

Resource Summary Report

Generated by [nidm-terms](#) on Aug 11, 2026

[pH2rU3_ForInd_Omicron_sinobiological_Spiked21_T7_CMV_ZsGT2APurR](#)

RRID:Addgene_204146

Type: Plasmid

Proper Citation

RRID:Addgene_204146

Plasmid Information

URL: <http://www.addgene.org/204146>

Proper Citation: RRID:Addgene_204146

Insert Name: SARS-CoV-2 BA.1 variant spike

Organism: SARS-CoV-2

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:36868218](#)

Vector Backbone Description: Vector Backbone:pH2rU3; Vector Types:Mammalian Expression, Lentiviral; Bacterial Resistance:Ampicillin

Plasmid Name:

pH2rU3_ForInd_Omicron_sinobiological_Spiked21_T7_CMV_ZsGT2APurR

Relevant Mutation: 21 amino acid cytoplasmic tail deletion

Record Creation Time: 20230818T080452+0000

Record Last Update: 20260808T082849+0000

Ratings and Alerts

No rating or validation information has been found for
pH2rU3_ForInd_Omicron_sinobiological_Spiked21_T7_CMV_ZsGT2APurR.

No alerts have been found for
pH2rU3_ForInd_Omicron_sinobiological_Spiked21_T7_CMV_ZsGT2APurR.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Carr CR, et al. (2024) Deep mutational scanning reveals functional constraints and antigenic variability of Lassa virus glycoprotein complex. bioRxiv : the preprint server for biology.