

## Product datasheet for RC219594L4V

## OriGene Technologies, Inc.

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## PIAS1 (NM\_016166) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

Product Type: Lentiviral Particles

**Product Name:** PIAS1 (NM\_016166) Human Tagged ORF Clone Lentiviral Particle

Symbol: PIAS1

**Synonyms:** DDXBP1; GBP; GU/RH-II; ZMIZ3

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

**ACCN:** NM\_016166 **ORF Size:** 1953 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC219594).

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:** <u>NM 016166.1</u>

 RefSeq Size:
 2309 bp

 RefSeq ORF:
 1956 bp

 Locus ID:
 8554

 UniProt ID:
 075925

 Cytogenetics:
 15q23

**Domains:** SAP, zf-MIZ

**Protein Families:** Druggable Genome, Transcription Factors





## PIAS1 (NM\_016166) Human Tagged ORF Clone Lentiviral Particle - RC219594L4V

**Protein Pathways:** Jak-STAT signaling pathway, Pathways in cancer, Small cell lung cancer, Ubiquitin mediated

proteolysis

**MW:** 71.7 kDa

**Gene Summary:** This gene encodes a member of the protein inhibitor of activated STAT (PIAS) family. PIAS

proteins function as SUMO E3 ligases and play important roles in many cellular processes by

mediating the sumoylation of target proteins. This protein plays a central role as a

transcriptional coregulator of numerous cellular pathways includign the STAT1 and nuclear factor kappaB pathways. Alternate splicing results in multiple transcript variants. [provided

by RefSeq, Mar 2016]