

## Product datasheet for RC200064L1V

## OriGene Technologies, Inc.

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## ITGB3BP (NM\_014288) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** ITGB3BP (NM\_014288) Human Tagged ORF Clone Lentiviral Particle

Symbol: ITGB3BP

**Synonyms:** CENP-R; CENPR; HSU37139; NRIF3; TAP20

NM 014288

Mammalian Cell

Selection:

None

**Vector:** pLenti-C-Myc-DDK (PS100064)

Tag: Myc-DDK

ORF Size: 531 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC200064).

Sequence:

ACCN:

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 014288.3

 RefSeq Size:
 1019 bp

 RefSeq ORF:
 534 bp

 Locus ID:
 23421

 UniProt ID:
 Q13352

 Cytogenetics:
 1p31.3

**Protein Families:** Druggable Genome, Stem cell - Pluripotency, Transcription Factors

MW: 20.2 kDa







## **Gene Summary:**

This gene encodes a transcriptional coregulator that binds to and enhances the activity of members of the nuclear receptor families, thyroid hormone receptors and retinoid X receptors. This protein also acts as a corepressor of NF-kappaB-dependent signaling. This protein induces apoptosis in breast cancer cells through a caspase 2-mediated signaling pathway. This protein is also a component of the centromere-specific histone H3 variant nucleosome associated complex (CENP-NAC) and may be involved in mitotic progression by recruiting the histone H3 variant CENP-A to the centromere. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Sep 2011]