Resource Summary Report

Generated by FDI Lab - SciCrunch.org on May 5, 2024

pET28a-SENP2 (catalytic domain)

RRID:Addgene_16357 Type: Plasmid

Proper Citation

RRID:Addgene_16357

Plasmid Information

URL: http://www.addgene.org/16357

Proper Citation: RRID:Addgene_16357

Insert Name: human SENP2 catalytic domain

Organism: Homo sapiens

Bacterial Resistance: Kanamycin

Defining Citation: PMID:17591783

Vector Backbone Description: Backbone Marker:Novagen; Backbone Size:5369; Vector Backbone:pET28a; Vector Types:Bacterial Expression; Bacterial Resistance:Kanamycin

Plasmid Name: pET28a-SENP2 (catalytic domain)

Relevant Mutation: Amino Acid 365 to 590 catalytic domain of human SENP2.

Ratings and Alerts

No rating or validation information has been found for pET28a-SENP2 (catalytic domain).

No alerts have been found for pET28a-SENP2 (catalytic domain).

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 FDI Lab - SciCrunch.org.

Williams C, et al. (2023) The Ufd1 cofactor determines the linkage specificity of polyubiquitin chain engagement by the AAA+ ATPase Cdc48. Molecular cell, 83(5), 759.

Leonen CJA, et al. (2021) Sumoylation of the human histone H4 tail inhibits p300-mediated transcription by RNA polymerase II in cellular extracts. eLife, 10.