

Resource Summary Report

Generated by [RRID](#) on Aug 18, 2026

pLenti CMVtight Puro DEST (w768-1)

RRID:Addgene_26430

Type: Plasmid

Proper Citation

RRID:Addgene_26430

Plasmid Information

URL: <http://www.addgene.org/26430>

Proper Citation: RRID:Addgene_26430

Insert Name: no insert

Bacterial Resistance: Chloramphenicol and Ampicillin

Vector Backbone Description: Backbone Size:9402; Vector Backbone:p156RRL-sinPPT-CMV-GFP-PRE/Nhe I; Vector Types:Mammalian Expression, Lentiviral, Tet-On Advanced Destination vector; Bacterial Resistance:Chloramphenicol and Ampicillin

Comments: Need to be transduced in a Tet-On advanced cell line to obtain expression of the GOI.

Plasmid Name: pLenti CMVtight Puro DEST (w768-1)

Record Creation Time: 20220422T222118+0000

Record Last Update: 20260815T081252+0000

Ratings and Alerts

No rating or validation information has been found for pLenti CMVtight Puro DEST (w768-1).

No alerts have been found for pLenti CMVtight Puro DEST (w768-1).

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Lee H, et al. (2026) RB loss modulates chromatin organization by regulating cohesin-dependent loops and enhancer-promoter interactions. *Nature communications*, 17(1).

Kaler CJ, et al. (2022) BAP1 Loss Promotes Suppressive Tumor Immune Microenvironment via Upregulation of PROS1 in Class 2 Uveal Melanomas. *Cancers*, 14(15).

Sanidas I, et al. (2022) Chromatin-bound RB targets promoters, enhancers, and CTCF-bound loci and is redistributed by cell-cycle progression. *Molecular cell*, 82(18), 3333.

Junqueira Alves C, et al. (2021) Plexin-B2 orchestrates collective stem cell dynamics via actomyosin contractility, cytoskeletal tension and adhesion. *Nature communications*, 12(1), 6019.

Fenini G, et al. (2018) Genome Editing of Human Primary Keratinocytes by CRISPR/Cas9 Reveals an Essential Role of the NLRP1 Inflammasome in UVB Sensing. *The Journal of investigative dermatology*, 138(12), 2644.

Garstkiewicz M, et al. (2017) Opposing effects of Nrf2 and Nrf2-activating compounds on the NLRP3 inflammasome independent of Nrf2-mediated gene expression. *European journal of immunology*, 47(5), 806.

Morgan CW, et al. (2015) Engineered cellular gene-replacement platform for selective and inducible proteolytic profiling. *Proceedings of the National Academy of Sciences of the United States of America*, 112(27), 8344.