# Resource Summary Report

Generated by dkNET on Sep 8, 2026

## pRDA_052

RRID:Addgene_136474
Type: Plasmid

### Proper Citation

RRID:Addgene_136474

### Plasmid Information

**URL:** http://www.addgene.org/136474

**Proper Citation:** RRID:Addgene_136474

**Insert Name:** Puromycin resistance

**Organism:** Other

**Bacterial Resistance:** Ampicillin

**Defining Citation:** PMID:32661438

**Vector Backbone Description:** Vector Backbone:pLKO; Vector Types:Lentiviral, Synthetic Biology; Bacterial Resistance:Ampicillin

**Plasmid Name:** pRDA_052

**Record Creation Time:** 20220422T221752+0000

**Record Last Update:** 20260905T074819+0000

### Ratings and Alerts

No rating or validation information has been found for pRDA_052.

No alerts have been found for pRDA_052.

### Data and Source Information

**Source:** Addgene

### Usage and Citation Metrics

We found 7 mentions in open access literature.

**Listed below are recent publications.** The full list is available at dkNET .

Wolf G, et al. (2025) The genetic interaction map of the human solute carrier superfamily. Molecular systems biology, 21(6), 531.

Borck PC, et al. (2025) SKI complex loss renders 9p21.3-deleted or MSI-H cancers dependent on PELO. Nature, 638(8052), 1104.

Melore SM, et al. (2024) HyperCas12a enables highly-multiplexed epigenome editing screens. bioRxiv : the preprint server for biology.

Cetin R, et al. (2023) Optimized metrics for orthogonal combinatorial CRISPR screens. Scientific reports, 13(1), 7405.

Rebendenne A, et al. (2022) Bidirectional genome-wide CRISPR screens reveal host factors regulating SARS-CoV-2, MERS-CoV and seasonal HCoVs. Nature genetics, 54(8), 1090.

Li R, et al. (2022) Comparative optimization of combinatorial CRISPR screens. Nature communications, 13(1), 2469.

Lenoir WF, et al. (2021) Discovery of putative tumor suppressors from CRISPR screens reveals rewired lipid metabolism in acute myeloid leukemia cells. Nature communications, 12(1), 6506.