

## Product datasheet for RC220864L2V

## OriGene Technologies, Inc.

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## p73 (TP73) (NM\_005427) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** p73 (TP73) (NM\_005427) Human Tagged ORF Clone Lentiviral Particle

Symbol:p73Synonyms:P73

Mammalian Cell None

Selection:

NOTIC

**Vector:** pLenti-C-mGFP (PS100071)

Tag: mGFP

**ACCN:** NM\_005427 **ORF Size:** 1908 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC220864).

Sequence:

**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. 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 005427.1, NP 005418.1

 RefSeq Size:
 2234 bp

 RefSeq ORF:
 1911 bp

 Locus ID:
 7161

 UniProt ID:
 015350

 Cytogenetics:
 1p36.32

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

**Protein Pathways:** Neurotrophin signaling pathway, p53 signaling pathway





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**MW:** 69.4 kDa

**Gene Summary:** 

This gene encodes a member of the p53 family of transcription factors involved in cellular responses to stress and development. It maps to a region on chromosome 1p36 that is frequently deleted in neuroblastoma and other tumors, and thought to contain multiple tumor suppressor genes. The demonstration that this gene is monoallelically expressed (likely from the maternal allele), supports the notion that it is a candidate gene for neuroblastoma. Many transcript variants resulting from alternative splicing and/or use of alternate promoters have been found for this gene, but the biological validity and the full-length nature of some variants have not been determined. [provided by RefSeq, Feb 2011]