

OriGene Technologies, Inc.

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Product datasheet for RC222394L4V

ADAM15 (NM_207191) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type:	Lentiviral Particles
Product Name:	ADAM15 (NM_207191) Human Tagged ORF Clone Lentiviral Particle
Symbol:	ADAM15
Synonyms:	MDC15
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-mGFP-P2A-Puro (PS100093)
Tag:	mGFP
ACCN:	NM_207191
ORF Size:	2316 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC222394).
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 207191.2</u>
RefSeq Size:	2782 bp
RefSeq ORF:	2319 bp
Locus ID:	8751
UniProt ID:	<u>Q13444</u>
Cytogenetics:	1q21.3
Protein Families:	Druggable Genome, Protease, Transmembrane
MW:	83.7 kDa



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Gene Summary:The protein encoded by this gene is a member of the ADAM (a disintegrin and
metalloproteinase) protein family. ADAM family members are type I transmembrane
glycoproteins known to be involved in cell adhesion and proteolytic ectodomain processing
of cytokines and adhesion molecules. This protein contains multiple functional domains
including a zinc-binding metalloprotease domain, a disintegrin-like domain, as well as a EGF-
like domain. Through its disintegrin-like domain, this protein specifically interacts with the
integrin beta chain, beta 3. It also interacts with Src family protein-tyrosine kinases in a
phosphorylation-dependent manner, suggesting that this protein may function in cell-cell
adhesion as well as in cellular signaling. Multiple alternatively spliced transcript variants
encoding distinct isoforms have been observed. [provided by RefSeq, Jul 2008]

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