

## Product datasheet for RC231410L4V

## OriGene Technologies, Inc.

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## HIF1 beta (ARNT) (NM 001197325) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

Product Name: HIF1 beta (ARNT) (NM 001197325) Human Tagged ORF Clone Lentiviral Particle

Symbol: HIF1 beta

Synonyms: bHLHe2; HIF-1-beta; HIF-1beta; HIF1-beta; HIF1BETA; TANGO

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

**ACCN:** NM\_001197325

ORF Size: 2319 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC231410).

OTI Disclaimer:

Sequence:

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: <u>NM 001197325.1</u>

RefSeq ORF: 2322 bp Locus ID: 405

 UniProt ID:
 P27540

 Cytogenetics:
 1q21.3

Protein Families: Druggable Genome, Transcription Factors
Protein Pathways: Pathways in cancer, Renal cell carcinoma

**MW:** 85.3 kDa





## **Gene Summary:**

This gene encodes a protein containing a basic helix-loop-helix domain and two characteristic PAS domains along with a PAC domain. The encoded protein binds to ligand-bound aryl hydrocarbon receptor and aids in the movement of this complex to the nucleus, where it promotes the expression of genes involved in xenobiotic metabolism. This protein is also a co-factor for transcriptional regulation by hypoxia-inducible factor 1. Chromosomal translocation of this locus with the ETV6 (ets variant 6) gene on chromosome 12 have been described in leukemias. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Oct 2013]