

## Product datasheet for RC223854L1V

## OriGene Technologies, Inc.

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## RUNX1 (NM\_001001890) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

Product Name: RUNX1 (NM\_001001890) Human Tagged ORF Clone Lentiviral Particle

Symbol: RUNX1

Synonyms: AML1; AML1-EVI-1; AMLCR1; CBF2alpha; CBFA2; EVI-1; PEBP2aB; PEBP2alpha

Mammalian Cell

Selection:

None

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

Tag: Myc-DDK

**ACCN:** NM\_001001890

ORF Size: 1359 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC223854).

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 001001890.1

RefSeq Size: 7288 bp RefSeq ORF: 1362 bp

Locus ID: 861

 UniProt ID:
 Q01196

 Cytogenetics:
 21q22.12

Protein Families: Druggable Genome, ES Cell Differentiation/IPS, Transcription Factors

**Protein Pathways:** Acute myeloid leukemia, Chronic myeloid leukemia, Pathways in cancer





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MW: 48.6 kDa

**Gene Summary:** Core binding factor (CBF) is a heterodimeric transcription factor that binds to the core

element of many enhancers and promoters. The protein encoded by this gene represents the

alpha subunit of CBF and is thought to be involved in the development of normal

hematopoiesis. Chromosomal translocations involving this gene are well-documented and have been associated with several types of leukemia. Three transcript variants encoding

different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]