

WEEK 10

6.2 Sunyue Cai

Re-screening verification

We conducted a preliminary screening experiment of about 1300 monoclonals on 15 plates from No. 73 to 87, and selected 61 clones with high OD and fluorescence values for re-screening verification. The re-screening verification system is as follows:

Control Group	Test Group
220 μ L LB+0.1% 50 mg/mL ampicillin	220 μ L LB+0.1% 50 mg/mL ampicillin+1% BWPobR ^{mut}
220 μ L LB+0.1% 50 mg/mL ampicillin+1% BWPobR ^{WT} Bacteria	200 μ L LB+0.1% 50 mg/mL ampicillin+10% 0.6 g/L 4HB+1% BWPobR ^{mut}
200 μ L LB+0.1% 50 mg/mL ampicillin+10% 0.6 g/L 4HB+1% BWPobR ^{WT}	200 μ L LB+0.1% 50 mg/mL ampicillin+10% 0.6 g/L HMA+1% BWPobR ^{mut}

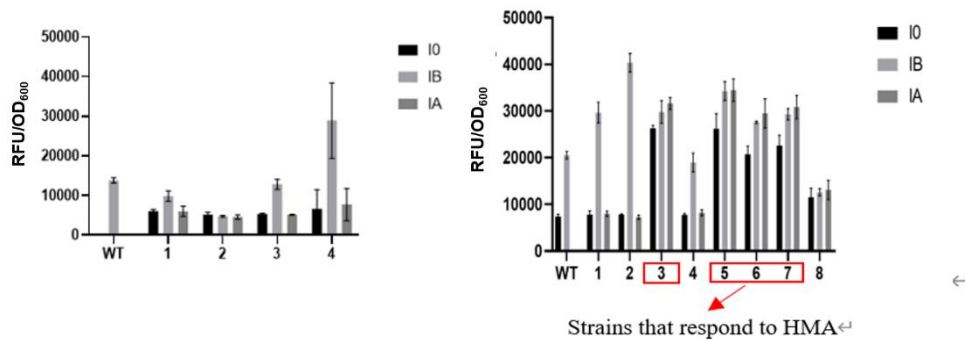


Figure 1 Data diagram of partial re-screening verification experiment
There are three parallel groups, but no strains screened meet the requirements.

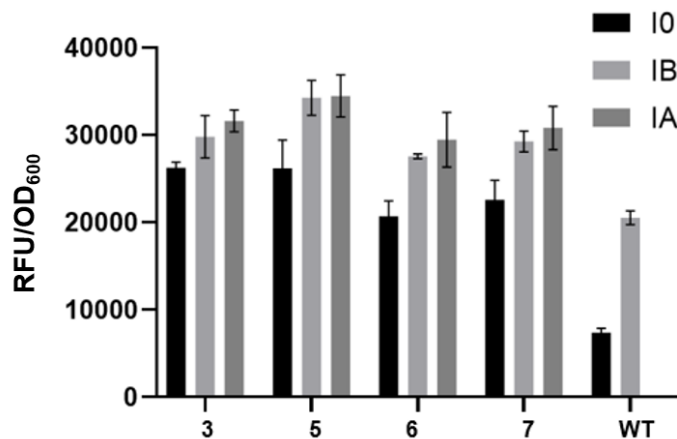


Figure 2 Re-screening verification diagrams of four strains with better responses to HMA

The above four strains have a high response to HMA. Due to the high I0 value, it is speculated that PobR protein mutants cannot inhibit the promoter. Repeated experiments are required to verify.

6.4 Jianing Li

Megawhop PCR

Our error-prone PCR system and PCR conditions are as follows:

PCR System		PCR Condition	
pYB1a-PobR-eGFP-Cmr	5 μ L / 50 ng	98 $^{\circ}$ C	3 min
2 \times Mut Random system	25 μ L	98 $^{\circ}$ C	30 s
Mut Enhancer	3 μ L	50 $^{\circ}$ C	30 s
pobR-mut-F	1 μ L	72 $^{\circ}$ C	30 s
pobR-mut-R	1 μ L	72 $^{\circ}$ C	2 min
ddW	19 μ L		

} X20

Then we use cycle-pure-kit kit for PCR product purification:

PCR System		PCR Condition	
pYB1a-PobR-eGFP-Cmr	5 μ L / 50 ng	95 $^{\circ}$ C	5 min
2x High fidelity mix	50 μ L	95 $^{\circ}$ C	30 s
Megaprimer	600 ng / 6 μ L	55 $^{\circ}$ C	30 s
ddW	39 μ L	72 $^{\circ}$ C	2 min30 s
		72 $^{\circ}$ C	10 min

} X30

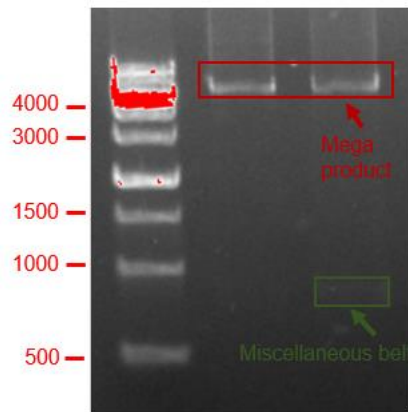


Figure 3 0528 mega results

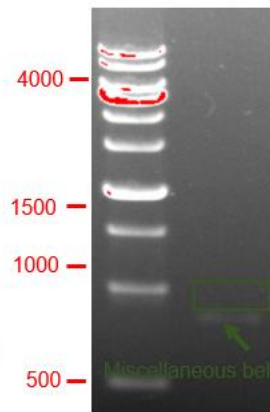


Figure 4 0601 mega results

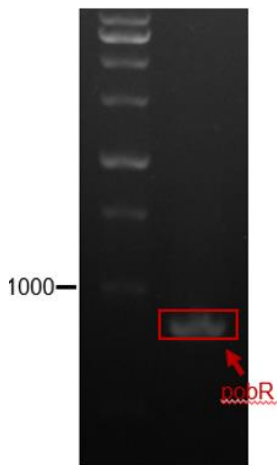


Figure 5 0528 Error-prone PCR results

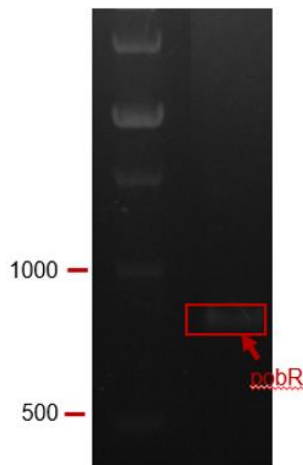


Figure 6 0601 Error-prone PCR results

6.6 Chao Chen

Preliminary test of flat plate screening

In order to determine whether the low-intensity induction can make the strain grow normally under the condition of adding 0.1% 20 mg/mL and 30 mg/mL chloramphenicol, the following experiment was done: antibiotic 1‰ + inducer 10% and 200 μ L were cultured in a shaker at 37°C. After the OD600 is about 4, the bacterial solution diluted 10^4 times is coated.



Figure7 BWPobR^{WT}+1% 50 mg/mL ampicillin

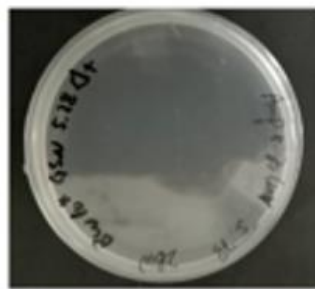


Figure 8 BWPobR^{WT}+1% 50 mg/mL ampicillin+1% 20 mg/mL chloramphenicol

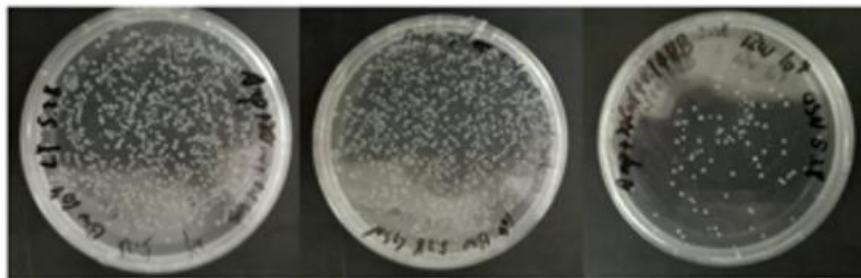


Figure 9 BWPobR^{WT}+1% 50 mg/mL ampicillin+1% 20 mg/mL chloramphenicol+0.1 g/L 4HB

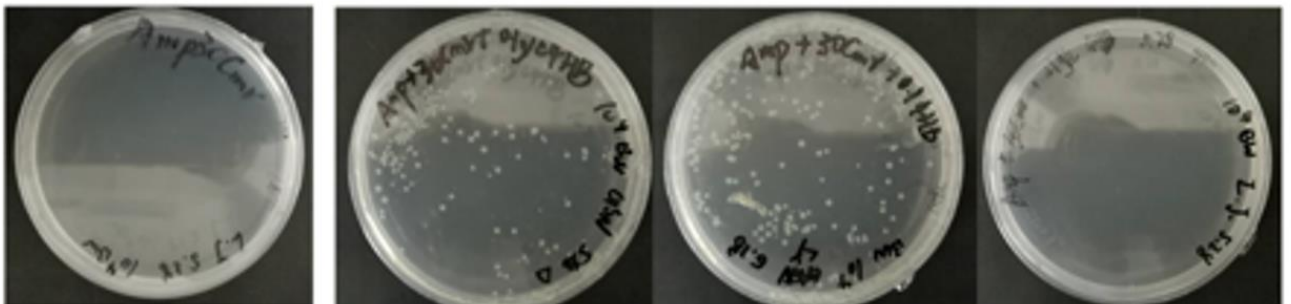


Figure 10 BWPobR^{WT}+1% 50 mg/mL ampicillin+1% 30 mg/mL chloramphenicol; 1% 50 mg/mL ampicillin+1% 30 mg/mL chloramphenicol+0.1 g/L 4HB

It is judged that under the condition of adding 0.1% 20 mg/mL and 30 mg/mL chloramphenicol, low-intensity induction can make the strain grow normally. In order to judge the random mutation library capacity and roughly judge the number of colonies that can grow in a single round of coating plate preliminary screening, the following experiments are carried out : After Megahop pcr, use DpnI endonuclease for digestion. The digestion system is as follows (4 tubes of 10 μ L digestion system) digestion at 37°C for 2 h, and digestion at 75°C for 20 min.

DpnI Enzyme	0.2 μ L
CutSmart	1 μ L
DNA	6 μ L (55 ng/ μ L)
DDW	2.8 μ L

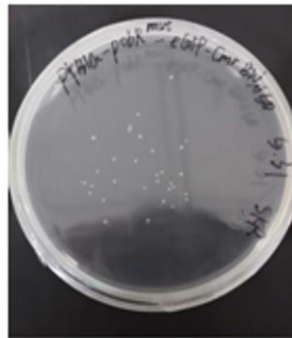


Figure12 50 μ L BWPobR^{WT}Bacteria+50 μ g/mL ampicillin

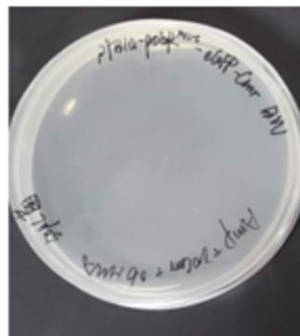


Figure13 50 μ L BWPobR^{WT} Bacteria+50 μ g/mL ampicillin + 30 μ g/mL chloramphenicol + 0.6 g/L HMA

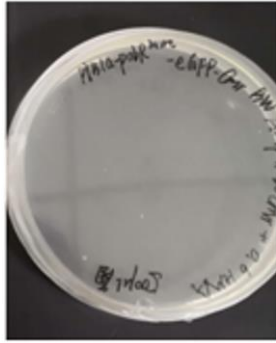


Figure14 500 μ L BWPobR^{WT} Bacteria+50 μ g/mL ampicillin + 20 μ g/mL chloramphenicol
+ 0.6 g/L HMA

Transfer 10 μ L digestion product into 50 μ L BW competence, add 600 μ L LB medium to 37°C shaker for 45 min, then take 50 μ L and 500 μ L bacterial solution for the preparation of coating engineering bacteria. The number of colonies was small, which was in line with expectations.