

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

pBABE-neo largeTgenomic

RRID:Addgene_10891

Type: Plasmid

Proper Citation

RRID:Addgene_10891

Plasmid Information

URL: <http://www.addgene.org/10891>

Proper Citation: RRID:Addgene_10891

Insert Name: large T and small T antigen

Organism: simian virus

Bacterial Resistance: Ampicillin

Vector Backbone Description: Backbone Size:4800; Vector Backbone:pBABE-neo; Vector Types:Mammalian Expression, Retroviral; Bacterial Resistance:Ampicillin

Plasmid Name: pBABE-neo largeTgenomic

Record Creation Time: 20220422T221528+0000

Record Last Update: 20260808T080413+0000

Ratings and Alerts

No rating or validation information has been found for pBABE-neo largeTgenomic.

No alerts have been found for pBABE-neo largeTgenomic.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Moquin-Beaudry G, et al. (2022) Autologous humanized mouse models of iPSC-derived tumors enable characterization and modulation of cancer-immune cell interactions. *Cell reports methods*, 2(1), 100153.

Moquin-Beaudry G, et al. (2019) The Tumor-Immune Response Is Not Compromised by Mesenchymal Stromal Cells in Humanized Mice. *Journal of immunology (Baltimore, Md. : 1950)*, 203(10), 2735.

Majerska J, et al. (2018) Transformation-induced stress at telomeres is counteracted through changes in the telomeric proteome including SAMHD1. *Life science alliance*, 1(4), e201800121.

Sousa CM, et al. (2016) Pancreatic stellate cells support tumour metabolism through autophagic alanine secretion. *Nature*, 536(7617), 479.

Cho S, et al. (2015) Design and Characterization of Bioengineered Cancer-Like Stem Cells. *PloS one*, 10(10), e0141172.

DeAngelis JT, et al. (2011) 2D difference gel electrophoresis analysis of different time points during the course of neoplastic transformation of human mammary epithelial cells. *Journal of proteome research*, 10(2), 447.