

# Resource Summary Report

Generated by [RRID](#) on Aug 24, 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:** 20260815T080450+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 [RRID](#) .

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.