



Sep 24, 2019

## Loop L2 (even level) type IIS cloning - pCs-ye vectors

Eftychis Frangedakis<sup>1</sup>, Susana Sauret-Gueto<sup>2</sup>, Anthony West<sup>2</sup>, marta tomaselli<sup>3</sup>, Nicola Patron<sup>2</sup>, Marius Rebmann<sup>2</sup>, Jim Haseloff<sup>2</sup>

<sup>1</sup>University of Cambridge, Plant Sciences , OpenPlant, <sup>2</sup>Plant Sciences, University of Cambridge, OpenPlant, <sup>3</sup>University of Cambridge, Open Plant





ABSTRACT

## Protocol based on

Pollak B, Cerda A, Delmans M, et al (2019) Loop assembly: a simple and open system for recursive fabrication of DNA circuits. New Phytol 222:628–640

## https://doi.org/10.1111/nph.15625

MATERIALS		
NAME ~	CATALOG #	VENDOR ~
Sterile water		
dATP, 100mM, 25uMoles	U1205	Promega
BSA, molecular biology grade, 20 mg/ml	B9000S	New England Biolabs
Tango Buffer	BY5	Thermo Fisher Scientific
T4 DNA ligase	15224041	Thermo Fisher Scientific
Lgul (Sapl)	ER1931	Thermo Fisher Scientific

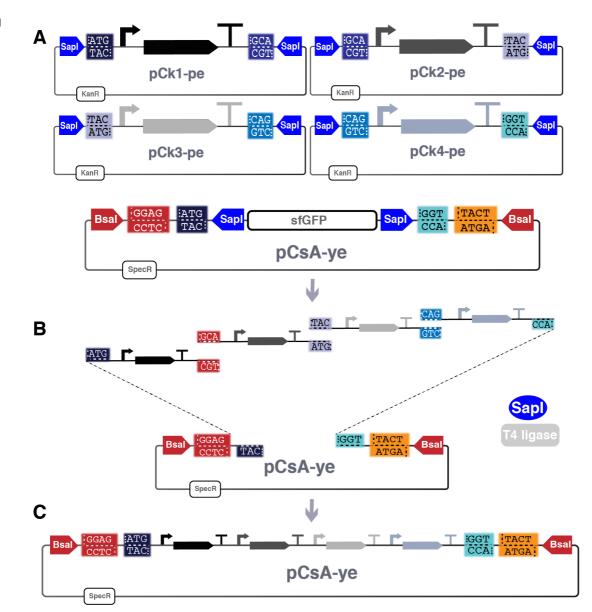
- 1 Determine DNA parts concentration with spectrophotometry (Nanodrop).
- 2 Prepare aliquots for DNA parts to be assembled at a concentration of 15 nM and of the pCs-pe vector at a concentration of 7.5 nM.

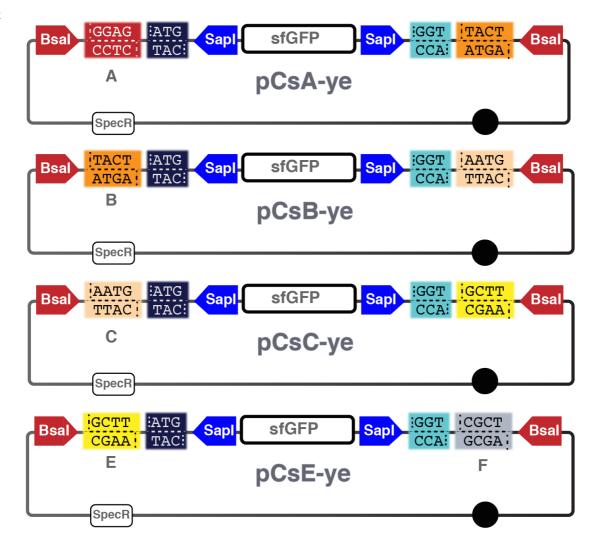
To calculate the concentration needed for each part to assemble (not the backbone) in  $ng/\mu L$ , divide the length of the plasmid where they are cloned by 100. For the backbone, divide the plasmid length by 200. In this way you can prepare aliquots of your parts and backbones and add 1  $\mu L$  of each one to the plasmid mix.

3 Prepare the Loop assembly Even Level reaction MM according to according to Table

Component	Volume (μL)
Sterile water	2
10x Tango buffer (Thermo Fisher)	1
1 mg/mL bovine serum albumin (NEB)	0.5
T4 DNA ligase (5 U/µL) (Thermo Fisher)	0.25
10mM ATP (SIGMA)	1
Sapl (Lgul) (5 U/µL) (Thermo Fisher)	0.25
Final volume	5

- 4 Prepare plasmids mix for each reaction, by adding in a 0.2 mL tube, 1  $\mu$ L of each DNA part aliquot (see step 2), 1  $\mu$ L of the pCs-pe vector and sterile water up to 5  $\mu$ L. Mix well.
- Add 5  $\mu$ L of master mix to the 5  $\mu$ L of plasmids mix, to a final volume of 10  $\mu$ L. Mix well. If more than 4 DNA parts are to be assembled into a pCk-pe vector, reduce the water volume in the MM by 1  $\mu$ L for each extra 1  $\mu$ L of DNA part added in the plasmids mix.
- 6 Place samples in a thermocycler and use the following program:
  Assembly: 26 cycles of 37 °C for 3 min and 16 °C for 4 min.
  Termination and enzyme denaturation: 50 °C for 5 min and 80 °C for 10 min.
- 7 Transform 20  $\mu$ L of chemically competent E. coli cells (transformation efficiency of 1 × 107 transformants/ $\mu$ g plasmid DNA) using 2  $\mu$ L of the Loop assembly reaction and then plate on LB agar plates containing 100  $\mu$ g/mL spectinomycin.
- 8 Incubate O/N at 37 °C.
- 9 Colonies with white color are likely to contain the vector with the insert while yellow color colonies will contain the empty vector.
- 10 Confirm the presence of the correct insert with Sanger sequencing using the primers pC\_F and pC\_R





This is an open access protocol distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited