

# Resource Summary Report

Generated by [RRID](#) on Aug 18, 2026

## pcDNA3.3\_SARS2\_XBB.1.16

RRID:Addgene\_201189

Type: Plasmid

### Proper Citation

RRID:Addgene\_201189

### Plasmid Information

**URL:** <http://www.addgene.org/201189>

**Proper Citation:** RRID:Addgene\_201189

**Insert Name:** Spike (XBB.1.16)

**Organism:** SARS-CoV-2

**Bacterial Resistance:** Ampicillin

**Vector Backbone Description:** Backbone Size:5500; Vector Backbone:pcDNA3.3;  
Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

**Plasmid Name:** pcDNA3.3\_SARS2\_XBB.1.16

**Relevant Mutation:** E180V, T(K)478R on the basis of XBB.1.5

**Record Creation Time:** 20230524T080411+0000

**Record Last Update:** 20260815T082632+0000

### Ratings and Alerts

No rating or validation information has been found for pcDNA3.3\_SARS2\_XBB.1.16.

No alerts have been found for pcDNA3.3\_SARS2\_XBB.1.16.

### 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 [RRID](#) .

Gonzales J, et al. (2025) Development of an ultrahigh affinity, trimeric ACE2 biologic as a universal SARS-CoV-2 antagonist. Communications biology, 8(1), 1428.