

# Resource Summary Report

Generated by [dkNET](#) on Sep 14, 2026

## NdeI – SpeI(NheI) – BamHI/BglII - ANXA2 (23-339)

RRID:Addgene\_136546

Type: Plasmid

### Proper Citation

RRID:Addgene\_136546

### Plasmid Information

**URL:** <http://www.addgene.org/136546>

**Proper Citation:** RRID:Addgene\_136546

**Bacterial Resistance:** Ampicillin

**Defining Citation:** [PMID:32442400](#)

**Vector Backbone Description:** Vector Backbone:pET15; Vector Types:Bacterial Expression; Bacterial Resistance:Ampicillin

**Comments:** More data in this paper: Structure determination of the transactivation domain of p53 in complex with S100A4 using annexin A2 as a crystallization chaperone

**Plasmid Name:** NdeI – SpeI(NheI) – BamHI/BglII - ANXA2 (23-339)

**Record Creation Time:** 20220422T221753+0000

**Record Last Update:** 20260912T091915+0000

### Ratings and Alerts

No rating or validation information has been found for NdeI – SpeI(NheI) – BamHI/BglII - ANXA2 (23-339).

No alerts have been found for NdeI – SpeI(NheI) – BamHI/BglII - ANXA2 (23-339).

### 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 [dkNET](#) .

Cousido-Siah A, et al. (2022) A scalable strategy to solve structures of PDZ domains and their complexes. *Acta crystallographica. Section D, Structural biology*, 78(Pt 4), 509.

Ecsédi P, et al. (2020) Structure Determination of the Transactivation Domain of p53 in Complex with S100A4 Using Annexin A2 as a Crystallization Chaperone. *Structure* (London, England : 1993), 28(8), 943.