

Resource Summary Report

Generated by [RRID](#) on Aug 19, 2026

pcDNA3.1 GW-NL-myc

RRID:Addgene_113447

Type: Plasmid

Proper Citation

RRID:Addgene_113447

Plasmid Information

URL: <http://www.addgene.org/113447>

Proper Citation: RRID:Addgene_113447

Bacterial Resistance: Chloramphenicol and Ampicillin

Defining Citation: [PMID:29997244](https://pubmed.ncbi.nlm.nih.gov/29997244/)

Vector Backbone Description: Backbone Size:5428; Vector Backbone:pcDNA3.1; Vector Types:Mammalian Expression, Luciferase, Gateway shuttle vector; Bacterial Resistance:Chloramphenicol and Ampicillin

Plasmid Name: pcDNA3.1 GW-NL-myc

Record Creation Time: 20220422T221554+0000

Record Last Update: 20260815T080212+0000

Ratings and Alerts

No rating or validation information has been found for pcDNA3.1 GW-NL-myc.

No alerts have been found for pcDNA3.1 GW-NL-myc.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Trepte P, et al. (2024) AI-guided pipeline for protein-protein interaction drug discovery identifies a SARS-CoV-2 inhibitor. *Molecular systems biology*, 20(4), 428.

Brenner D, et al. (2024) A TBK1 variant causes autophagolysosomal and motoneuron pathology without neuroinflammation in mice. *The Journal of experimental medicine*, 221(5).

Lee CY, et al. (2024) Systematic discovery of protein interaction interfaces using AlphaFold and experimental validation. *Molecular systems biology*, 20(2), 75.

Trepte P, et al. (2023) AI-guided pipeline for protein-protein interaction drug discovery identifies a SARS-CoV-2 inhibitor. *bioRxiv : the preprint server for biology*.

Greco TM, et al. (2022) Dynamics of huntingtin protein interactions in the striatum identifies candidate modifiers of Huntington disease. *Cell systems*, 13(4), 304.