

## Product datasheet for RC202128L2V

## OriGene Technologies, Inc.

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## GTF2IRD1 (NM\_016328) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** GTF2IRD1 (NM\_016328) Human Tagged ORF Clone Lentiviral Particle

Symbol: GTF2IRD1

Synonyms: BEN; CREAM1; GTF3; hMusTRD1alpha1; MUSTRD1; RBAP2; WBS; WBSCR11; WBSCR12

Mammalian Cell

Selection:

None

**Vector:** pLenti-C-mGFP (PS100071)

Tag: mGFP

**ACCN:** NM\_016328 **ORF Size:** 2877 bp

**ORF Nucleotide** 

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Sequence:

The ORF insert of this clone is exactly the same as(RC202128).

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

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 016328.1

RefSeq Size: 3471 bp
RefSeq ORF: 2880 bp
Locus ID: 9569
UniProt ID: Q9UHL9

Cytogenetics: 7q11.23

Domains: GTF21

**Protein Families:** Druggable Genome, Transcription Factors





## GTF2IRD1 (NM\_016328) Human Tagged ORF Clone Lentiviral Particle - RC202128L2V

**Protein Pathways:** Basal transcription factors

**MW:** 106 kDa

**Gene Summary:** The protein encoded by this gene contains five GTF2I-like repeats and each repeat possesses

a potential helix-loop-helix (HLH) motif. It may have the ability to interact with other HLH-proteins and function as a transcription factor or as a positive transcriptional regulator under the control of Retinoblastoma protein. This gene plays a role in craniofacial and cognitive development and mutations have been associated with Williams-Beuren syndrome, a multisystem developmental disorder caused by deletion of multiple genes at 7q11.23. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Nov 2010]