

# Resource Summary Report

Generated by [RRID](#) on Aug 23, 2026

## pHRdSV40-scFv-GCN4-sfGFP-VP64-GB1-NLS

RRID:Addgene\_60904

Type: Plasmid

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### Proper Citation

RRID:Addgene\_60904

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### Plasmid Information

**URL:** <http://www.addgene.org/60904>

**Proper Citation:** RRID:Addgene\_60904

**Insert Name:** scFv-GCN4

**Bacterial Resistance:** Ampicillin

**Defining Citation:** [PMID:25307933](#)

**Vector Backbone Description:** Vector Backbone:pHR; Vector Types:Mammalian Expression, Lentiviral; Bacterial Resistance:Ampicillin

**Comments:** For more information, visit <https://valelab4.ucsf.edu/external/research/suntag.html> Note this plasmid was used with the CRISPRa library, as described in Gilbert et al. Cell 2014 PubMed ID: 25307932.

**Plasmid Name:** pHRdSV40-scFv-GCN4-sfGFP-VP64-GB1-NLS

**Record Creation Time:** 20220422T222350+0000

**Record Last Update:** 20260815T081737+0000

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### Ratings and Alerts

No rating or validation information has been found for pHRdSV40-scFv-GCN4-sfGFP-VP64-GB1-NLS.

No alerts have been found for pHRdSV40-scFv-GCN4-sfGFP-VP64-GB1-NLS.

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### Data and Source Information

**Source:** [Addgene](#)

## Usage and Citation Metrics

We found 59 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](#) .

Shi L, et al. (2026) Mapping kinase-dependent tumor immune adaptation with multiplexed single-cell CRISPR screens. *bioRxiv : the preprint server for biology*.

Dutta S, et al. (2026) In vivo CRISPR-mediated activation of cardiogenic genes to reprogram cardiac fibroblasts. *Frontiers in cell and developmental biology*, 14, 1812941.

Xiao N, et al. (2025) Type A cholesterol-dependent cytolysins translocate to the trans-Golgi network for NLRP3 inflammasome activation. *Nature immunology*, 26(10), 1673.

Liu J, et al. (2025) SunTag-PE: a modular prime editing system enables versatile and efficient genome editing. *Communications biology*, 8(1), 452.

Zhao T, et al. (2025) Derepressing nuclear pyruvate dehydrogenase induces therapeutic cancer cell reprogramming. *Cell metabolism*, 37(8), 1667.

Liu J, et al. (2024) Involvement of ribosomal protein L17 and Y-box binding protein 1 in the assembly of hepatitis C virus potentially via their interaction with the 3' untranslated region of the viral genome. *Journal of virology*, 98(7), e0052224.

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.

Mu S, et al. (2024) Enhancing prime editor flexibility with coiled-coil heterodimers. *Genome biology*, 25(1), 108.

Conger KO, et al. (2024) ASCT2 is a major contributor to serine uptake in cancer cells. *Cell reports*, 43(8), 114552.

McFaline-Figueroa JL, et al. (2024) Multiplex single-cell chemical genomics reveals the kinase dependence of the response to targeted therapy. *Cell genomics*, 4(2), 100487.

Jung P, et al. (2023) CRISPR Activation/Interference Screen to Identify Genetic Networks in HDAC-Inhibitor-Resistant Cells. *Methods in molecular biology (Clifton, N.J.)*, 2589, 429.

Peng H, et al. (2023) Epigenome-wide association study identifies novel genes associated with ischemic stroke. *Clinical epigenetics*, 15(1), 106.

Conger KO, et al. (2023) ASCT2 is the primary serine transporter in cancer cells. *bioRxiv : the preprint server for biology*.

McFaline-Figueroa JL, et al. (2023) Multiplex single-cell chemical genomics reveals the kinase dependence of the response to targeted therapy. *bioRxiv : the preprint server for biology*.

Liu X, et al. (2023) IMPDH inhibition activates TLR-VCAM1 pathway and suppresses the development of MLL-fusion leukemia. *EMBO molecular medicine*, 15(1), e15631.

Yabushita T, et al. (2023) Mitotic perturbation is a key mechanism of action of decitabine in myeloid tumor treatment. *Cell reports*, 42(9), 113098.

Liu X, et al. (2022) G-quadruplex-guided RNA engineering to modulate CRISPR-based genomic regulation. *Nucleic acids research*, 50(19), 11387.

Wood RK, et al. (2022) Secretory defects in pediatric osteosarcoma result from downregulation of selective COPII coatomer proteins. *iScience*, 25(4), 104100.

Danziger O, et al. (2022) Inducible CRISPR activation screen for interferon-stimulated genes identifies OAS1 as a SARS-CoV-2 restriction factor. *PLoS pathogens*, 18(4), e1010464.

Manj??n AG, et al. (2022) Unexpected gene activation following CRISPR-Cas9-mediated genome editing. *EMBO reports*, 23(2), e53902.