

Resource Summary Report

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

MAC-tag-C

RRID:Addgene_108077

Type: Plasmid

Proper Citation

RRID:Addgene_108077

Plasmid Information

URL: <http://www.addgene.org/108077>

Proper Citation: RRID:Addgene_108077

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-C

Record Creation Time: 20220422T221524+0000

Record Last Update: 20260815T080116+0000

Ratings and Alerts

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

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

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Sottnik JL, et al. (2024) WNT4 Regulates Cellular Metabolism via Intracellular Activity at the Mitochondria in Breast and Gynecologic Cancers. *Cancer research communications*, 4(1), 134.

Turkki P, et al. (2024) Tensin-2 interactomics reveals interaction with GAPDH and a phosphorylation-mediated regulatory role in glycolysis. *Scientific reports*, 14(1), 19862.

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

Gawriyski L, et al. (2023) Comprehensive characterization of the embryonic factor LEUTX. *iScience*, 26(3), 106172.

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

Vuoristo S, et al. (2022) DUX4 is a multifunctional factor priming human embryonic genome activation. *iScience*, 25(4), 104137.

Salokas K, et al. (2022) Physical and functional interactome atlas of human receptor tyrosine kinases. *EMBO reports*, 23(6), e54041.

Reint G, et al. (2021) Rapid genome editing by CRISPR-Cas9-POLD3 fusion. *eLife*, 10.

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.