

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 14, 2025

3xHA-miniTurbo-NLS_pCDNA3

RRID:Addgene_107172

Type: Plasmid

Proper Citation

RRID:Addgene_107172

Plasmid Information

URL: <http://www.addgene.org/107172>

Proper Citation: RRID:Addgene_107172

Insert Name: miniTurbo (BirA mutant)

Organism: Other

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:30125270](https://pubmed.ncbi.nlm.nih.gov/30125270/)

Vector Backbone Description: Vector Backbone:pCDNA3; Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

Comments: Please visit <https://www.biorxiv.org/content/early/2017/10/02/196980> for BioRxiv preprint

Plasmid Name: 3xHA-miniTurbo-NLS_pCDNA3

Relevant Mutation: aa1-63 deleted; Q65P, I87V, R118S, E140K, Q141R, S150G, L151P, V160A, T192A, K194I, M209V, I305V

Record Creation Time: 20220422T221519+0000

Record Last Update: 20220422T221629+0000

Ratings and Alerts

No rating or validation information has been found for 3xHA-miniTurbo-NLS_pCDNA3.

No alerts have been found for 3xHA-miniTurbo-NLS_pCDNA3.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Stockhammer A, et al. (2024) When less is more - a fast TurboID knock-in approach for high-sensitivity endogenous interactome mapping. *Journal of cell science*, 137(16).

Gillis NE, et al. (2022) Thyroid hormone dependent transcriptional programming by TR? requires SWI/SNF chromatin remodelers. *Nucleic acids research*, 50(3), 1382.

Liu Y, et al. (2022) Spatiotemporal Proximity Labeling Tools to Track GlcNAc Sugar-Modified Functional Protein Hubs during Cellular Signaling. *ACS chemical biology*, 17(8), 2153.

Piët ACA, et al. (2022) Proximity Ligation Mapping of Microcephaly Associated SMPD4 Shows Association with Components of the Nuclear Pore Membrane. *Cells*, 11(4).

Estell C, et al. (2021) ZC3H4 restricts non-coding transcription in human cells. *eLife*, 10.