

## Product datasheet for RC201790L3V

## OriGene Technologies, Inc.

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## PSMC3 (NM\_002804) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

Product Type: Lentiviral Particles

Product Name: PSMC3 (NM 002804) Human Tagged ORF Clone Lentiviral Particle

Symbol: PSMC3

**Synonyms:** DCIDP; RPT5; TBP1

Mammalian Cell

Selection:

Puromycin

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

 Tag:
 Myc-DDK

 ACCN:
 NM\_002804

 ORF Size:
 1317 bp

**ORF Nucleotide** 

OTI Disclaimer:

cloatida The C

Sequence:

The ORF insert of this clone is exactly the same as (RC201790).

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 002804.4, NP 002795.2

 RefSeq Size:
 1618 bp

 RefSeq ORF:
 1320 bp

 Locus ID:
 5702

 UniProt ID:
 P17980

 Cytogenetics:
 11p11.2

Domains: AAA, AAA

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







**Protein Pathways:** Proteasome

RefSeq, Jul 2008]

**MW:** 49 kDa

**Gene Summary:** The 26S proteasome is a multicatalytic proteinase complex with a highly ordered structure

composed of 2 complexes, a 20S core and a 19S regulator. The 20S core is composed of 4 rings of 28 non-identical subunits; 2 rings are composed of 7 alpha subunits and 2 rings are composed of 7 beta subunits. The 19S regulator is composed of a base, which contains 6 ATPase subunits and 2 non-ATPase subunits, and a lid, which contains up to 10 non-ATPase subunits. Proteasomes are distributed throughout eukaryotic cells at a high concentration and cleave peptides in an ATP/ubiquitin-dependent process in a non-lysosomal pathway. An essential function of a modified proteasome, the immunoproteasome, is the processing of class I MHC peptides. This gene encodes one of the ATPase subunits, a member of the triple-A family of ATPases that have chaperone-like activity. This subunit may compete with PSMC2 for binding to the HIV tat protein to regulate the interaction between the viral protein and the transcription complex. A pseudogene has been identified on chromosome 9. [provided by

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