

Resource Summary Report

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

pCSF107mT-GATEWAY-3'-FLAG

RRID:Addgene_67619

Type: Plasmid

Proper Citation

RRID:Addgene_67619

Plasmid Information

URL: <http://www.addgene.org/67619>

Proper Citation: RRID:Addgene_67619

Bacterial Resistance: Chloramphenicol and Ampicillin

Defining Citation: [PMID:26391338](#)

Vector Backbone Description: Backbone Size:4181; Vector Backbone:PCS2+ was modified to generate pCSf107mT; Vector Types:Mammalian Expression, in vitro transcription or translation off of an SP6 promoter, Xenopus expression; Bacterial Resistance:Chloramphenicol and Ampicillin

Plasmid Name: pCSF107mT-GATEWAY-3'-FLAG

Record Creation Time: 20220422T222422+0000

Record Last Update: 20260815T081840+0000

Ratings and Alerts

No rating or validation information has been found for pCSF107mT-GATEWAY-3'-FLAG.

No alerts have been found for pCSF107mT-GATEWAY-3'-FLAG.

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](#) .

Mattioli F, et al. (2025) Bi-allelic variants in BRF2 are associated with perinatal death and craniofacial anomalies. *Genome medicine*, 17(1), 38.

Aaltonen MJ, et al. (2022) Serine palmitoyltransferase assembles at ER-mitochondria contact sites. *Life science alliance*, 5(2).

Stomberski CT, et al. (2022) A multienzyme S-nitrosylation cascade regulates cholesterol homeostasis. *Cell reports*, 41(4), 111538.

Hargreaves CE, et al. (2022) Resolving the polygenic aetiology of a late onset combined immune deficiency caused by NFKB1 haploinsufficiency and modified by PIK3R1 and TNFRSF13B variants. *Clinical immunology (Orlando, Fla.)*, 234, 108910.

Hogan AK, et al. (2021) UBR7 acts as a histone chaperone for post-nucleosomal histone H3. *The EMBO journal*, 40(24), e108307.

Linares JF, et al. (2021) PKC δ inhibition activates an ULK2-mediated interferon response to repress tumorigenesis. *Molecular cell*, 81(21), 4509.

Ware TB, et al. (2020) Reprogramming fatty acyl specificity of lipid kinases via C1 domain engineering. *Nature chemical biology*, 16(2), 170.

Franks CE, et al. (2017) The Ligand Binding Landscape of Diacylglycerol Kinases. *Cell chemical biology*, 24(7), 870.

Grant IM, et al. (2015) The Xenopus ORFeome: A resource that enables functional genomics. *Developmental biology*, 408(2), 345.