

Resource Summary Report

Generated by [RRID](#) on Sep 1, 2026

pIND-EGFP

RRID:Addgene_15770

Type: Plasmid

Proper Citation

RRID:Addgene_15770

Plasmid Information

URL: <http://www.addgene.org/15770>

Proper Citation: RRID:Addgene_15770

Insert Name: mH2A.Z-EGFP

Organism: Mus musculus

Bacterial Resistance: Ampicillin

Vector Backbone Description: Backbone Marker:Invitrogen; Backbone Size:5000;
Vector Backbone:pIND; Vector Types:Mammalian Expression; Bacterial
Resistance:Ampicillin

Plasmid Name: pIND-EGFP

Record Creation Time: 20220422T221854+0000

Record Last Update: 20260829T075759+0000

Ratings and Alerts

No rating or validation information has been found for pIND-EGFP.

No alerts have been found for pIND-EGFP.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Wright JL, et al. (2025) The INO80 Chromatin Remodeling Complex Regulates Histone H2A.Z Mobility and the G1-S Transition in Oligodendrocyte Precursors. *Glia*, 73(6), 1307.

Jung H, et al. (2024) Structural and Biochemical Characterization of the Nucleosome Containing Variants H3.3 and H2A.Z. *Epigenomes*, 8(2).

Sokolova V, et al. (2023) Structural mechanism of HP1??-dependent transcriptional repression and chromatin compaction. *bioRxiv : the preprint server for biology*.

Lewis TS, et al. (2021) Structural basis of chromatin regulation by histone variant H2A.Z. *Nucleic acids research*, 49(19), 11379.