

OriGene Technologies, Inc.

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Product datasheet for RC212656L3V

PDLIM7 (NM_203352) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type:	Lentiviral Particles
Product Name:	PDLIM7 (NM_203352) Human Tagged ORF Clone Lentiviral Particle
Symbol:	PDLIM7
Synonyms:	LMP1; LMP3
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
Tag:	Myc-DDK
ACCN:	NM_203352
ORF Size:	1269 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC212656).
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 203352.1</u>
RefSeq Size:	1607 bp
RefSeq ORF:	1272 bp
Locus ID:	9260
UniProt ID:	<u>Q9NR12</u>
Cytogenetics:	5q35.3
Protein Families:	Druggable Genome
MW:	46.3 kDa



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Gene Summary:The protein encoded by this gene is representative of a family of proteins composed of
conserved PDZ and LIM domains. LIM domains are proposed to function in protein-protein
recognition in a variety of contexts including gene transcription and development and in
cytoskeletal interaction. The LIM domains of this protein bind to protein kinases, whereas the
PDZ domain binds to actin filaments. The gene product is involved in the assembly of an
actin filament-associated complex essential for transmission of ret/ptc2 mitogenic signaling.
The biological function is likely to be that of an adapter, with the PDZ domain localizing the
LIM-binding proteins to actin filaments of both skeletal muscle and nonmuscle tissues.
Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Jul
2008]

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