## **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on May 10, 2025

# pCRISPRia-v2

RRID:Addgene\_84832 Type: Plasmid

#### **Proper Citation**

RRID:Addgene\_84832

#### **Plasmid Information**

URL: http://www.addgene.org/84832

Proper Citation: RRID:Addgene\_84832

Insert Name: sgRNA

Bacterial Resistance: Ampicillin

Defining Citation: PMID:27661255

**Vector Backbone Description:** Backbone Marker:Addgene; Backbone Size:8200; Vector Backbone:pSICO derivative; Vector Types:Mammalian Expression, Lentiviral, CRISPR; Bacterial Resistance:Ampicillin

Plasmid Name: pCRISPRia-v2

**Record Creation Time:** 20220422T222546+0000

Record Last Update: 20231115T081052+0000

### **Ratings and Alerts**

No rating or validation information has been found for pCRISPRia-v2.

No alerts have been found for pCRISPRia-v2.

Data and Source Information

#### **Usage and Citation Metrics**

We found 30 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Khoroshkin M, et al. (2024) Systematic identification of post-transcriptional regulatory modules. Nature communications, 15(1), 7872.

Yang Y, et al. (2024) Large tandem duplications in cancer result from transcription and DNA replication collisions. medRxiv : the preprint server for health sciences.

Fernandez SG, et al. (2024) Ribosome rescue factor PELOTA modulates translation start site choice for C/EBP? protein isoforms. Life science alliance, 7(7).

Umhoefer JM, et al. (2024) Deciphering regulation of FOXP3 expression in human conventional T cells. bioRxiv : the preprint server for biology.

Ivancevic A, et al. (2024) Endogenous retroviruses mediate transcriptional rewiring in response to oncogenic signaling in colorectal cancer. Science advances, 10(29), eado1218.

Wu R, et al. (2024) Disruption of nuclear speckle integrity dysregulates RNA splicing in C9ORF72-FTD/ALS. Neuron, 112(20), 3434.

Zhao Y, et al. (2024) Long noncoding RNA Malat1 protects against osteoporosis and bone metastasis. Nature communications, 15(1), 2384.

Villagomez FR, et al. (2024) Claudin-4 stabilizes the genome via nuclear and cell cycle remodeling to support ovarian cancer cell survival. Cancer research communications.

Aiken J, et al. (2024) Spastin locally amplifies microtubule dynamics to pattern the axon for presynaptic cargo delivery. Current biology : CB, 34(8), 1687.

Zirak B, et al. (2024) Revealing the grammar of small RNA secretion using interpretable machine learning. Cell genomics, 4(4), 100522.

Mowery CT, et al. (2024) Systematic decoding of cis gene regulation defines contextdependent control of the multi-gene costimulatory receptor locus in human T cells. Nature genetics, 56(6), 1156.

Lai JD, et al. (2024) KCNJ2 inhibition mitigates mechanical injury in a human brain organoid model of traumatic brain injury. Cell stem cell, 31(4), 519.

Chakrabarty Y, et al. (2024) The HRI branch of the integrated stress response selectively triggers mitophagy. Molecular cell, 84(6), 1090.

Hazan JM, et al. (2024) Integration of transcription regulation and functional genomic data reveals IncRNA SNHG6's role in hematopoietic differentiation and leukemia. Journal of biomedical science, 31(1), 27.

Luteijn RD, et al. (2024) The activation of the adaptor protein STING depends on its interactions with the phospholipid PI4P. Science signaling, 17(827), eade3643.

Jones-Tabah J, et al. (2024) The Parkinson's disease risk gene cathepsin B promotes fibrillar alpha-synuclein clearance, lysosomal function and glucocerebrosidase activity in dopaminergic neurons. Research square.

Vasudevan HN, et al. (2024) Functional interactions between neurofibromatosis tumor suppressors underlie Schwann cell tumor de-differentiation and treatment resistance. Nature communications, 15(1), 477.

Khoroshkin M, et al. (2024) A systematic search for RNA structural switches across the human transcriptome. Nature methods, 21(9), 1634.

Cheng Y, et al. (2024) A non-canonical role for a small nucleolar RNA in ribosome biogenesis and senescence. Cell, 187(17), 4770.

Sheedy CJ, et al. (2024) PEX1G843D remains functional in peroxisome biogenesis but is rapidly degraded by the proteasome. bioRxiv : the preprint server for biology.