

# Resource Summary Report

Generated by [RRID](#) on Aug 19, 2026

## pRK1043

RRID:Addgene\_8835

Type: Plasmid

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### Proper Citation

RRID:Addgene\_8835

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### Plasmid Information

**URL:** <http://www.addgene.org/8835>

**Proper Citation:** RRID:Addgene\_8835

**Insert Name:** tobacco etch virus protease, catalytic domain

**Organism:** tobacco etch virus

**Bacterial Resistance:** Chloramphenicol and Ampicillin

**Defining Citation:** [PMID:11809930](#)

**Vector Backbone Description:** Backbone Marker:New England Biolabs; Backbone Size:6700; Vector Backbone:pMal-C2; Vector Types:Bacterial Expression; Bacterial Resistance:Chloramphenicol and Ampicillin

**Comments:** pRK1043 overproduces the catalytic domain of tobacco etch virus (TEV) protease in the form of an affinity sandwich, with E. coli maltose-binding protein fused to its N-terminus and a polyarginine tag fused to its C-terminus. The TEV protease produced by this strain is the autoinactivation-resistant mutant S219V. The pRIL plasmid (Stratagene) improves the yield of TEV protease by increasing the supply of argU tRNA in the cell (for AGG and AGA codons). The catalytic activity of the S219V mutant is approximately 2-fold greater than that of the wild-type protease. Shifting the temperature from 37°C to 30°C during induction maximizes the yield of soluble MBP-TEV-Arg fusion protein. Bacterial strain: E.coli BL21(DE3)-RIL (Stratagene).

**Plasmid Name:** pRK1043

**Relevant Mutation:** Autolysis-resistant S219V mutant

**Record Creation Time:** 20220422T222604+0000

**Record Last Update:** 20260815T082150+0000

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## Ratings and Alerts

No rating or validation information has been found for pRK1043.

No alerts have been found for pRK1043.

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## Data and Source Information

**Source:** [Addgene](#)

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Liang Q, et al. (2025) Rational design yields RNA-binding zinc finger domains with altered sequence specificity. RNA (New York, N.Y.), 31(2), 150.

Dingal PCDP, et al. (2023) Molecular mechanisms controlling the biogenesis of the TGF- $\beta$  signal Vg1. Proceedings of the National Academy of Sciences of the United States of America, 120(43), e2307203120.

Kipniss NH, et al. (2017) Engineering cell sensing and responses using a GPCR-coupled CRISPR-Cas system. Nature communications, 8(1), 2212.

Brockly F, et al. (2016) Production and Purification of Recombinant SUMOylated Proteins Using Engineered Bacteria. Methods in molecular biology (Clifton, N.J.), 1475, 55.

Mentrup T, et al. (2015) A Cell-Based Assay Reveals Nuclear Translocation of Intracellular Domains Released by SPPL Proteases. Traffic (Copenhagen, Denmark), 16(8), 871.

Jun Y, et al. (2007) Sec18p and Vam7p remodel trans-SNARE complexes to permit a lipid-anchored R-SNARE to support yeast vacuole fusion. The EMBO journal, 26(24), 4935.