

Resource Summary Report

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

pCIBN-hTRF2-tagRFP-T

RRID:Addgene_103812

Type: Plasmid

Proper Citation

RRID:Addgene_103812

Plasmid Information

URL: <http://www.addgene.org/103812>

Proper Citation: RRID:Addgene_103812

Insert Name: CIBN-hTRF2-tagRFP-T

Organism: Homo sapiens

Bacterial Resistance: Kanamycin

Defining Citation: [PMID:29122982](#)

Vector Backbone Description: Backbone Marker:Clontech; Backbone Size:3932; Vector Backbone:pEGFP-C1; Vector Types:Mammalian Expression; Bacterial Resistance:Kanamycin

Comments: EGFP was cut out from the backbone (NheI/BamHI cuts)

Plasmid Name: pCIBN-hTRF2-tagRFP-T

Relevant Mutation: hTRF2: deletion of amino acids 1-42 and 476 compared to NP_005643.2

Record Creation Time: 20220422T221503+0000

Record Last Update: 20260808T080328+0000

Ratings and Alerts

No rating or validation information has been found for pCIBN-hTRF2-tagRFP-T.

No alerts have been found for pCIBN-hTRF2-tagRFP-T.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Dopie J, et al. (2020) Tyramide signal amplification mass spectrometry (TSA-MS) ratio identifies nuclear speckle proteins. The Journal of cell biology, 219(9).