Resource Summary Report

Generated by RRID on Aug 19, 2026

pFGET19_Ulp1

RRID:Addgene_64697
Type: Plasmid

Proper Citation

RRID:Addgene_64697

Plasmid Information

URL: http://www.addgene.org/64697

Proper Citation: RRID:Addgene_64697

Insert Name: ULP1

Organism: Saccharomyces cerevisiae

Bacterial Resistance: Kanamycin

Defining Citation: PMID:26297996

Vector Backbone Description: Backbone Marker:Novagen; Backbone Size:5368; Vector Backbone:pET-28b(+); Vector Types:Bacterial Expression; Bacterial Resistance:Kanamycin

Plasmid Name: pFGET19_Ulp1

Relevant Mutation: Ulp1(403-621)

Record Creation Time: 20220422T222409+0000

Record Last Update: 20260815T081812+0000

Ratings and Alerts

No rating or validation information has been found for pFGET19_Ulp1.

No alerts have been found for pFGET19_Ulp1.

Data and Source Information

Source: Addgene

Usage and Citation Metrics

We found 33 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Sousa T, et al. (2026) Lipid composition controls the huntingtin exon 1 membrane-association and differentially modulates its flanking regions' dynamics. Protein science : a publication of the Protein Society, 35(6), e70612.

Arkinson C, et al. (2025) NUB1 traps unfolded FAT10 for ubiquitin-independent degradation by the 26S proteasome. Nature structural & molecular biology, 32(9), 1752.

Bratt AS, et al. (2025) Pharmacologic interrogation of USP28 cellular function in p53 signaling. Cell chemical biology.

Gemeinhardt TM, et al. (2025) A disordered linker in the Polycomb protein Polyhomeotic tunes phase separation and oligomerization. Molecular cell, 85(11), 2128.

Greene E, et al. (2025) Product-stabilized filamentation by human glutamine synthetase allosterically tunes metabolic activity. bioRxiv : the preprint server for biology.

Cáceres JC, et al. (2025) Nitric Oxide Inhibition of Glycyl Radical Enzymes and Their Activases. Journal of the American Chemical Society, 147(14), 11777.

Lambrechts LS, et al. (2025) The BRD4-nucleosome interaction is enhanced modestly and non-selectively by histone acetylation. Nucleic acids research, 53(22).

Solodkov PP, et al. (2024) Serial Llama Immunization with Various SARS-CoV-2 RBD Variants Induces Broad Spectrum Virus-Neutralizing Nanobodies. Vaccines, 12(2).

Arkinson C, et al. (2024) Nub1 traps unfolded FAT10 for ubiquitin-independent degradation by the 26S proteasome. bioRxiv : the preprint server for biology.

Lin J, et al. (2024) Design principles to tailor Hsp104 therapeutics. Cell reports, 43(12), 115005.

Singh AK, et al. (2023) Calcium-assisted sortase A cleavage of SUMOylated metallothionein constructs leads to high-yield production of human MT3. Microbial cell factories, 22(1), 125.

Demay F, et al. (2023) Simple purification and characterization of soluble and homogenous ABC-F translation factors from Enterococcus faecium. Protein expression and purification, 210, 106325.

Meeks KR, et al. (2023) Expression and kinetic characterization of PYCR3. Archives of biochemistry and biophysics, 733, 109468.

Hamaker NK, et al. (2023) High-efficiency and multilocus targeted integration in CHO cells using CRISPR-mediated donor nicking and DNA repair inhibitors. Biotechnology and bioengineering, 120(9), 2419.

Martinez-D'Alto A, et al. (2023) Characterization of a unique polysaccharide monooxygenase from the plant pathogen Magnaporthe oryzae. Proceedings of the National Academy of Sciences of the United States of America, 120(8), e2215426120.

Mani S, et al. (2023) Targeting DPP4-RBD interactions by sitagliptin and linagliptin delivers a potential host-directed therapy against pan-SARS-CoV-2 infections. International journal of biological macromolecules, 245, 125444.

Gemeinhardt TM, et al. (2023) How a disordered linker in the Polycomb protein Polyhomeotic tunes phase separation and oligomerization. bioRxiv : the preprint server for biology.

Tovar-Ramírez ME, et al. (2023) ATCUN-like Copper Site in ?B2-Crystallin Plays a Protective Role in Cataract-Associated Aggregation. Inorganic chemistry, 62(27), 10592.

Chen Z, et al. (2022) Protein purification, crystallization, and structure determination of human DEAD-box RNA helicase DDX21 in different unwinding states. STAR protocols, 3(3), 101642.

Kozielski F, et al. (2022) Identification of fragments binding to SARS-CoV-2 nsp10 reveals ligand-binding sites in conserved interfaces between nsp10 and nsp14/nsp16. RSC chemical biology, 3(1), 44.