

# Resource Summary Report

Generated by [RRID](#) on Aug 23, 2026

## MadMyc-HA

RRID:Addgene\_16557

Type: Plasmid

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### Proper Citation

RRID:Addgene\_16557

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### 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

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### Ratings and Alerts

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

No alerts have been found for MadMyc-HA.

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### Data and Source Information

Source: [Addgene](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.