

Resource Summary Report

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

pDEST-N2H-F1-empty vector

RRID:Addgene_125551

Type: Plasmid

Proper Citation

RRID:Addgene_125551

Plasmid Information

URL: <http://www.addgene.org/125551>

Proper Citation: RRID:Addgene_125551

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:31467278](https://pubmed.ncbi.nlm.nih.gov/31467278/)

Vector Backbone Description: Backbone Marker:we built; Backbone Size:8332; Vector Backbone:custom made; Vector Types:Mammalian Expression, Yeast Expression, in vitro expression; Bacterial Resistance:Ampicillin

Comments: Please visit <https://www.biorxiv.org/content/10.1101/530790v1.full> for bioRxiv preprint.

Plasmid Name: pDEST-N2H-F1-empty vector

Record Creation Time: 20220422T221658+0000

Record Last Update: 20260725T073723+0000

Ratings and Alerts

No rating or validation information has been found for pDEST-N2H-F1-empty vector.

No alerts have been found for pDEST-N2H-F1-empty vector.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Lambourne L, et al. (2025) Widespread variation in molecular interactions and regulatory properties among transcription factor isoforms. *Molecular cell*, 85(7), 1445.

Trepte P, et al. (2024) AI-guided pipeline for protein-protein interaction drug discovery identifies a SARS-CoV-2 inhibitor. *Molecular systems biology*, 20(4), 428.

Trepte P, et al. (2023) AI-guided pipeline for protein-protein interaction drug discovery identifies a SARS-CoV-2 inhibitor. *bioRxiv : the preprint server for biology*.