

Product datasheet for RC200345L3V

OriGene Technologies, Inc.

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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:

CART1; MLN62; RNF83 Synonyms:

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK

NM 004295 ACCN:

ORF Size: 1410 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC200345).

OTI Disclaimer:

Sequence:

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. More info

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: NM 004295.3

RefSeq Size: 2902 bp RefSeq ORF: 1413 bp Locus ID: 9618 **UniProt ID:** Q9BUZ4

Cytogenetics: 17q11.2

Domains: zf-TRAF, RING, MATH

Protein Families: Druggable Genome





TRAF4 (NM_004295) Human Tagged ORF Clone Lentiviral Particle - RC200345L3V

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]