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Optimize pyck cloning strategy

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Andreea S¹ ¹University of Groningen

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Protocol status: Other Protocol has been developed based on literature, but hasn't been tested yet.

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Abstract

Introducing NLP14a in the genome of B. mycoides: In order to create plasmid pYCR-gamyNLP, the backbone of Pycr will be digested with the PCR product of gamy_Fw and gamy_Rv to produce the pYCR_gamy (CRISPR vector containing the gRNA). The NLP14a sequence will be ordered from Twist Biosciences And PCR-amplified with the primers NLP_Fw and NLP_Rv. The resulting PCR product, as well as the pYCR_gamy vector will be digested with Sfil. The digested mixtures will be ligated using T4 ligase.

Kill switch cloning: The suggested kill switch mechanism is based on Trp auxotrophic strains as well as a toxinantitoxin mechanism. The toxin will be introduced in the genome using crispr in place of trpE. The antitoxin will be maintained as a cytoplasmic plasmid (pAD-YqcF) and expressed only in the presence of solanine. Unfortunately, to the best of our knowledge, no operator that binds solanine of B. mycoides has been described in literature. If it is provebn that no such operator is present, we suggest on oly keeping the tryptophan dependence strategy.

In order to create plasmid pYCR-gtrpEYqcG plasmid, the backbone of Pycr will be digested with Bsal and ligated with the annealing product of gtrpe_Fw and gtrpe_Rv to produce the pYCR_gtrp (CRISPR vector containing the gRNA). Synthetic dna for *yqcG* will be ordered from Twist Biosciences and PCR-amplified with the primers Yqcg_Fw and Ygcg_Rv. The resulting PCR product, as well as the pYCR_gtrpE vector will be digested with Sfil. The digested mixtures will be ligated using T4 ligase

Protocol was adapted after a paper of Yi et al., 2018

1. Yi, Y., Li, Z., Song, C. & Kuipers, O. P. Exploring plant-microbe interactions of the rhizobacteria Bacillus subtilis and Bacillus mycoides by use of the CRISPR-Cas9 system. Environ. Microbiol. 20, 4245–4260 (2018).

Clone sgRNA sequence into pYCR		
1	Design sgRNA spacer sequence (~20 nts) using "Benchling'" and choose B. mycoides M2E_15 genome Choose the gRNA with the highest on-target score and the lowest off-target score.	
2	Order the 2 complementary oligos flanked by overhang containing Bsal restriction site	
3	 Anneal oligos by mixing the following components: - Δ 70 μL Nuclease Free water - Δ 10 μL Oligo DNA annealing buffer 10x (IMJ 100 millimolar (mM)) Tris H 8 , IMJ 500 millimolar (mM) NaCl, IMJ 10 millimolar (mM) EDTA) - Δ 10 μL DNA oligos A and B (IMJ 10 micromolar (μM) each) 	
4	Mix well and incubate for 😒 00:05:00 at 🖁 90 °C 5m	
5	Cool slowly to room temperature (aprox. O1:00:00) by removing the heat block from the apparatus	
6	Run agarose gel (1%) to check the annealing product	
7	Digest pYCR with Bsal and inactivate the enzyme by PCR clean up	
8	Ligate the annealing product with the precut pYCR: MIx: - Δ 10 μL I of annealing product - Δ 60 ng of digested pYCR - Δ 1 μL T4 ligase	
9	Transform the ligation mixture into E. coli MC1061 using the heat-shock methos.	

Grow the cells on LB agar plates + Erithromycin [M] 0.1 mg/mL

10 To select the transformants perform a colony PCR with Fw primer binding upstream of PvanP and Rv primer downstream Sfil cloning sites.

Expected result

Succesful transformation will have a size of 427 pb

11 Miniprep and sequence the constructs

Clone flanking regions into pYCR containing sgRNA

Select the repair fragment of choice:

 for knock-out: PCR amplify aprox. 1 kpb upstream and downstream of the sgRNA targeted region
 for knock-in: sequence of the desired inserction sequence flanked by Sfil restriction sites compatible with the backbone of pYCR

- 13 Digest the pYCR(+sgRNA) and the repair fragments with Sfil Inactivate Sfil by PCR clean up
- 14 Ligate the components using T4 ligase Heat inactivate the ligase (incubate 🚫 00:20:00 😮 60 °C)

20m

- 15 Perform Smal digestion to get rid of the original pYCR
- 16 Transform the ligation mixture into E. coli MC1061 using the heat-shock methos. Grow the cells on LB agar plates + erythromycin
- 17 To select the transformants perform a colony PCR with Fw primer binding upstream of PvanP and Rv primer downstream Sfil cloning sites.
- 18 Miniprep and sequence the constructs

Transform final construct in B. mycoides M2E_15			
19	Transform using an electroporation protocol using $\boxed{25 \ \mu L}$ plasmid DNA		
	Protocol		
	Cloning of <i>Bacillus mycoides</i>		
	CREATED BY a.stan.6 PREVIEW		
19.1	Pick 1 colony of B. mycoides M2E_15 and inoculate it in BHIS		
19.2	When OD600nm reached 0.85 add 2% glycine and 2% threonine in order to weaken the cell wall.		
19.3	Grow the cells overnight at (5 200 rpm, 30°C		
19.4	Dilute the overnight culture 50 times in LBSP medium until the OD600nm reaches 0.65		
19.5	Collect the cells by		
19.6	Wash cell pellet with pre-chilled electroporation buffer (10% glycerol, 0.25 M sorbitol, 0.25M trehalose) x4		
19.7	Suspended in 1 ml electroporation buffer (10% glycerol, 0.25 M sorbitol, 0.25M trehalose).		

19.8 Snap-freeze the electrocompetent cells in liquid nitrogen and sore at **3** -80 °C

- 19.9 For the PCR mix pipette:
 - Δ 5 µL primers mixture Fw + Rv (5 uM) (see table "Primers")
 - 👗 10 μL HF buffer
 - <u>Δ</u> 1 μL dNTP (10mM)
 - DNA template (0.1 ng)
 - 😃 0.25 μL phusion polymerase
 - MIIiQ to 🛛 🕹 50 μL
- 19.10 PCR reaction is performed with the following protocol:

	Temperature (*C)	Time (mm: ss)
	98	05:00
	98	10:00
	Variable	00:30
	72	00:30
	repeat above steps	30x
Γ	72	10:00
	11	Infinit e

- ^{19.11} Load $4 5 \mu L$ of sample plus $4 1 \mu L$ of staining solution 6x. Check the length of the band (bp). Purify sample : PCR cleanup/ gel extraction according to manufacturer.
- 19.12 Digest the polymerized insert and the desired vector with 2U of apropriate restriction enzyme. See table ''Primers'' in order to choose the enzyme.

19.13	Digestion mixture ($_$ 20 µL): -Sfil 2 U - Plasmid ($_$ 50 ng) -Insert ($_$ 20 ng) - $_$ 2 µL CutSmart 10x (or appropriate buffer) -MiliQ to $_$ 20 µL
19.14	Incubate O1:00:00 at the temperature recommended by the manufacturer of the restriction enzyme
19.15	Inactivate the restriction enzyme by incubationg 20 min 65 °C **Sfil can't be heat inactivated so in order to inactivate it a PCR clean up is necesary
19.16	Ligate_the digested insert and vector. Ligation mixture : -10 ul digestion product -1 ul T4 -2 ul T4 ligase buffer -7 ul MQ
19.17	Incubate 02:00:00 From temperature
19.18	Thaw on ice $\boxed{100 \ \mu L}$ of electrocompetent cells
19.19	Add $\underline{A} 2 \mu g$ of plasmid and the aliquot of electrocompetent cells to ice-cold electroporation cuvettes
19.20	Electroporate 25 uF, 10 kV/cm, 200 Ohms
19.21	Add 1 ml of BHIS and incubate (5 100 rpm, 30°C, 05:00:00 for recovery
19.22	Plate on LB + Agar (1.5%) + 100 μg/ml spectinomycin. For crispr cloning, to activate cas9 expression, add 0.2% mannose.

19.23 Incubate at 🖁 30 °C 🚫 Overnight

- 19.24 Randomly pick potential mutants and test by colony PCR (for primers use the ones that flank the region of insertion sites).
- 19.25 After selecting colonies that show the expected band size, purify the desired plasmid (miniprep) and transform into B. mycoides.
- 20 Plate on LB agar containing erythromycin Grow at 28 °C overnight
- 21 Pick colonies and test them by colony PCR If the expected band size is present, purify the plasmid and sequence
- Grow the colonies with the expected sequence on liquid LB + erythromycin + manose
 Incubate (5 200 rpm, 28°C, 12:00:00
- 23 Next day pick colonies in BHI medium (without antibiotic) and grow at 37 °C overnight
- 24 Dilute the liquid cultures and plate them on LN (no antibiotic) grow overnight st **§** 37 °C