

# Resource Summary Report

Generated by [dkNET](#) on Aug 20, 2026

## [pYPQ265C](#)

RRID:Addgene\_164715

Type: Plasmid

### Proper Citation

RRID:Addgene\_164715

### Plasmid Information

**URL:** <http://www.addgene.org/164715>

**Proper Citation:** RRID:Addgene\_164715

**Insert Name:** hAID-zCas9(D10A)-UGI

**Bacterial Resistance:** Spectinomycin

**Defining Citation:** [PMID:34042262](#)

**Vector Backbone Description:** Vector Backbone:pYPQ255; Vector Types:Plant Expression, CRISPR, Gateway compatible hAID-CBE\_V01 entry clone; Bacterial Resistance:Spectinomycin

**Plasmid Name:** pYPQ265C

**Record Creation Time:** 20220422T221930+0000

**Record Last Update:** 20260815T080923+0000

### Ratings and Alerts

No rating or validation information has been found for pYPQ265C.

No alerts have been found for pYPQ265C.

### Data and Source Information

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

### Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Jeong YY, et al. (2025) Identification of optimal adenine and cytosine base editors for genome editing in Arabidopsis and soybean. BMB reports, 58(7), 288.

Randall LB, et al. (2021) Genome- and transcriptome-wide off-target analyses of an improved cytosine base editor. Plant physiology, 187(1), 73.