

Resource Summary Report

Generated by [dkNET](#) on Aug 29, 2026

pRDA_479

RRID:Addgene_179099

Type: Plasmid

Proper Citation

RRID:Addgene_179099

Plasmid Information

URL: <http://www.addgene.org/179099>

Proper Citation: RRID:Addgene_179099

Insert Name: ABE8e (SpG)

Organism: Synthetic

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:35288574](#)

Vector Backbone Description: Vector Backbone:pRDA_426; Vector Types:Lentiviral, CRISPR; Bacterial Resistance:Ampicillin

Plasmid Name: pRDA_479

Record Creation Time: 20220422T222030+0000

Record Last Update: 20260829T080111+0000

Ratings and Alerts

No rating or validation information has been found for pRDA_479.

No alerts have been found for pRDA_479.

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](#) .

Greenwood E, et al. (2025) Haplotype rather than single causal variants effects contribute to regulatory gene expression associations in human myeloid cells. bioRxiv : the preprint server for biology.

Merz ML, et al. (2025) Orthogonal resistance mechanisms of classical- and induced-proximity inhibitors. bioRxiv : the preprint server for biology.

Sokirniy I, et al. (2025) A side-by-side comparison of variant function measurements using deep mutational scanning and base editing. Nucleic acids research, 53(14).

Meeusen B, et al. (2025) A functional map of phosphoprotein phosphatase regulation identifies an evolutionary conserved reductase for the catalytic metal ions. bioRxiv : the preprint server for biology.

Ingham A, et al. (2025) CRAMP1-dependent histone H1 biogenesis is essential for topoisomerase II inhibitor tolerance. Molecular cell, 85(13), 2487.

Yeo MJR, et al. (2025) UM171 glues asymmetric CRL3-HDAC1/2 assembly to degrade CoREST corepressors. Nature, 639(8053), 232.

Sangree AK, et al. (2022) Benchmarking of SpCas9 variants enables deeper base editor screens of BRCA1 and BCL2. Nature communications, 13(1), 1318.