

Product datasheet for RC216724L2V

OriGene Technologies, Inc.

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HIF1 beta (ARNT) (NM_001668) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: HIF1 beta (ARNT) (NM_001668) Human Tagged ORF Clone Lentiviral Particle

Symbol: HIF1 beta

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

Mammalian Cell

Selection:

None

Vector: pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_001668 **ORF Size:** 2367 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC216724).

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.

RefSeg: NM 001668.2

RefSeq Size: 4846 bp
RefSeq ORF: 2370 bp
Locus ID: 405

 UniProt ID:
 P27540

 Cytogenetics:
 1q21.3

Domains: PAS, HLH, PAC

Protein Families: Druggable Genome, Transcription Factors





Protein Pathways: Pathways in cancer, Renal cell carcinoma

MW: 86.5 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]