

## Product datasheet for RC230546L3V

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## Mps1 (TTK) (NM\_001166691) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** Mps1 (TTK) (NM\_001166691) Human Tagged ORF Clone Lentiviral Particle

Symbol: TTK

Synonyms: CT96; ESK; MPH1; MPS1; MPS1L1; PYT

**Mammalian Cell** 

Selection:

Puromycin

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

Tag: Myc-DDK

**ACCN:** NM\_001166691

ORF Size: 2568 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC230546).

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.

**RefSeq:** NM 001166691.1, NP 001160163.1

 RefSeq Size:
 3019 bp

 RefSeq ORF:
 2571 bp

 Locus ID:
 7272

 UniProt ID:
 P33981

 Cytogenetics:
 6q14.1

**Protein Families:** Druggable Genome, Protein Kinase



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Protein Pathways: Cell cycle, Oocyte meiosis, TGF-beta signaling pathway, Ubiquitin mediated proteolysis, Wnt

signaling pathway

**MW:** 96.9 kDa

**Gene Summary:** This gene encodes a dual specificity protein kinase with the ability to phosphorylate tyrosine,

serine and threonine. Associated with cell proliferation, this protein is essential for

chromosome alignment at the centromere during mitosis and is required for centrosome

duplication. It has been found to be a critical mitotic checkpoint protein for accurate segregation of chromosomes during mitosis. Tumorigenesis may occur when this protein fails

to degrade and produces excess centrosomes resulting in aberrant mitotic spindles.

Alternative splicing results in multiple transcript variants. [provided by RefSeq, Nov 2009]