

## Product datasheet for RC215782L3V

### OriGene Technologies, Inc.

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# KIF14 (NM\_014875) Human Tagged ORF Clone Lentiviral Particle

#### **Product data:**

**Product Type:** Lentiviral Particles

**Product Name:** KIF14 (NM\_014875) Human Tagged ORF Clone Lentiviral Particle

Symbol: KIF14

Synonyms: MCPH20; MKS12

Mammalian Cell

Selection:

Puromycin

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

Tag: Myc-DDK
ACCN: NM 014875

ORF Size: 4944 bp

**ORF Nucleotide** 

Sequence:

The ORF insert of this clone is exactly the same as(RC215782).

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 014875.2

 RefSeq Size:
 7293 bp

 RefSeq ORF:
 4947 bp

 Locus ID:
 9928

 UniProt ID:
 Q15058

 Cytogenetics:
 1q32.1

**Protein Families:** Druggable Genome

**MW:** 186.5 kDa







#### **Gene Summary:**

This gene encodes a member of the kinesin-3 superfamily of microtubule motor proteins. These proteins are involved in numerous processes including vesicle transport, chromosome segregation, mitotic spindle formation, and cytokinesis. In human HeLa-S3 and 293T cells, this protein is localized to the cytoplasm during interphase, to the spindle poles and spindle microtubules during mitosis, and to the midbody during cytokinesis. An internal motor domain displays microtubule-dependent ATPase activity, consistent with its function as a microtubule motor protein. Knockdown of this gene results in failed cytokinesis with endoreplication, which results in multinucleated cells. This gene has been identified as a likely oncogene in breast, lung and ovarian cancers, as well as retinoblastomas and gliomas. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2015]