

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](http://FDI Lab - SciCrunch.org) on Apr 14, 2025

## p2T-CMV-AIDmax-BlastR

RRID:Addgene\_152994

Type: Plasmid

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### Proper Citation

RRID:Addgene\_152994

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### Plasmid Information

**URL:** <http://www.addgene.org/152994>

**Proper Citation:** RRID:Addgene\_152994

**Insert Name:** AID-nSpCas9-UGI-UGI

**Organism:** Synthetic

**Bacterial Resistance:** Ampicillin

**Defining Citation:** [PMID:32533916](https://pubmed.ncbi.nlm.nih.gov/32533916/)

**Vector Backbone Description:** Vector Backbone:p2T-CMV-BlastR; Vector Types:Mammalian Expression, CRISPR; Bacterial Resistance:Ampicillin

**Plasmid Name:** p2T-CMV-AIDmax-BlastR

**Record Creation Time:** 20220422T221837+0000

**Record Last Update:** 20220422T222816+0000

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### Ratings and Alerts

No rating or validation information has been found for p2T-CMV-AIDmax-BlastR.

No alerts have been found for p2T-CMV-AIDmax-BlastR.

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### Data and Source Information

Source: [Addgene](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](#).

Ryu J, et al. (2024) Joint genotypic and phenotypic outcome modeling improves base editing variant effect quantification. *Nature genetics*, 56(5), 925.