

## **Product datasheet for TL315481**

## OriGene Technologies, Inc.

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## **CREBL2 Human shRNA Plasmid Kit (Locus ID 1389)**

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** CREBL2 Human shRNA Plasmid Kit (Locus ID 1389)

**Locus ID:** 1389

**Vector:** pGFP-C-shLenti (TR30023)

E. coli Selection: Chloramphenicol (34 ug/ml)

Mammalian Cell

Selection:

Puromycin

Format: Lentiviral plasmids

Components: CREBL2 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 1389).

5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.

RefSeq: NM 001310, NM 001310.1, NM 001310.2, NM 001310.3, BC106052, BC106052.1, BC104873,

BC112157, NM 001310.4

UniProt ID: <u>060519</u>

Summary: cAMP response element (CRE)-binding protein-like-2 (CREBL2) was identified in a search to

find genes in a commonly deleted region on chromosome 12p13 flanked by ETV6 and CDKN1B genes, frequently associated with hematopoietic malignancies, as well as breast, non-small-cell lung and ovarian cancers. CREBL2 shares a 41% identity with CRE-binding protein (CREB) over a 48-base long region which encodes the bZip domain of CREB. The bZip domain consists of about 30 amino acids rich in basic residues involved in DNA binding, followed by a leucine zipper motif involved in protein dimerization. This suggests that CREBL2

encodes a protein with DNA binding capabilities. The occurance of CREBL2 deletion in

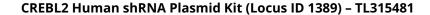
malignancy suggests that CREBL2 may act as a tumor suppressor gene. [provided by RefSeq,

Jul 2008]

**shRNA Design:** These shRNA constructs were designed against multiple splice variants at this gene locus. To

be certain that your variant of interest is targeted, please contact <u>techsupport@origene.com</u>. If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u>.







## Performance Guaranteed:

OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).