

## OriGene Technologies, Inc.

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## Product datasheet for RC200345L4V

## TRAF4 (NM\_004295) Human Tagged ORF Clone Lentiviral Particle

## **Product data:**

Product Type:	Lentiviral Particles
Product Name:	TRAF4 (NM_004295) Human Tagged ORF Clone Lentiviral Particle
Symbol:	TRAF4
Synonyms:	CART1; MLN62; RNF83
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-mGFP-P2A-Puro (PS100093)
Tag:	mGFP
ACCN:	NM_004295
ORF Size:	1410 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC200345).
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 004295.3</u>
RefSeq Size:	2902 bp
RefSeq ORF:	1413 bp
Locus ID:	9618
UniProt ID:	<u>Q9BUZ4</u>
Cytogenetics:	17q11.2
Domains:	zf-TRAF, RING, MATH
Protein Families:	Druggable Genome



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<b>CRIGENE</b> TRAF4 (NM_004295) Human Tagged ORF Clone Lentiviral Particle – RC200345L4V	
Protein Pathways:	Pathways in cancer, Small cell lung cancer
MW:	53.5 kDa
Gene Summary:	This gene encodes a member of the TNF receptor associated factor (TRAF) family. TRAF proteins are associated with, and mediate signal transduction from members of the TNF receptor superfamily. The encoded protein has been shown to interact with neurotrophin receptor, p75 (NTR/NTSR1), and negatively regulate NTR induced cell death and NF-kappa B activation. This protein has been found to bind to p47phox, a cytosolic regulatory factor included in a multi-protein complex known as NAD(P)H oxidase. This protein thus, is thought to be involved in the oxidative activation of MAPK8/JNK. Alternatively spliced transcript variants have been observed but the full-length nature of only one has been determined. [provided by RefSeq, Jul 2008]

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