Resource Summary Report

Generated by dkNET on Aug 8, 2026

pIVM_26

RRID:Addgene_204501
Type: Plasmid

Proper Citation

RRID:Addgene_204501

Plasmid Information

URL: http://www.addgene.org/204501

Proper Citation: RRID:Addgene_204501

Insert Name: EloB (1-104) and EloC (17-112)

Organism: Homo sapiens

Bacterial Resistance: Streptomycin

Defining Citation: PMID:28591624

Vector Backbone Description: Backbone Marker:Novagen; Backbone Size:3781; Vector Backbone:pCDF-1b DUET; Vector Types:Bacterial Expression; Bacterial Resistance:Streptomycin

Comments: Gadd MS, Testa A, Lucas X, Chan KH, Chen W, Lamont DJ, Zengerle M, Ciulli A. Structural basis of PROTAC cooperative recognition for selective protein degradation. Nat Chem Biol. 2017 May;13(5):514-521. doi: 10.1038/nchembio.2329. Epub 2017 Mar 13. PMID: 28288108; PMCID: PMC5392356.

Plasmid Name: pIVM_26

Record Creation Time: 20230907T080451+0000

Record Last Update: 20260808T082858+0000

Ratings and Alerts

No rating or validation information has been found for pIVM_26.

No alerts have been found for pIVM_26.

Data and Source Information

Source: Addgene

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Samelson AJ, et al. (2026) CRISPR screens in iPSC-derived neurons reveal principles of tau proteostasis. Cell, 189(5), 1517.

Glassman CR, et al. (2026) Virome-wide ubiquitin ligase discovery reveals diverse mechanisms of immune evasion. Science (New York, N.Y.).

Lu X, et al. (2025) Optimization of the PROTAC linker region of the proteasome substrate receptor hRpn13 rationalized by structural modeling with molecular dynamics. The Journal of biological chemistry, 301(6), 108520.