

Resource Summary Report

Generated by [RRID](#) on Aug 21, 2026

pSBbi-RB

RRID:Addgene_60522

Type: Plasmid

Proper Citation

RRID:Addgene_60522

Plasmid Information

URL: <http://www.addgene.org/60522>

Proper Citation: RRID:Addgene_60522

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:25650551](#)

Vector Backbone Description: Backbone Size:2021; Vector Backbone:pUC19; Vector Types:Mammalian Expression, Transposon; Bacterial Resistance:Ampicillin

Plasmid Name: pSBbi-RB

Record Creation Time: 20220422T222348+0000

Record Last Update: 20260815T081733+0000

Ratings and Alerts

No rating or validation information has been found for pSBbi-RB.

No alerts have been found for pSBbi-RB.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Catanzaro NJ, et al. (2025) ACE2 from *Pipistrellus abramus* bats is a receptor for HKU5 coronaviruses. *Nature communications*, 16(1), 4932.

Streb P, et al. (2023) How chromosomal translocations arise to cause cancer: Gene proximity, trans-splicing, and DNA end joining. *iScience*, 26(6), 106900.

Bednash JS, et al. (2023) Inhibiting the Deubiquitinase UCHL1 Reduces SARS-CoV-2 Viral Uptake by ACE2. *American journal of respiratory cell and molecular biology*, 68(5), 566.

Jarman EJ, et al. (2023) DKK1 drives immune suppressive phenotypes in intrahepatic cholangiocarcinoma and can be targeted with anti-DKK1 therapeutic DKN-01. *Liver international : official journal of the International Association for the Study of the Liver*, 43(1), 208.

Tsai M, et al. (2023) Influenza virus reduces ubiquitin E3 ligase MARCH10 expression to decrease ciliary beat frequency. *American journal of physiology. Lung cellular and molecular physiology*, 324(5), L666.

Johnson BS, et al. (2022) MicroID2: A Novel Biotin Ligase Enables Rapid Proximity-Dependent Proteomics. *Molecular & cellular proteomics : MCP*, 21(7), 100256.

Erwood S, et al. (2022) Saturation variant interpretation using CRISPR prime editing. *Nature biotechnology*, 40(6), 885.

Lee DH, et al. (2021) Genome wide CRISPR screening reveals a role for sialylation in the tumorigenesis and chemoresistance of acute myeloid leukemia cells. *Cancer letters*, 510, 37.

Jin HS, et al. (2020) CD226hiCD8+ T Cells Are a Prerequisite for Anti-TIGIT Immunotherapy. *Cancer immunology research*, 8(7), 912.

Kim D, et al. (2019) Molecular mechanism for the multiple sclerosis risk variant rs17594362. *Human molecular genetics*, 28(21), 3600.