

Resource Summary Report

Generated by [dkNET](#) on Sep 11, 2026

[pQLinkN](#)

RRID:Addgene_13670

Type: Plasmid

Proper Citation

RRID:Addgene_13670

Plasmid Information

URL: <http://www.addgene.org/13670>

Proper Citation: RRID:Addgene_13670

Insert Name: multi cloning site

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:17311810](#)

Vector Backbone Description: Backbone Size:4794; Vector Backbone:pQE-2; Vector Types:Bacterial Expression, Co-expression; Bacterial Resistance:Ampicillin

Comments: Please note that the chloramphenicol resistance gene in this plasmid is incomplete. It is lacking a promoter, therefore it is supposed to be inactive.

Plasmid Name: pQLinkN

Record Creation Time: 20220422T221753+0000

Record Last Update: 20260905T074821+0000

Ratings and Alerts

No rating or validation information has been found for pQLinkN.

No alerts have been found for pQLinkN.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Zhao Z, et al. (2026) Structure of E. coli twin-arginine translocase (Tat) complex with bound cargo. *Molecular cell*, 86(13), 2572.

DAmico KA, et al. (2024) Structure of a membrane tethering complex incorporating multiple SNAREs. *Nature structural & molecular biology*, 31(2), 246.

DAmico KA, et al. (2023) Structure of a Membrane Tethering Complex Incorporating Multiple SNAREs. *bioRxiv : the preprint server for biology*.

Zhu L, et al. (2022) Structural Basis of the Immunity Mechanisms of Pediocin-like Bacteriocins. *Applied and environmental microbiology*, 88(13), e0048122.

Weiffert T, et al. (2018) Protein stabilization with retained function of monellin using a split GFP system. *Scientific reports*, 8(1), 12763.

Tripathi A, et al. (2017) Two alternative binding mechanisms connect the protein translocation Sec71-Sec72 complex with heat shock proteins. *The Journal of biological chemistry*, 292(19), 8007.

Baker RW, et al. (2013) Crystal Structures of the Sec1/Munc18 (SM) Protein Vps33, Alone and Bound to the Homotypic Fusion and Vacuolar Protein Sorting (HOPS) Subunit Vps16*. *PLoS one*, 8(6), e67409.

Tripathi A, et al. (2009) Structural characterization of Tip20p and Dsl1p, subunits of the Dsl1p vesicle tethering complex. *Nature structural & molecular biology*, 16(2), 114.

Ren Y, et al. (2009) A structure-based mechanism for vesicle capture by the multisubunit tethering complex Dsl1. *Cell*, 139(6), 1119.