

OriGene Technologies, Inc.

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Product datasheet for RC231324L4V

RUNX1T1 (NM_001198633) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type:	Lentiviral Particles
Product Name:	RUNX1T1 (NM_001198633) Human Tagged ORF Clone Lentiviral Particle
Symbol:	RUNX1T1
Synonyms:	AML1-MTG8; AML1T1; CBFA2T1; CDR; ETO; MTG8; ZMYND2
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-mGFP-P2A-Puro (PS100093)
Tag:	mGFP
ACCN:	NM_001198633
ORF Size:	1752 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC231324).
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. <u>More info</u>
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 001198633.1, NP 001185562.1</u>
RefSeq ORF:	1755 bp
Locus ID:	862
UniProt ID:	<u>Q06455</u>
Cytogenetics:	8q21.3
Protein Families:	Transcription Factors
Protein Pathways:	Acute myeloid leukemia, Pathways in cancer
MW:	65.5 kDa



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Gene Summary:This gene encodes a member of the myeloid translocation gene family which interact with
DNA-bound transcription factors and recruit a range of corepressors to facilitate
transcriptional repression. The t(8;21)(q22;q22) translocation is one of the most frequent
karyotypic abnormalities in acute myeloid leukemia. The translocation produces a chimeric
gene made up of the 5'-region of the runt-related transcription factor 1 gene fused to the 3'-
region of this gene. The chimeric protein is thought to associate with the nuclear
corepressor/histone deacetylase complex to block hematopoietic differentiation. Alternative
splicing results in multiple transcript variants. [provided by RefSeq, Nov 2010]

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