

Resource Summary Report

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

pCMV-Neo-Bam DP1

RRID:Addgene_37968

Type: Plasmid

Proper Citation

RRID:Addgene_37968

Plasmid Information

URL: <http://www.addgene.org/37968>

Proper Citation: RRID:Addgene_37968

Insert Name: DP1

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:8657117](#)

Vector Backbone Description: Backbone Marker:Vogelstein Lab ; Backbone Size:6600; Vector Backbone:pCMV-Neo-Bam; Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

Comments: DP1 functions as a heterodimer with several members of the E2F family of transcription factors.

Plasmid Name: pCMV-Neo-Bam DP1

Record Creation Time: 20220422T222200+0000

Record Last Update: 20260815T081418+0000

Ratings and Alerts

No rating or validation information has been found for pCMV-Neo-Bam DP1.

No alerts have been found for pCMV-Neo-Bam DP1.

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 [RRID](#) .

Saad D, et al. (2021) High Conformational Flexibility of the E2F1/DP1/DNA Complex. Journal of molecular biology, 433(18), 167119.

Clijsters L, et al. (2019) Cyclin F Controls Cell-Cycle Transcriptional Outputs by Directing the Degradation of the Three Activator E2Fs. Molecular cell, 74(6), 1264.

Terr?? B, et al. (2016) GEMC1 is a critical regulator of multiciliated cell differentiation. The EMBO journal, 35(9), 942.