

Resource Summary Report

Generated by [kravitz2](#) on Jul 29, 2026

pET His6 TEV LIC cloning vector (1B)

RRID:Addgene_29653

Type: Plasmid

Proper Citation

RRID:Addgene_29653

Plasmid Information

URL: <http://www.addgene.org/29653>

Proper Citation: RRID:Addgene_29653

Insert Name: None

Bacterial Resistance: Kanamycin

Vector Backbone Description: Backbone Size:5343; Vector Backbone:pET; Vector Types:Bacterial Expression; Bacterial Resistance:Kanamycin

Comments: This plasmid is an empty vector to be used with a LIC cloning protocol. It has a TEV-cleavable His6 fusion tag on its N-terminus. To clone into this vector, add LIC fusion tags to the 5' end of your PCR primers. Forward - 5'TACTTCCAATCCAATGCA3' Reverse - 5'TTATCCACTTCCAATGTTATTA3' Linearize the plasmid with SspI and gel purify. When digesting the DNA with T4 polymerase, use dCTP for insert and dGTP for vector. More information on this vector can be found through <http://qb3.berkeley.edu/qb3/macrolab/>

Plasmid Name: pET His6 TEV LIC cloning vector (1B)

Record Creation Time: 20220422T222131+0000

Record Last Update: 20260725T074533+0000

Ratings and Alerts

No rating or validation information has been found for pET His6 TEV LIC cloning vector (1B).

No alerts have been found for pET His6 TEV LIC cloning vector (1B).

Data and Source Information

Usage and Citation Metrics

We found 64 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Burger M, et al. (2025) Encapsulation of protein/DNA complexes into unilamellar liposomes via annexin-mediated membrane recruitment and sonication. *Cell reports methods*, 5(6), 101073.

Cuevas-Navarro A, et al. (2025) Pharmacological restoration of GTP hydrolysis by mutant RAS. *Nature*, 637(8044), 224.

Chen MZ, et al. (2025) A versatile antibody capture system drives specific in vivo delivery of mRNA-loaded lipid nanoparticles. *Nature nanotechnology*, 20(9), 1273.

Zoabi FH, et al. (2025) Inhibiting MARCH5/Mfn2 signaling as an alternative strategy to protect cardiomyocytes from hypoxia-induced mitochondrial dysfunction. *Computational and structural biotechnology journal*, 27, 3045.

K?l?ç G, et al. (2025) Expression of a Tuberculosis-Associated Immunogenic Protein in *Escherichia coli*. *Life (Basel, Switzerland)*, 15(9).

Kaus-Drobek M, et al. (2025) From discovery to potential application: engineering a novel M23 peptidase to combat *Listeria monocytogenes*. *Scientific reports*, 15(1), 15628.

Hoogstra SJ, et al. (2025) Biotransformation of Deoxynivalenol to the Novel Metabolite Deoxynivalenol-8,15-hemiketal-7-glucoside by the *Bacillus subtilis* Glycosyltransferase YjiC. *ACS omega*, 10(14), 14496.

Gjindali A, et al. (2025) Two cowpea Rubisco activase isoforms for crop thermotolerance. *The New phytologist*, 247(3), 1199.

Feder JS, et al. (2025) A fluorescent-protein spin qubit. *Nature*, 645(8079), 73.

Herrera-Asmat O, et al. (2025) Pleomorphic effects of three small-molecule inhibitors on transcription elongation by *Mycobacterium tuberculosis* RNA polymerase. *eLife*, 14.

Herrera-Asmat O, et al. (2025) Pleomorphic effects of three small-molecule inhibitors on transcription elongation by *Mycobacterium tuberculosis* RNA polymerase. *bioRxiv : the preprint server for biology*.

Dawes S, et al. (2024) Chaperone BiP controls ER stress sensor Ire1 through interactions with its oligomers. *Life science alliance*, 7(10).

Rivera-de-Torre E, et al. (2024) Discovery of broadly-neutralizing antibodies against brown recluse spider and Gadim scorpion sphingomyelinases using consensus toxins as antigens. *Protein science : a publication of the Protein Society*, 33(3), e4901.

Mevisse TET, et al. (2024) STK19 positions TFIIH for cell-free transcription-coupled DNA repair. *Cell*, 187(25), 7091.

Liu Y, et al. (2024) MECP2 directly interacts with RNA polymerase II to modulate transcription in human neurons. *Neuron*, 112(12), 1943.

Saha A, et al. (2024) An alpha-helical lid guides the target DNA toward catalysis in CRISPR-Cas12a. *Nature communications*, 15(1), 1473.

Yin JZ, et al. (2024) The HisRS-like domain of GCN2 is a pseudoenzyme that can bind uncharged tRNA. *Structure (London, England : 1993)*, 32(6), 795.

Kokic G, et al. (2024) Structural basis for RNA polymerase II ubiquitylation and inactivation in transcription-coupled repair. *Nature structural & molecular biology*, 31(3), 536.

Hajam IA, et al. (2023) Functional divergence of a bacterial enzyme promotes healthy or acneic skin. *Nature communications*, 14(1), 8061.

Hendricks JM, et al. (2023) Identification of structurally diverse FSP1 inhibitors that sensitize cancer cells to ferroptosis. *Cell chemical biology*, 30(9), 1090.