

Product datasheet for RC211276L2V

OriGene Technologies, Inc.

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CD153 (TNFSF8) (NM_001244) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: CD153 (TNFSF8) (NM_001244) Human Tagged ORF Clone Lentiviral Particle

Symbol: CD153

Synonyms: CD30L; CD30LG; CD153; TNLG3A

Mammalian Cell

Selection:

None

Vector: pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_001244

ORF Size: 702 bp

ORF Nucleotide

OTI Disclaimer:

The ORF insert of this clone is exactly the same as(RC211276).

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.

RefSeg: NM 001244.2

 RefSeq Size:
 3797 bp

 RefSeq ORF:
 705 bp

 Locus ID:
 944

 UniProt ID:
 P32971

Cytogenetics: 9q32-q33.1

Domains: TNF

Protein Families: Druggable Genome, Transmembrane





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Protein Pathways: Cytokine-cytokine receptor interaction

MW: 26 kDa

Gene Summary: The protein encoded by this gene is a cytokine that belongs to the tumor necrosis factor

(TNF) ligand family. This cytokine is a ligand for TNFRSF8/CD30, which is a cell surface antigen and a marker for Hodgkin lymphoma and related hematologic malignancies. The engagement of this cytokine expressed on B cell surface plays an inhibitory role in modulating Ig class switch. This cytokine was shown to enhance cell proliferation of some lymphoma cell lines, while to induce cell death and reduce cell proliferation of other lymphoma cell lines. The pleiotropic biologic activities of this cytokine on different CD30+ lymphoma cell lines may play a pathophysiologic role in Hodgkin's and some non-Hodgkin's lymphomas. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Nov