Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 15, 2025

hCRISPRi-v2, genome-wide, top 5 sgRNAs/gene

RRID:Addgene_83969

Type: Plasmid

Proper Citation

RRID:Addgene_83969

Plasmid Information

URL: http://www.addgene.org/83969

Proper Citation: RRID:Addgene_83969

Bacterial Resistance: Ampicillin

Defining Citation: PMID:27661255

Vector Backbone Description: Vector Backbone:pCRISPRia-v2; Vector Types:Lentiviral;

Bacterial Resistance: Ampicillin

Plasmid Name: hCRISPRi-v2, genome-wide, top 5 sgRNAs/gene

Record Creation Time: 20220422T222542+0000

Record Last Update: 20220422T225135+0000

Ratings and Alerts

No rating or validation information has been found for hCRISPRi-v2, genome-wide, top 5 sgRNAs/gene.

No alerts have been found for hCRISPRi-v2, genome-wide, top 5 sgRNAs/gene.

Data and Source Information

Source: Addgene

Usage and Citation Metrics

We found 13 mentions in open access literature.

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

Vu JT, et al. (2024) A genome-wide screen links peroxisome regulation with Wnt signaling through RNF146 and tankyrase. bioRxiv: the preprint server for biology.

Muthukumar G, et al. (2024) Triaging of ?-helical proteins to the mitochondrial outer membrane by distinct chaperone machinery based on substrate topology. Molecular cell, 84(6), 1101.

Choi S, et al. (2024) The Role of IRF9 Upregulation in Modulating Sensitivity to Olaparib and Platinum-Based Chemotherapies in Breast Cancer. Genes, 15(7).

Heo SJ, et al. (2024) Compact CRISPR genetic screens enabled by improved guide RNA library cloning. Genome biology, 25(1), 25.

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

Arriaga JM, et al. (2024) In vivo genome-wide CRISPR screening identifies CITED2 as a driver of prostate cancer bone metastasis. Oncogene, 43(17), 1303.

Carnie CJ, et al. (2024) Transcription-coupled repair of DNA-protein cross-links depends on CSA and CSB. Nature cell biology, 26(5), 797.

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

Zelceski A, et al. (2023) MND1 and PSMC3IP control PARP inhibitor sensitivity in mitotic cells. Cell reports, 42(5), 112484.

Wei TT, et al. (2022) Cannabinoid receptor 1 antagonist genistein attenuates marijuana-induced vascular inflammation. Cell, 185(10), 1676.

Coukos R, et al. (2021) An engineered transcriptional reporter of protein localization identifies regulators of mitochondrial and ER membrane protein trafficking in high-throughput CRISPRi screens. eLife, 10.

Palmer AC, et al. (2019) A curative combination cancer therapy achieves high fractional cell killing through low cross-resistance and drug additivity. eLife, 8.

Tesar PJ, et al. (2007) New cell lines from mouse epiblast share defining features with human embryonic stem cells. Nature, 448(7150), 196.