

Resource Summary Report

Generated by [nidm-terms](#) on Aug 9, 2026

pGZ110 (pFA6a-3FLAG-MNase-TRP1)

RRID:Addgene_70233

Type: Plasmid

Proper Citation

RRID:Addgene_70233

Plasmid Information

URL: <http://www.addgene.org/70233>

Proper Citation: RRID:Addgene_70233

Insert Name: 3xFLAG tag and aa 83-231 of Mnase

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:26490019](#)

Vector Backbone Description: Vector Backbone:pFA6a-3HA-Trp1; Vector Types:Yeast genomic targeting; Bacterial Resistance:Ampicillin

Plasmid Name: pGZ110 (pFA6a-3FLAG-MNase-TRP1)

Record Creation Time: 20220422T222434+0000

Record Last Update: 20260808T082046+0000

Ratings and Alerts

No rating or validation information has been found for pGZ110 (pFA6a-3FLAG-MNase-TRP1).

No alerts have been found for pGZ110 (pFA6a-3FLAG-MNase-TRP1).

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 [nidm-terms](#) .

Evatt JM, et al. (2024) Centromere pairing enables correct segregation of meiotic chromosomes. *Current biology* : CB, 34(10), 2085.

Gr??nberg S, et al. (2017) Genome-wide Mapping of Protein-DNA Interactions with ChEC-seq in *Saccharomyces cerevisiae*. *Journal of visualized experiments* : JoVE(124).