

Resource Summary Report

Generated by [NIF](#) on Jul 29, 2026

pMSCVhyg-H3.3WT-FLAG/HA

RRID:Addgene_128740

Type: Plasmid

Proper Citation

RRID:Addgene_128740

Plasmid Information

URL: <http://www.addgene.org/128740>

Proper Citation: RRID:Addgene_128740

Insert Name: H3.3 WT

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:30335158](#)

Vector Backbone Description: Backbone Marker:Clontech; Backbone Size:553; Vector Backbone:pMSCVhyg; Vector Types:Retroviral; Bacterial Resistance:Ampicillin

Plasmid Name: pMSCVhyg-H3.3WT-FLAG/HA

Record Creation Time: 20220422T221716+0000

Record Last Update: 20260725T073753+0000

Ratings and Alerts

No rating or validation information has been found for pMSCVhyg-H3.3WT-FLAG/HA.

No alerts have been found for pMSCVhyg-H3.3WT-FLAG/HA.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Tsukahara T, et al. (2025) Division of labor among H3K4 methyltransferases defines distinct facets of homeostatic plasticity. Cell reports, 44(6), 115746.

Tsukahara T, et al. (2023) Division of labor among H3K4 Methyltransferases Defines Distinct Facets of Homeostatic Plasticity. bioRxiv : the preprint server for biology.