

# **Product datasheet for SC335389**

## TAL1 (NM 001290403) Human Untagged Clone

#### **Product data:**

**Product Type:** Expression Plasmids

Product Name: TAL1 (NM\_001290403) Human Untagged Clone

Tag: Tag Free Symbol: TAL1

Synonyms: bHLHa17; SCL; tal-1; TCL5

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC335389 representing NM\_001290403.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

CCTGCCGCCGATGGAGCCGGCCCTCGGTGA

Restriction Sites: Sgfl-Mlul



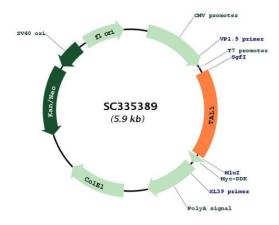
**OriGene Technologies, Inc.** 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



#### Plasmid Map:



**ACCN:** NM\_001290403

**Insert Size:** 996 bp

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

**Reconstitution Method:** 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

**RefSeq:** NM 001290403.1



### TAL1 (NM\_001290403) Human Untagged Clone - SC335389

 RefSeq Size:
 4774 bp

 RefSeq ORF:
 996 bp

 Locus ID:
 6886

 UniProt ID:
 P17542

 Cytogenetics:
 1p33

**Protein Families:** Adult stem cells, Druggable Genome, ES Cell Differentiation/IPS, Transcription Factors

MW: 34.3 kDa

**Gene Summary:** Implicated in the genesis of hemopoietic malignancies. It may play an important role in

hemopoietic differentiation. Serves as a positive regulator of erythroid differentiation (By

similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) lacks an internal exon in the 5' UTR, compared to variant 1. Variants 1, 2, 3, 4 and 5 encode the same isoform 1. Sequence Note: The RefSeq transcript and protein were derived from transcript and genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the

transcript record were based on alignments.