

Resource Summary Report

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

MAC-tag-N

RRID:Addgene_108078

Type: Plasmid

Proper Citation

RRID:Addgene_108078

Plasmid Information

URL: <http://www.addgene.org/108078>

Proper Citation: RRID:Addgene_108078

Insert Name: CCDB

Organism: Other

Bacterial Resistance: Chloramphenicol and Ampicillin

Defining Citation: [PMID:29568061](#)

Vector Backbone Description: Vector Backbone:pcDNA5/FRT/TO; Vector Types:Mammalian Expression, Bacterial Expression; Bacterial Resistance:Chloramphenicol and Ampicillin

Plasmid Name: MAC-tag-N

Record Creation Time: 20220422T221524+0000

Record Last Update: 20260815T080116+0000

Ratings and Alerts

No rating or validation information has been found for MAC-tag-N.

No alerts have been found for MAC-tag-N.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Novosolova N, et al. (2025) Oxidative modulation of Piezo1 channels. *Redox biology*, 86, 103797.

Norppa AJ, et al. (2024) Distinct functions for the paralogous RBM41 and U11/U12-65K proteins in the minor spliceosome. *Nucleic acids research*, 52(7), 4037.

Gawriyski L, et al. (2024) Interaction network of human early embryonic transcription factors. *EMBO reports*, 25(3), 1589.

Nguyen-Dien GT, et al. (2023) FBXL4 suppresses mitophagy by restricting the accumulation of NIX and BNIP3 mitophagy receptors. *The EMBO journal*, 42(13), e112767.

Kinnunen M, et al. (2023) The Impact of ETV6-NTRK3 Oncogenic Gene Fusions on Molecular and Signaling Pathway Alterations. *Cancers*, 15(17).

Covelo-Molares H, et al. (2021) The comprehensive interactomes of human adenosine RNA methyltransferases and demethylases reveal distinct functional and regulatory features. *Nucleic acids research*, 49(19), 10895.

Liu X, et al. (2021) SARS-CoV-2-host proteome interactions for antiviral drug discovery. *Molecular systems biology*, 17(11), e10396.

Liu X, et al. (2020) Combined proximity labeling and affinity purification-mass spectrometry workflow for mapping and visualizing protein interaction networks. *Nature protocols*, 15(10), 3182.

Zhao Q, et al. (2018) Structure and mechanogating mechanism of the Piezo1 channel. *Nature*, 554(7693), 487.