

Product datasheet for SC334957

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

PRUNE (PRUNE1) (NM_001303229) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: PRUNE (PRUNE1) (NM_001303229) Human Untagged Clone

Tag: Tag Free
Symbol: PRUNE1

Synonyms: DRES-17; DRES17; H-PRUNE; HTCD37; NMIHBA; PRUNE

Mammalian Cell

Neomycin

Selection:

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

Fully Sequenced ORF: >NCBI ORF sequence for NM_001303229, the custom clone sequence may differ by one or

more nucleotides

GATCTCACTGTCACAGTCTACCACAGCCTCCCTGTCCAAGAAGTGA

Restriction Sites: Sgfl-Mlul

ACCN: NM 001303229

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: NM 001303229.1, NP 001290158.1

 RefSeq Size:
 2898 bp

 RefSeq ORF:
 816 bp

 Locus ID:
 58497

 UniProt ID:
 Q86TP1

 Cytogenetics:
 1q21.3

Protein Pathways: Purine metabolism

Gene Summary: This gene encodes a member of the DHH protein superfamily of phosphoesterases. This

protein has been found to function as both a nucleotide phosphodiesterase and an

exopolyphosphatase. This protein is believed to stimulate cancer progression and metastases through the induction of cell motility. A pseuodgene has been identified on chromosome 13. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Dec 2014] Transcript Variant: This variant (2) lacks one alternate internal exon resulting in a distinct 5' UTR and the use of an in-frame downstream start codon compared to variant 1. The resulting

protein (isoform 2) has a shorter N-terminus than isoform 1.