

Before 1st August, 2014

- Devised practical experiments.
- Amplified parts(T7 promoter, Terminator).
- Got AxyPrep Plasmid MiniprepKit(50-prep) and GBclonart Seamless Assembly Kit(GB2002-24).
- 100 times dilution of linear plasmid backbones, designed the primers.
- Placed an order for standard primers used for amplifying standard parts.
- Prepared competent Escherichia coli cells successfully.

2nd August, 2014 Sat

- Read papers about data and protocol.
- Determined detailed protocol of lambda red.
- Extracted plasmids of terminator.
- The bacteria had been transformed plasmids of T7 promoter did not form any colony.
- Amplified parts(T7 promoter, GFP,RFP,blue chromoprotein).

3rd August, 2014 Sun

- Prepared 4L Luria-Bertani medium.
- Prepared solid medium (25 with chloramphenicol, 10 with ampicillin, 10 without any antibiotic).
- Prepared materials of growth curve experiment, include 8*300ml Luria-Bertani medium.
- Shook and extracted several parts(GFP, RFP, blue chromoprotein, T7 promoter).
- Tested T7 promoter by endonuclease digest and electrophoresis.
- Read papers about recombinase.

- Amplified several parts(BBa_K137058, BBa_K880005, BBa_K404113, BBa_E0420, BBa_K592101).
- Placed an order for PkD46, got a good bargain relatively.
- Determined the homologous sequences of lacZ.

total: 3075bp

A: 42bp:

CGGCAGTAAGGCGGTCTGGGATAGTTTTCTTGCGGCCCT
AATC

B: 42bp:

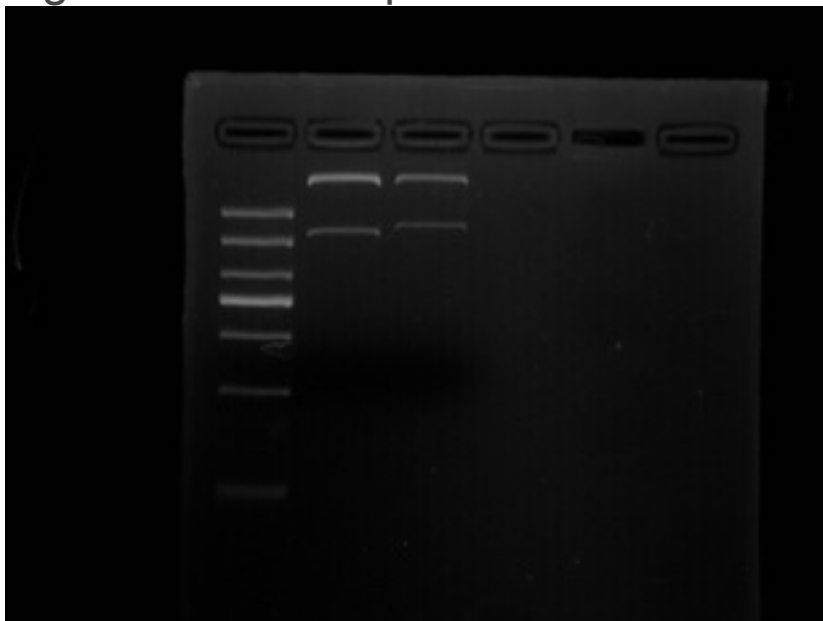
ATGTCGTTATCCAGCGGTGCACGGGTGAACTGATCGCG
CAGC

inter: 317bp

10. Designed primers of the firststep

4th August, 2014 Mon

- Placed an order of genome extraction kit and primers.
- Determined T7 promoter, GFP,RFP by endonuclease digest and electrophoresis.



- Proved that T7 promoter have to be redone. It is not your fault @Sophie.

- Extracted several parts.
- Transformed several parts(BBa_K137058, BBa_K880005, BBa_K404113, BBa_E0420, BBa_K592101)without activation, but failed. Proved that parts with chloromycetin resistance have to be activated before transformation.
- Activated BL21(A+ and A-)
- VR, VF2, pSB1K3-F, pSB1K3-R had been delivered, diluted to 100μmol and 10μmol before use, stored at -20°C.
- PCR: linear plasmid backbones(pSB1K3, pSB1A3, pSB1C3, pSB1T3), but failed.
- Determined the PCR procedure of PrimerSTAR HS polymerase.
*The procedure named 20140804, username is CF. If you want to amplify other sequence, you have to change the annealing temperature and time of elongation (elongating speed of PrimerSTAR is about 1K/min)
- Prepared 12 boxes of pipette tips, autoclave at 121°C for 20 minutes.
- A thorough cleanup of the super clean bench.

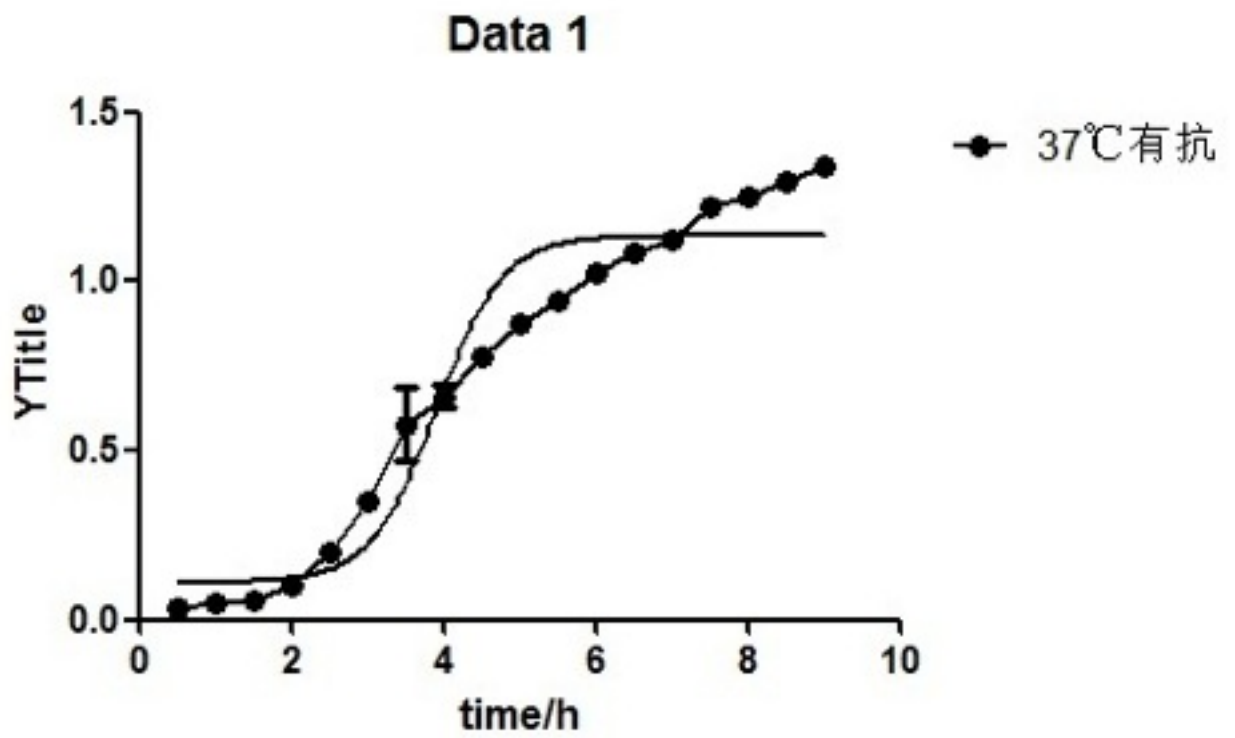
5th August, 2014 Tue

- Prepared 14 boxes of pipette tips, including 2 boxes of 5ml specification and 2*150ml conical flask. Autoclave at 121°C for 20 minutes.
- Prepared 8*250ml of Luria-Bertani medium. Autoclave at 121°C for 20minutes.
- Shook and extracted several parts (BBa_K137058, BBa_K880005, BBa_K404113, BBa_E0420, BBa_K592101)

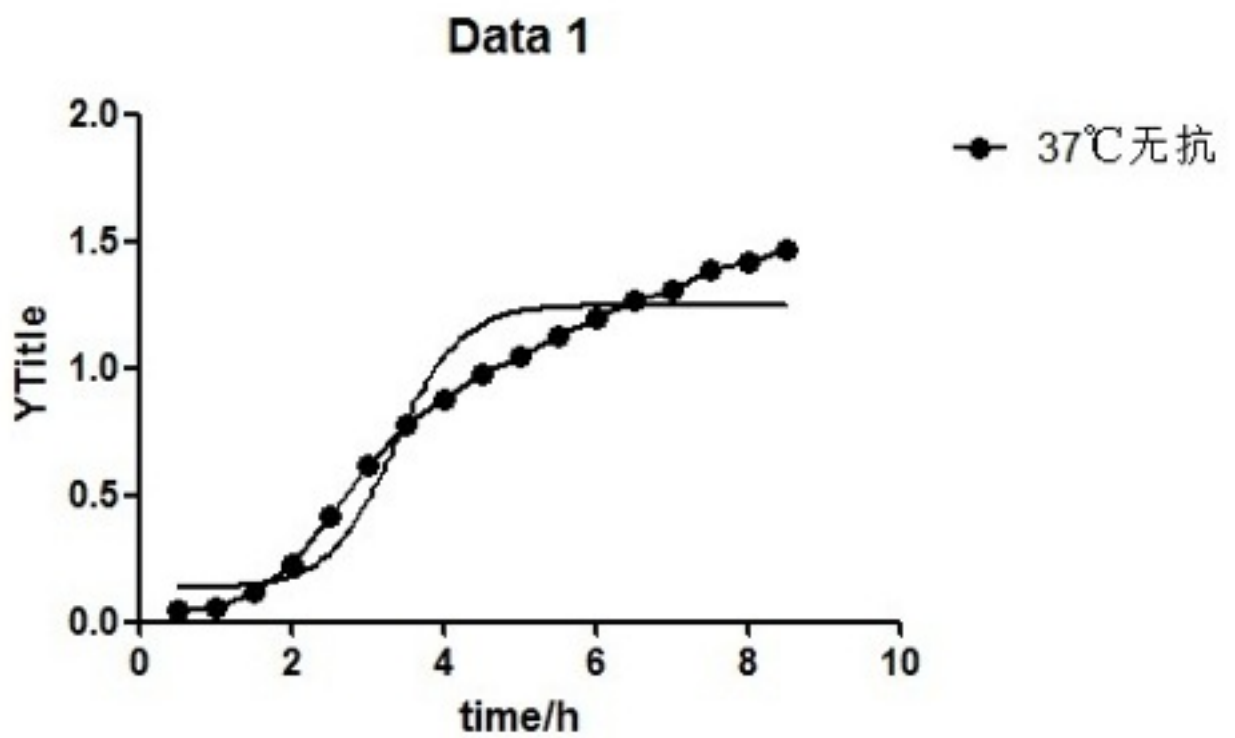
- Determined the PCR procedure of TaKaRa LA Taq polymerase.
*The procedure named 2014 taq, username is CF. If you want to amplify other sequence, you have to change the annealing temperature and time for elongation(elongating speed of LA Taq is about 1K/min)
- Amplified four linear plasmid backbones (pSB1K3, pSB1A3, pSB1C3, pSB1T3), but failed.
- To investigate why, we change the polymerase(borrowed two kinds of enzymes from 411) and template(plasmid of 2012, 2014, 2013), but still failed. This investigation will be continued.
- Bought 2 bottles(can be used insuper clean bench), a packet of tubes used in shaking bacteria, a packet of 1ml pipette tips and 3 lab coats.
- Prepared 75% ethanol solution, 1*TAE solution. Added rayon balls in the bottle of super clean bench.
- A thorough cleanup of 413.

6th August, 2014 Wed

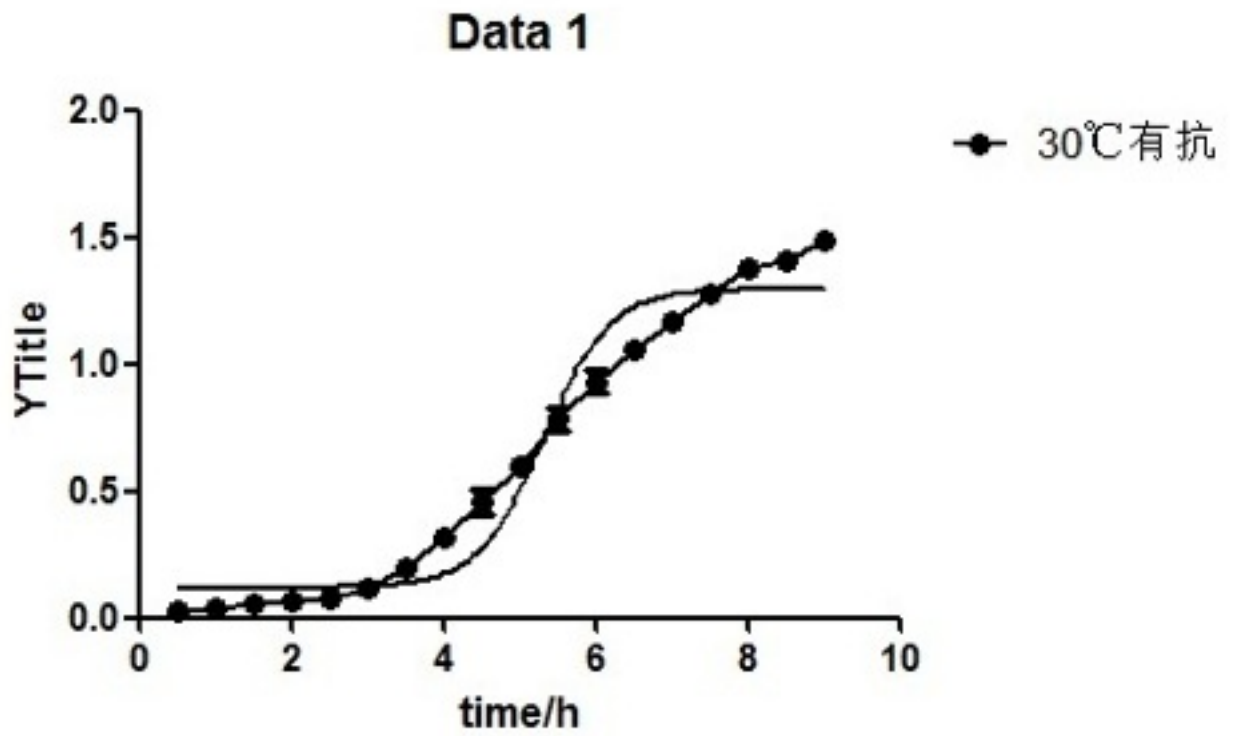
- Explored growth curves of E.coli in different environments. (37°C with ampicillin, 37°C without ampicillin, 30°C with ampicillin, 30°C without ampicillin)
*8 bottles in total. Measure the OD value of every bottle in every 30 minutes. This experiment will last about 16 hours, we will start at 8am.
37°C with ampicillin



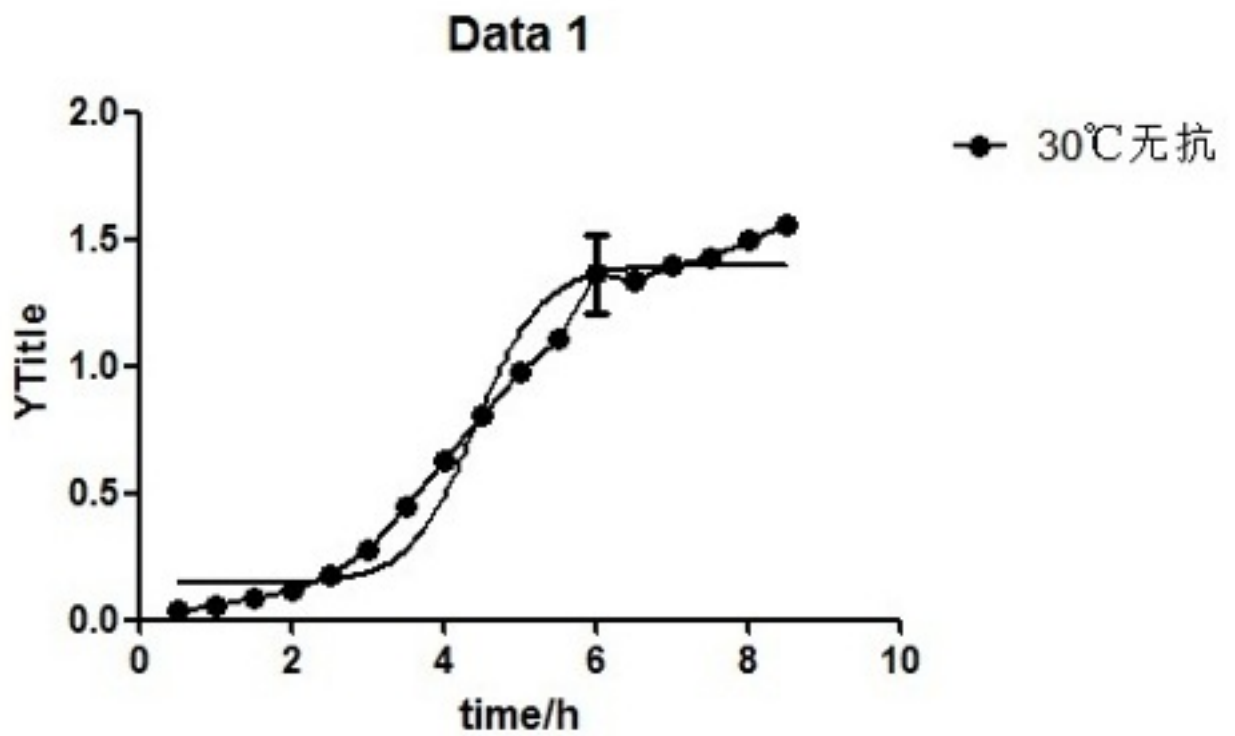
37°C without ampicillin



30°C with ampicillin



30°C without ampicillin



- Prepared materials for competent cell.

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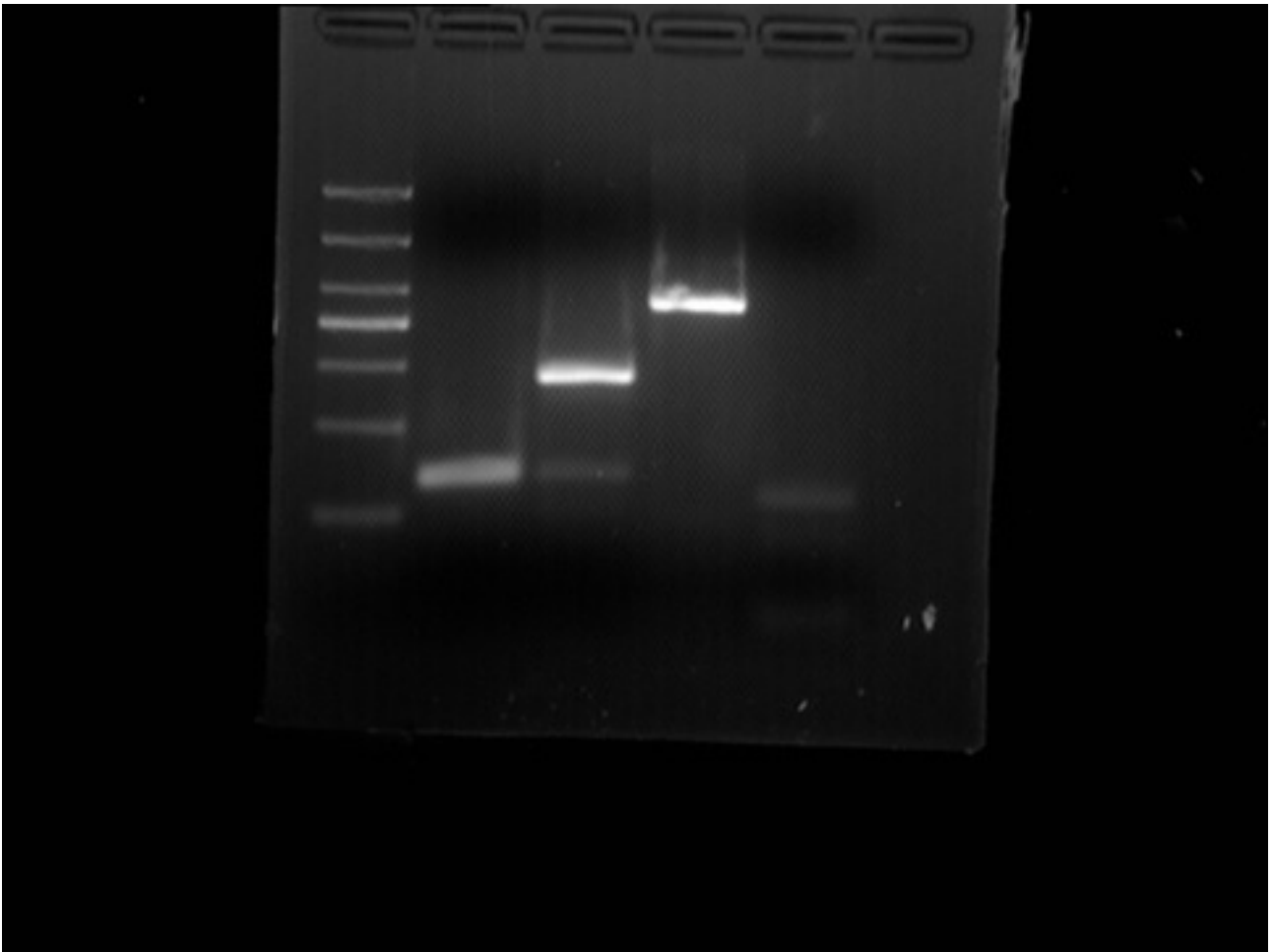
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7th August, 2014 Thu

- Prepared competent cell(BL21,DH5 α).
- Got the reason that why PCR always fail.
Plasmid backbone of RFB



Sequences with 84bp homologous arms(50bp, 150bp, 300bp)



- PKD46 delivered, Pick 8 colony and shook.
- Primers delivered, diluted to 100 μ mol and 10 μ mol before use, stored at -20°C.

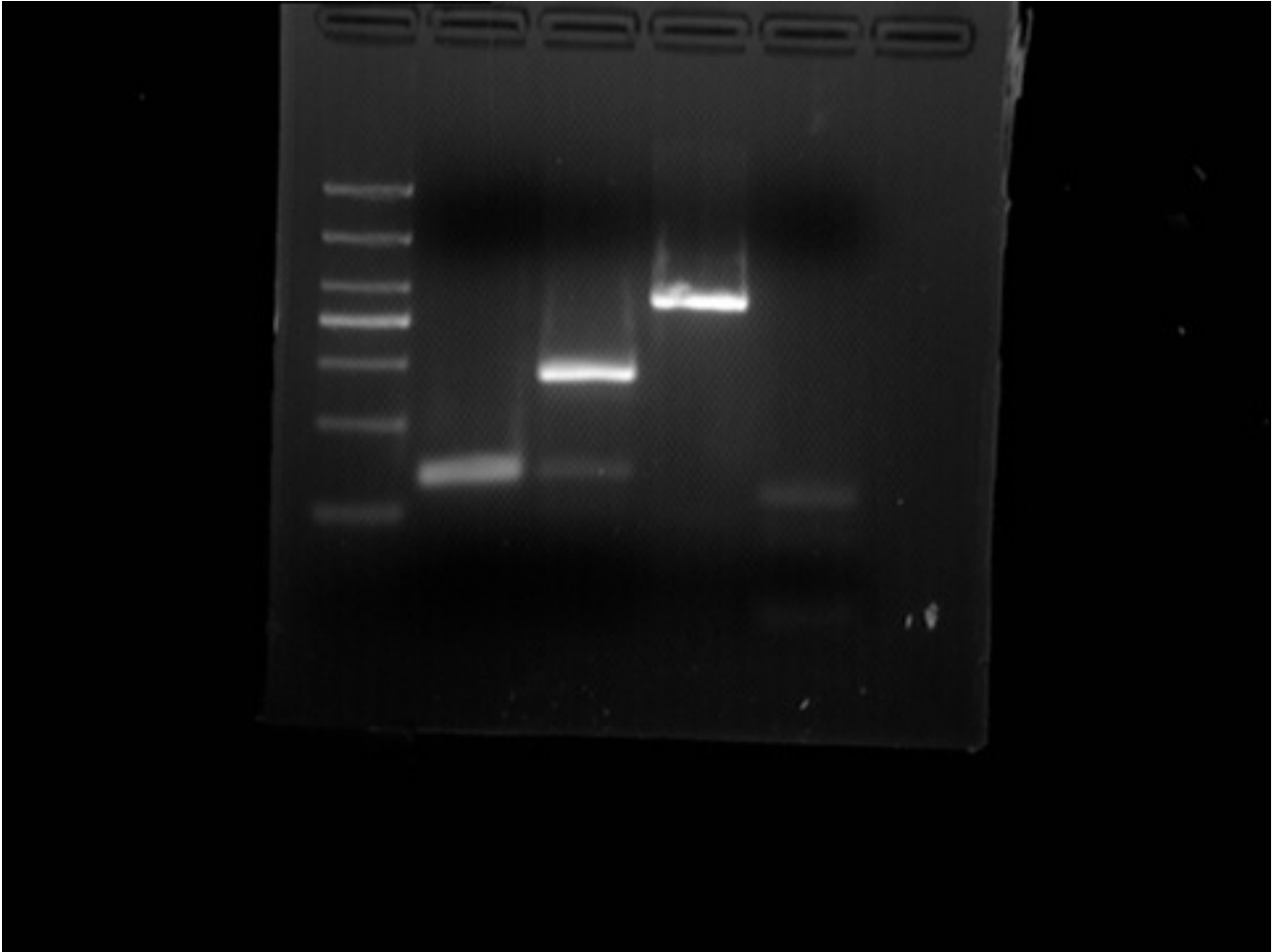
8th August, 2014 Fri

- Extracted plasmids of PKD46.
- Transformed plasmids of PKD46.
- PrimerSTAR HS polymerase delivered.

9th August, 2014 Sat

1. new competent E.coli DH5 α prepared
2. PCR & electrophoresis

3. BL21/pKD46 plate incubation 150bp, 300bp)



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- Primers delivered, diluted to 100 μ mol and 10 μ mol before use, stored at -20°C.

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8th August, 2014 Fri

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- Transformed plasmids of PKD46.
- PrimerSTAR HS polymerase delivered.

● **9th August, 2014 Sat**

- 1. new competent E.coli DH5 α prepared
- 2. PCR & electrophoresis
- 3. BL21/pKD46 plate incubation

10th August, 2014 Sun

- PCR& electrophoresis(1500~5000bp).
- Prepared competent cell of DH5 α .
- Transformed DH5 α (Test) andPKD46.
- Shook PKD46.

11th August, 2014 Mon

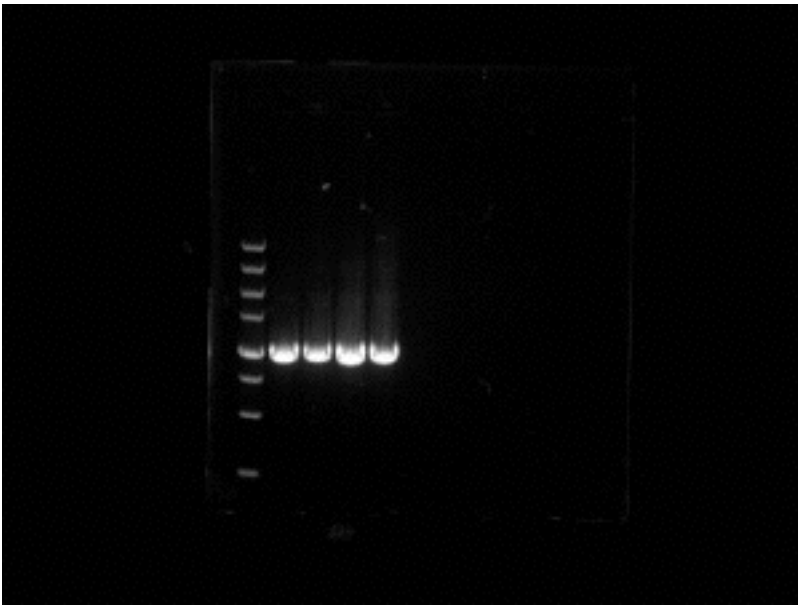
- Designed primers of circuit.
- Prepared LB culture and solidculture with antibiotic.
- Extracted PKD46.
- Digest PKD46 with endonuclease& electrophoresis.

12th August, 2014 Tue

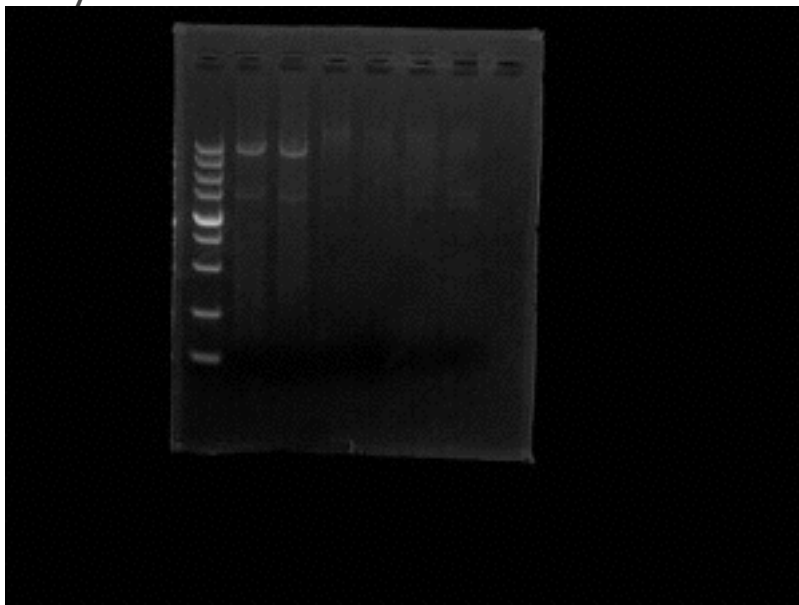
- Prepare competent cell of DH5 α : HanqingLiu, Chaofan Zhang.
- Extract, digest with endonuclease & electrophoresis and transform PKD46: Liangxi Wang, Chuanyun Xu.
- PCR of 50bp, 150bp, 300bp.

20th, August

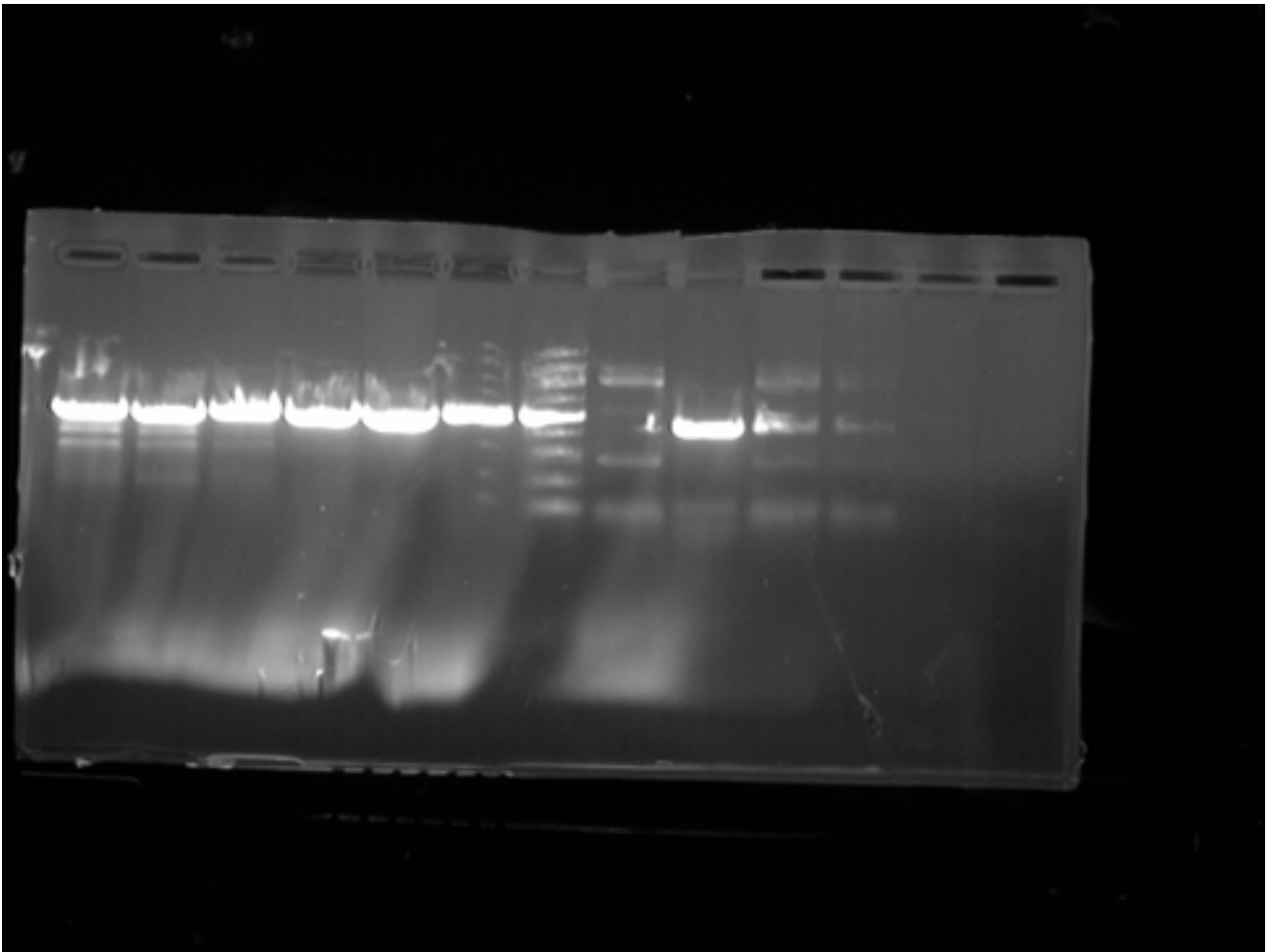
- 1.BL21/pkD46 confirmation:
colony PCR confirmation: check!



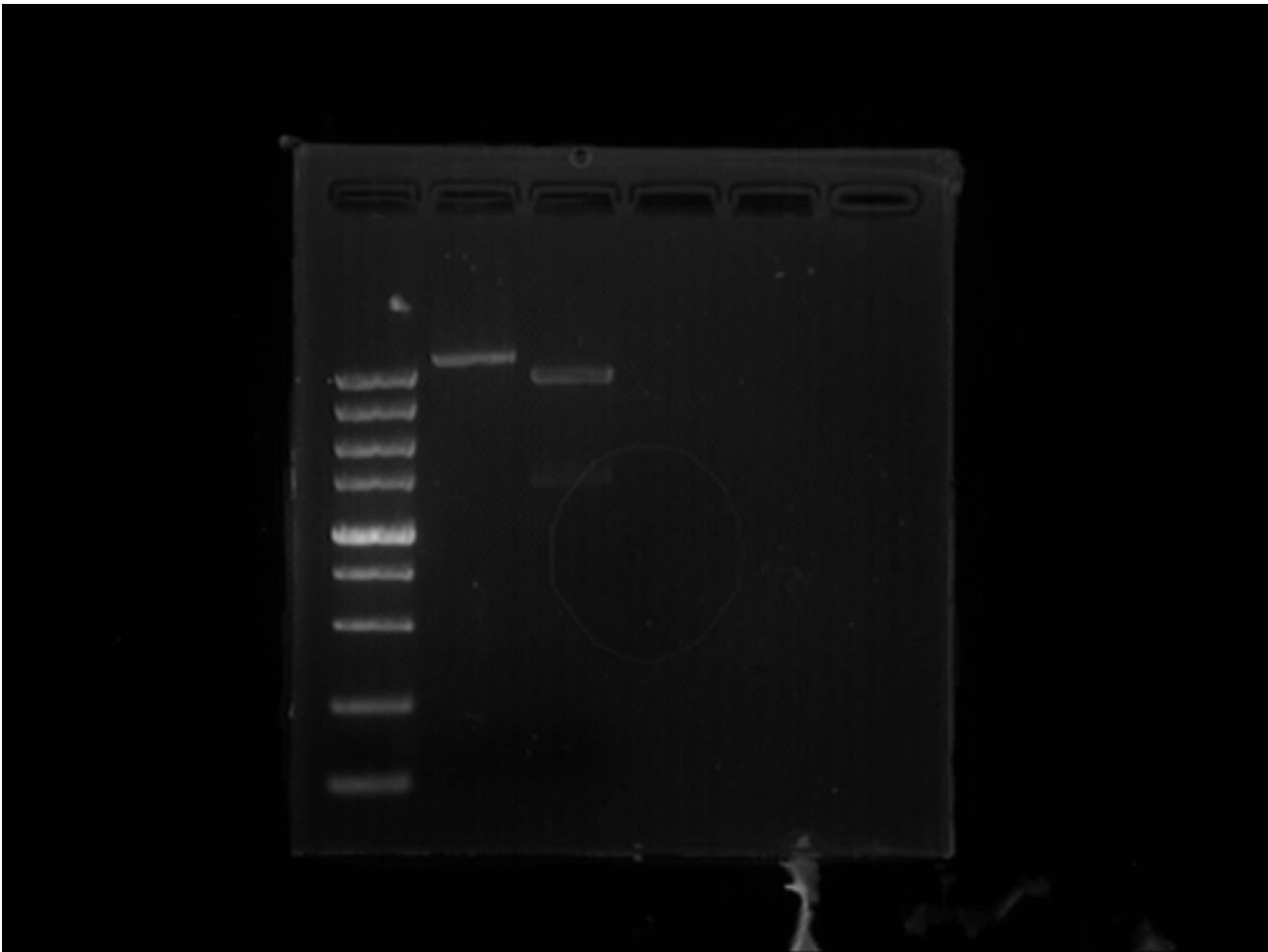
enzyme confirmation: check!



2.pKD46 confirmation:
PCR confirmation: check!



restriction enzyme confirmation: check!



sequence analysis confirmation: check!

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Sequence ID: ic(37639) Length: 508 Number of Matches: 1

Score	Expect	Identifies	Gaps	Strand
1709 bits(925)	0.0	928/928(99%)	1/929(0%)	Plus/Plus
Query 5070	CTGACTTTTACTGTGACGAGTCCGCGCCCTCTGATTTCCAGTCTGACCACTTGGGT			5129
Subject 1	CTG-CTTTTACTGTGACGAGTCCGCGCCCTCTGATTTCCAGTCTGACCACTTGGGT			59
Query 5130	TATCCCTGACGCTTATTGAACTGCTTACTGACTTAAAGCAGCCGTATGACAA			5189
Subject 60	TATCCCTGACGCTTATTGAACTGCTTACTGACTTAAAGCAGCCGTATGACAA			319
Query 5190	CGGGTCTGACCTCGCGGACGAAACTCACTTAAAGGATTTTGGTCAAGAAATAT			5249
Subject 120	CGGGTCTGACCTCGCGGACGAAACTCACTTAAAGGATTTTGGTCAAGAAATAT			379
Query 5250	CAGAAAGGACTTCACTGAGTCCCTTTTAAATGAAATGAGGTTTAAATGAAATGA			5309
Subject 180	CAGAAAGGACTTCACTGAGTCCCTTTTAAATGAAATGAGGTTTAAATGAAATGA			239
Query 5310	GTATATATGCTAAACTGCTTCTGACGCTACTGATCTTAACTTAAAGCAGCCGTAT			5369
Subject 240	GTATATATGCTAAACTGCTTCTGACGCTACTGATCTTAACTTAAAGCAGCCGTAT			299
Query 5370	CAGCAGCTGCTTATTTGCTCAGTCCGAGTACTGACTCCCGCTCTAGAGAACTA			5429
Subject 300	CAGCAGCTGCTTATTTGCTCAGTCCGAGTACTGACTCCCGCTCTAGAGAACTA			359
Query 5430	CGTACGGGAGGCTTACGCTTGGCCGAAATGCTGCAAGGACTCGGAGACCCGACT			5489
Subject 360	CGTACGGGAGGCTTACGCTTGGCCGAAATGCTGCAAGGACTCGGAGACCCGACT			419
Query 5490	CACCGCTTCAGATTTATGACTAAAGCAGTCACTGAGAGGCTGAGGCTCAGAGGT			5549
Subject 420	CACCGCTTCAGATTTATGACTAAAGCAGTCACTGAGAGGCTGAGGCTCAGAGGT			479

Download - Graphics

Sequence ID: I058157 Length: 956 Number of Matches: 1

Range 1: 1 to 956 Graphics Next Match Previous Match

Score	Expect	Identities	Gaps	Strand
1755 bits(950)	0.0	954/956(99%)	0/956(0%)	Plus/Minus
Query 2001	TACTATGATTCMAAACCGAATGATGATCTTAACTCAGGATGATTTTCAA	3000		
Subject 950	TACTATGATTCMAAACCGAATGATGATCTTAACTCAGGATGATTTTCAA	997		
Query 2021	GCATTAACATGACTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	3080		
Subject 096	GCATTAACATGACTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	827		
Query 2041	TTTGTGATGCTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	4040		
Subject 026	TTTGTGATGCTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	777		
Query 4041	ACTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	4100		
Subject 776	ACTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	717		
Query 4301	TCTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	4180		
Subject 736	TCTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	897		
Query 4361	GTAACTCAGGATGATCTTAACTCAGGATGATTTTC	4220		
Subject 056	GTAACTCAGGATGATCTTAACTCAGGATGATTTTC	597		
Query 4201	CTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	4280		
Subject 586	CTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	537		
Query 4281	CTTAACTCAGGATGATCTTAACTCAGGATGATTTTC	4340		
Subject 4281	CTTAACTCAGGATGATCTTAACTCAGGATGATTTTC			

3. blue/white selection--- arabinose induced gradient experiment picked out 3 possible positive clones!





waiting for colony PCR confirmation.....

4.established a practical & appliable colony PCR system