epidermal Autoregulatory Factor 1 (DEAF1) (NM_021008) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Deformed Epidermal Autoregulatory Factor 1 (DEAF1) (NM_021008) Human Tagged ORF

Clone Lentiviral Particle

Symbol: Deformed Epidermal Autoregulatory Factor 1
Synonyms: MRD24; NEDHELS; NUDR; SPN; VSVS; ZMYND5

Mammalian Cell

Selection:

Puromycin

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

Tag: mGFP

ACCN: NM_021008 **ORF Size:** 1695 bp

ORF Nucleotide

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Sequence:

The ORF insert of this clone is exactly the same as(RC216583).

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: <u>NM 021008.2</u>

 RefSeq Size:
 2737 bp

 RefSeq ORF:
 1698 bp

 Locus ID:
 10522

 UniProt ID:
 075398

 Cytogenetics:
 11p15.5

Domains: SAND, zf-MYND





Deformed Epidermal Autoregulatory Factor 1 (DEAF1) (NM_021008) Human Tagged ORF Clone Lentiviral Particle - RC216583L4V

Protein Families: Secreted Protein, Transcription Factors, Transmembrane

MW: 59.1 kDa

Gene Summary: This gene encodes a zinc finger domain-containing protein that functions as a regulator of

transcription. The encoded proteins binds to its own promoter as well as to that of several target genes. Activity of this protein is important in the regulation of embryonic development. Mutations in this gene have been found in individuals with autosomal dominant cognitive disability. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jun

2014]