

Resource Summary Report

Generated by [NIF](#) on Aug 15, 2026

MadMyc-HA

RRID:Addgene_16557

Type: Plasmid

Proper Citation

RRID:Addgene_16557

Plasmid Information

URL: <http://www.addgene.org/16557>

Proper Citation: RRID:Addgene_16557

Insert Name: MADMYC

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:10688915](#)

Vector Backbone Description: Backbone Size:0; Vector Backbone:n/a; Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

Comments: MADMYC: the amino terminal transactivation domain of c-Myc (aa 1-263) was removed and replaced with the amino terminal transcriptional repression domain (aa 1-38, the Sin3-interaction region) of Mad.

Plasmid Name: MadMyc-HA

Relevant Mutation: Dominant negative mutant

Record Creation Time: 20220422T221934+0000

Record Last Update: 20260815T080930+0000

Ratings and Alerts

No rating or validation information has been found for MadMyc-HA.

No alerts have been found for MadMyc-HA.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Xin B, et al. (2017) Critical role of Myc activation in mouse hepatocarcinogenesis induced by the activation of AKT and RAS pathways. *Oncogene*, 36(36), 5087.

Nakajima-Koyama M, et al. (2015) Induction of Pluripotency in Astrocytes through a Neural Stem Cell-like State. *The Journal of biological chemistry*, 290(52), 31173.