

OriGene Technologies, Inc.

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Product datasheet for MR217991L3V

Papd5 (NM_001164497) Mouse Tagged ORF Clone Lentiviral Particle

Product data:

Product Type:	Lentiviral Particles
Product Name:	Papd5 (NM_001164497) Mouse Tagged ORF Clone Lentiviral Particle
Symbol:	Papd5
Synonyms:	5730445M16Rik; 5830428A09
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
Tag:	Myc-DDK
ACCN:	NM_001164497
ORF Size:	2040 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(MR217991).
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. <u>More info</u>
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 001164497.1</u>
RefSeq Size:	4642 bp
RefSeq ORF:	2043 bp
Locus ID:	214627
UniProt ID:	<u>Q68ED3</u>
Cytogenetics:	8 C3



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Gene Summary:Terminal nucleotidyltransferase that catalyzes preferentially the transfert of ATP and GTP on
RNA 3' poly(A) tail creating a heterogeneous 3' poly(A) tail leading to mRNAs stabilization by
protecting mRNAs from active deadenylation (By similarity). Also functions as a catalytic
subunit of a TRAMP-like complex which has a poly(A) RNA polymerase activity and is involved
in a post-transcriptional quality control mechanism. Polyadenylation with short oligo(A) tails is
required for the degradative activity of the exosome on several of its nuclear RNA substrates.
Doesn't need a cofactor for polyadenylation activity (in vitro). Plays a role in replication-
dependent histone mRNA degradation, probably through terminal uridylation of mature
histone mRNAs. May play a role in sister chromatid cohesion (By similarity).[UniProtKB/Swiss-
Prot Function]

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