

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 7, 2025

pCMV_AncBE4max

RRID:Addgene_112094

Type: Plasmid

Proper Citation

RRID:Addgene_112094

Plasmid Information

URL: <http://www.addgene.org/112094>

Proper Citation: RRID:Addgene_112094

Insert Name: AncBE4max

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:29813047](https://pubmed.ncbi.nlm.nih.gov/29813047/)

Vector Backbone Description: Vector Backbone:pCMV; Vector Types:Mammalian Expression, CRISPR; Bacterial Resistance:Ampicillin

Plasmid Name: pCMV_AncBE4max

Record Creation Time: 20220422T221546+0000

Record Last Update: 20220422T221759+0000

Ratings and Alerts

No rating or validation information has been found for pCMV_AncBE4max.

No alerts have been found for pCMV_AncBE4max.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

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

Zhang G, et al. (2024) nCas9 Engineering for Improved Target Interaction Presents an Effective Strategy to Enhance Base Editing. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 11(31), e2405426.

Park JC, et al. (2024) Enhancing genome editing in hPSCs through dual inhibition of DNA damage response and repair pathways. *Nature communications*, 15(1), 4002.

Wang L, et al. (2024) MYH7 R453C induced cardiac remodelling via activating TGF- β /Smad2/3, ERK1/2 and Nox4/ROS/NF- κ B signalling pathways. *Open biology*, 14(6), 230427.

An M, et al. (2024) Systematic identification of pathogenic variants of non-small cell lung cancer in the promoters of DNA-damage repair genes. *EBioMedicine*, 110, 105480.

Deforz E, et al. (2023) HOXDeRNA activates a cancerous transcription program and super-enhancers genome-wide. *bioRxiv : the preprint server for biology*.

Cornean A, et al. (2022) Precise in vivo functional analysis of DNA variants with base editing using ACEofBASEs target prediction. *eLife*, 11.

Rosello M, et al. (2021) Precise base editing for the in vivo study of developmental signaling and human pathologies in zebrafish. *eLife*, 10.

Wang X, et al. (2020) Cas12a Base Editors Induce Efficient and Specific Editing with Low DNA Damage Response. *Cell reports*, 31(9), 107723.