

## Product datasheet for RC202088L3V

## OriGene Technologies, Inc.

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## TRIM21 (NM\_003141) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

Product Type: Lentiviral Particles

Product Name: TRIM21 (NM 003141) Human Tagged ORF Clone Lentiviral Particle

Symbol: TRIM21

Synonyms: RNF81; Ro/SSA; RO52; SSA; SSA1

Mammalian Cell

Selection:

Puromycin

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

 Tag:
 Myc-DDK

 ACCN:
 NM\_003141

 ORF Size:
 1425 bp

**ORF Nucleotide** 

OTI Disclaimer:

Sequence:

The ORF insert of this clone is exactly the same as(RC202088).

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 003141.3

 RefSeq Size:
 1946 bp

 RefSeq ORF:
 1428 bp

 Locus ID:
 6737

 UniProt ID:
 P19474

 Cytogenetics:
 11p15.4

**Domains:** zf-B\_box, RING, SPRY, PRY

**Protein Families:** Druggable Genome





## TRIM21 (NM\_003141) Human Tagged ORF Clone Lentiviral Particle - RC202088L3V

**Protein Pathways:** Systemic lupus erythematosus

**MW:** 54.2 kDa

**Gene Summary:** This gene encodes a member of the tripartite motif (TRIM) family. The TRIM motif includes

three zinc-binding domains, a RING, a B-box type 1 and a B-box type 2, and a coiled-coil region. The encoded protein is part of the RoSSA ribonucleoprotein, which includes a single polypeptide and one of four small RNA molecules. The RoSSA particle localizes to both the cytoplasm and the nucleus. RoSSA interacts with autoantigens in patients with Sjogren syndrome and systemic lupus erythematosus. Alternatively spliced transcript variants for this gene have been described but the full-length nature of only one has been determined.

[provided by RefSeq, Jul 2008]