

# Notebook of molecular cloning



Team: GreatBay\_SCIE 2019

Record date: from 7.12-8.31

7.12 (Fri)

## Number of single colonies on plates (operator: Diana)

1 on SM011

2 on SM004, SM006

0 on SM005, SM007

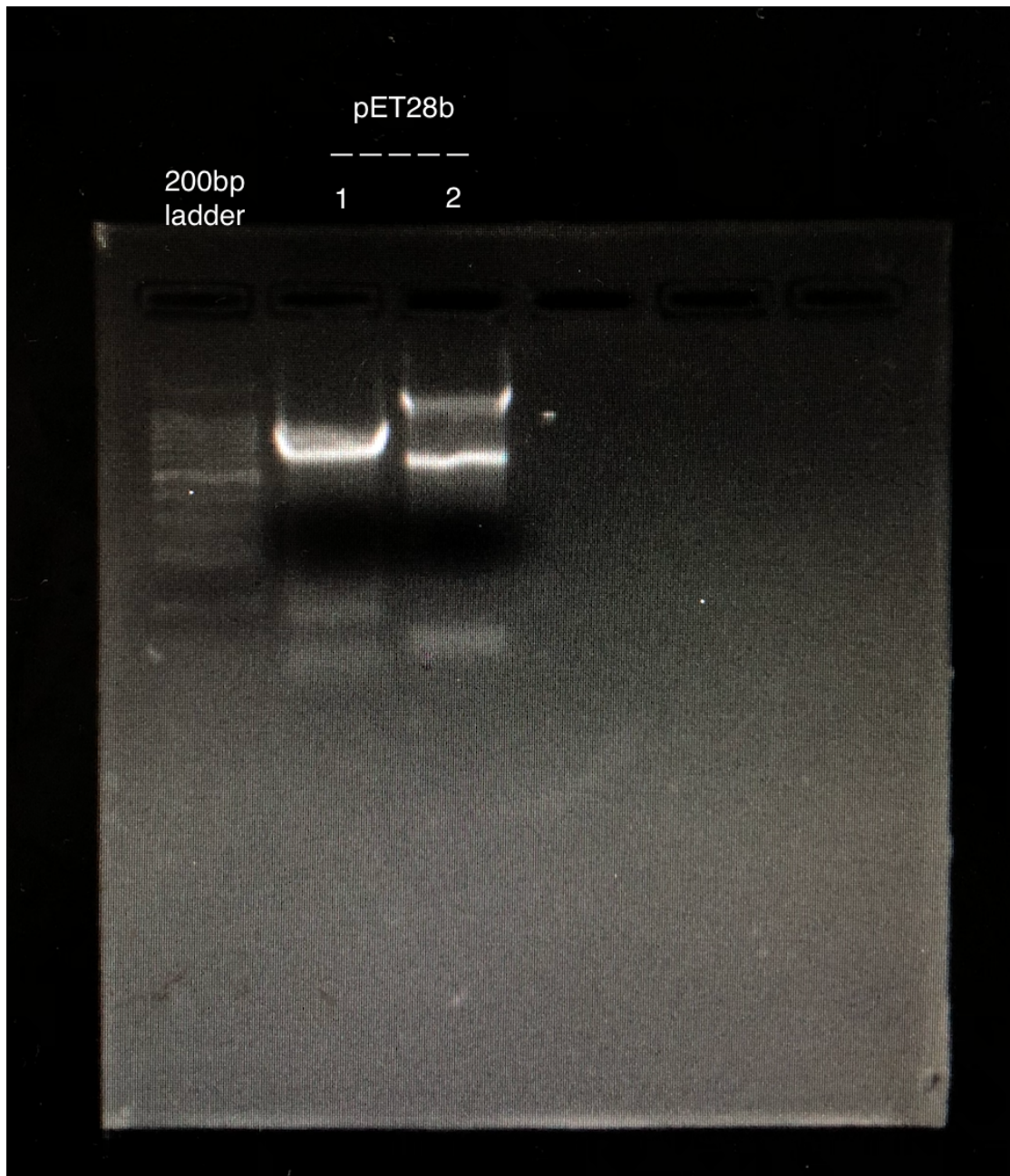
## PCR Template (operator: Diana, Aislinn)

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR		
Primer 1	SP002-F	2.5
Primer 2	SP002-R	2.5
Template	SM010	0.5 (77.9ng/ ul)
H2O		19.5
DNA polymerase	rTaq	25
All		50
		50/tubex1
Product length	1900bp	

PCR		
Primer 1	SP003-F	2.5
Primer 2	SP003-R	2.5
Template	SM010	0.5
H2O		19.5
DNA polymerase	rTaq	25
All		50
		50/tubex1
Product length	3400bp	

Process		
1	94	60s
2	94	30s
3	55	30s
4	72	105s
Go to 2	35 cycle	
5	72	5min
6	16	$\infty$



Gel extraction (operator: Diana, Aislinn)

- Gel extraction
- Liquid extraction

Elution buffer

- EB
- H2O

Elution volume: 40ul

Name: pET28b-1,2

Concentration(A260/280nm): 64.85ng/ $\mu$ L(pET28b-1), 42.95ng/ $\mu$ L(pET28b-2)

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### Reconstruction of SM005, SM007

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	pET28b-1	1
Fragment 2	pET28b-2	1.5
Fragment 3	SM005 Fr3	0.5
Fragment 4	SM005 Fr4	1
Fragment 5	SM005 Fr5	1
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Backbone		
Fragment 1	pET28b-1	1.3
Fragment 2	pET28b-2	2
Fragment 3	SM007 Fr3	0.7
Fragment 4	SM007 Fr4	1
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Process		
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1	50	60min
2	12	∞

### Transformation (operator: Diana)

- Electroporation 1 $\mu$ L
- Chemical transformation 10 $\mu$ L

Host: DH5 alpha  
 Plasmid to construct: SM005, SM007

Recovery medium: LB  
 Volume of bacterial liquid to spread: 150ul  
 Agar plate and antibiotic to use: LB Cm  
 Transformation conditions: heat shock at 42°C for 90s

### 7.13 (Sat)

#### Number of single colonies on plates (operator: Diana)

0 on all.

### 7.15 (Mon)

#### Construction of SM004, SM006, SM011

#### Colony selection (operator: Diana)

- From
- Agar plates
  - Glycerol stock
  - Liquid starter

Strain	SM004, SM006, SM011
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Antibiotics resistance	Kan
Medium	LB
Volume	Inoculating single colonies on plates
Temperature	37
Number	5

SM004: 1, 2

SM006: 3, 4

SM011: 5

**PCR Template: (operator: Diana)**

- Purified plasmid
- Purified genome
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR for verification		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM004 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1551bp	

PCR for verification		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM006 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1779bp	

PCR for verification		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM011colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1446bp	

Process		
1	94	10min
2	94	30s
3	55	30s
4	72	1min
Go to 2	35 cycle	
5	72	5min

6	16	$\infty$
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control: pET28b, 585bp



### Constructing SM003, SM012, SM013

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	pET28b-1	1.2
Fragment 2	pET28b-2	1.8



Fragment 3	SM003 Fr3	1
Fragment 4	SM003 Fr4	1
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Backbone		
Fragment 1	pET28b-1	1.5
Fragment 2	pET28b-2	2.5
Fragment 3	SM012 Fr3	1
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Backbone		
Fragment 1	pET28b-1	1
Fragment 2	pET28b-2	1.5
Fragment 3	SM003 Fr3	0.8
Fragment 4	SM005 Fr4	1
Fragment 5	SM004 Fr4	0.7
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Process		
1	50	60min
2	12	∞

## Transformation (operator: Diana, Uranium)

Electroporation 1 $\mu$ L

Chemical transformation 10 $\mu$ L

Host: DH5 alpha

Plasmid to construct: SM003, SM012, SM013

Recovery medium: LB

Volume of bacterial liquid to spread: 200ul

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

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## Construction of SM005 and SM007

<u>Gibson assembly</u>	<u>(operator: XYZ)</u>	
Backbone		
Fragment 1	pET28b-1	1
Fragment 2	pET28b-2	2
Fragment 3	SM005 Fr3	2
Fragment 4	SM005 Fr4	2.5
Fragment 5	SM005 Fr5	2.5
H <sub>2</sub> O		Add to 10
GA mix		10
All		20

Backbone		
Fragment 1	pET28b-1	0.5
Fragment 2	pET28b-2	1.5
Fragment 3	SM005 Fr3	1

Fragment 4	SM005 Fr4	3.5
Fragment 5	SM005 Fr5	3.5
H <sub>2</sub> O		Add to 10
GA mix		10
All		20

Backbone		
Fragment 1	pET28b-1	1
Fragment 2	pET28b-2	2
Fragment 3	SM007 Fr3	1
Fragment 4	SM007 Fr4	6
H <sub>2</sub> O		Add to 10
GA mix		10
All		20

Process		
1	50	60min
2	12	∞

### **Transformation (operator: XYZ)**

- Electroporation 1 $\mu$ L
- Chemical transformation 20 $\mu$ L

Host: DH5 alpha

Plasmid to construct: SM005, SM007

Recovery medium: LB

Volume of bacterial liquid to spread: 300ul

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

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## Culture preservation of Se004 (Start)

### Inoculation (operator: Diana)

From

Agar plates

Glycerol stock

Liquid starter

Strain	Se004
Antibiotics	Amp
Medium	LB
Volume	3mL in 15mL tube, single colony
Temperature	37
Number	1

## 7.16 (Tues)

### Construction of SM004, SM006, SM011

#### PCR Template: (operator: Diana)

Purified plasmid

Purified genome

Single colony

Bacteria liquid

Boiled Bacteria cell lysate

PCR for verification		
Primer 1	SP004-F	1

Primer 2	SP004-R	1
Template	SM004 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1551bp	

PCR for verification		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM006 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1779bp	

PCR for verification		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM011colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1446bp	

Process		
1	94	10min
2	94	30s
3	55	30s
4	72	1min
Go to 2	35 cycle	
5	72	5min
6	16	$\infty$

control: pET28b, 585bp



## Inoculation (operator: Diana)

From

Agar plates

Glycerol stock

Liquid starter

Strain	DH5alpha {SM004(1, 2), SM006(3, 4), SM011(5)}
Antibiotics	Kan
Medium	LB
Volume	4mL in 15mL tube, single colony
Temperature	37
Number	5

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## **Construction of SM003, SM012, SM013**

## **Construction of SM005, SM007**

### Number of single colonies on plates

SM003: 9

SM012: 6

SM013: 9

SM007: 5

SM005: 0

SM011: 0

---

## **Culture preservation of Se004 (Done)**

### Preparing bacterial glycerol stocks(operator: Diana)

Strain name: DH5 $\alpha$  (pMV-Cfa[C]-Cfa[N])

Number in stocks: S014

## 7.17 (Wed)

### Construction of SM003, SM012, SM013

### Construction of SM005, SM007

#### Colony selection (operator: Diana)

From

√Agar plates

Glycerol stock

Liquid starter

Strain	DH5alpha{SM003, SM012, SM013, SM007}
Antibiotics resistance	Kan
Medium	LB
Volume	Inoculating single colonies on plates
Temperature	37
Number	2 for each, 8 in total

#### PCR Template (operator: Diana)

Purified plasmid

Purified genome

√Single colony

Bacteria liquid

Boiled Bacteria cell lysate

PCR for verification		
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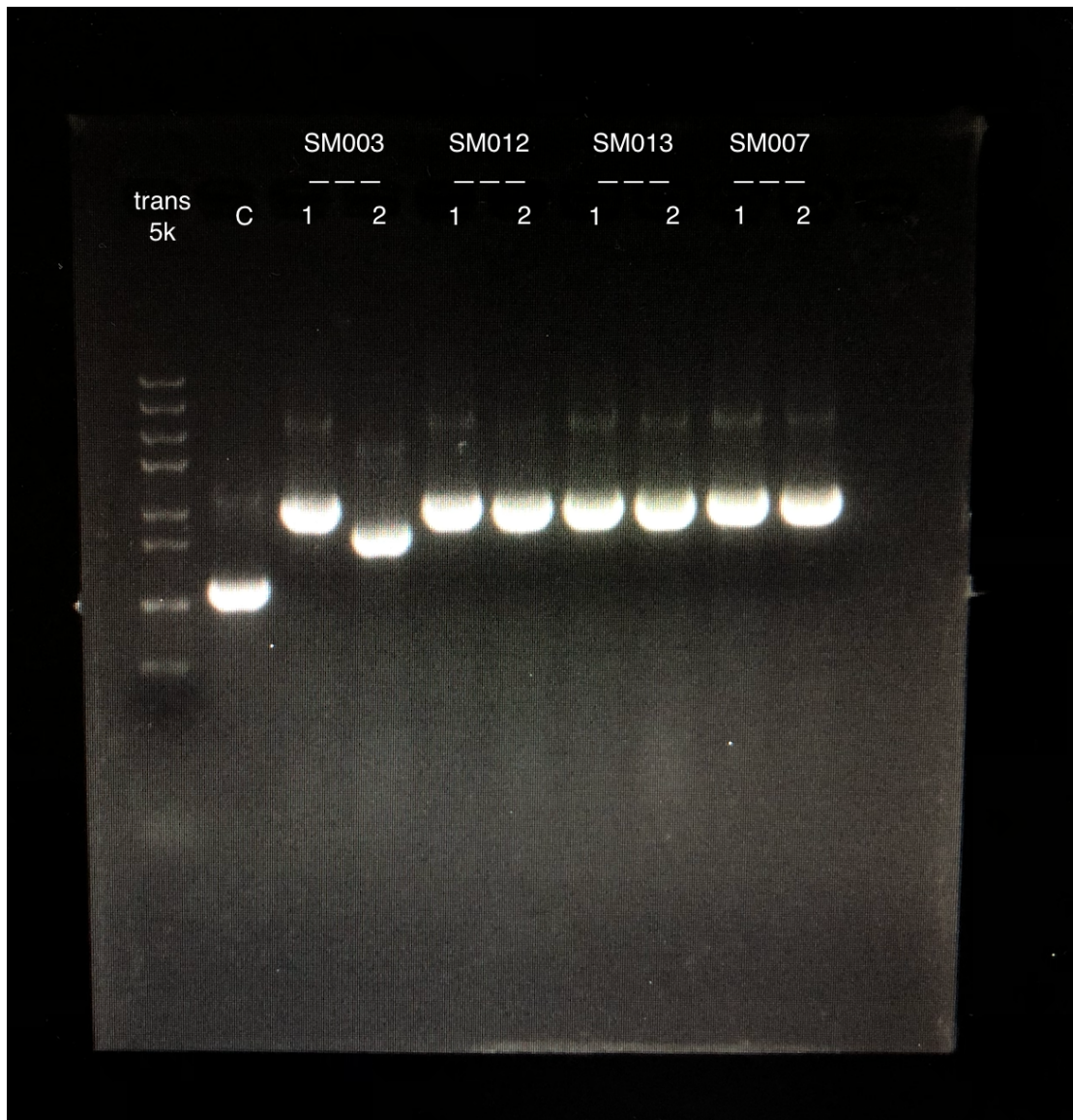
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM003 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	852bp	

Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM012 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1473bp	

Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM013 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1143bp	

Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM007 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex2
Product length	1209bp	

Process		
1	94	10min
2	94	30s
3	55	30s
4	72	1min
Go to 2	35 cycle	
5	72	5min
6	16	∞



## Construction of SM015 (Start)

### PCR Template (Operator: Adonie)

- Purified plasmid
- Purified genome
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

Fragment 1, rTaq

Components		
Primer 1	SP020	1.5
Primer 2	SP021	1.5
Template	SM015 (pACYC-duet1- mTyr-CNK)	0.5
H2O		10.5
DNA polymerase	Taq Mix	15
All		30
		30
Product length	1973 bp	

#### Fragment 2, rTaq

Components		
Primer 1	SP019	1.5
Primer 2	SP022	1.5
Template	SM015 (pACYC-duet1- mTyr-CNK)	0.5
H2O		10.5
DNA polymerase	Taq Mix	15
All		30
		30
Product length	2952 bp	

#### Fragment 1, FastPfu

Components		
Primer 1	SP020	1.5
Primer 2	SP021	1.5
Template	SM015 (pACYC-duet1- mTyr-CNK)	0.5

H2O		10.5
DNA polymerase	FastPfu 2x Mix	15
All		30
		30
Product length	1973 bp	

Fragment 2, FastPfu

Components		
Primer 1	SP019	1.5
Primer 2	SP022	1.5
Template	SM015 (pACYC-duet1- mTyr-CNK)	0.5
H2O		10.5
DNA polymerase	FastPfu 2x Mix	15
All		30
		30
Product length	2952 bp	



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## Transformation of Sy003, SM002, SM008, SM010 into Rosetta (Start)

### Transformation (operator: Diana)

- Electroporation 1  $\mu$ L
- Chemical transformation 1  $\mu$ L

Host: Rosetta(DE3)

Plasmid to construct: Sy003, SM002, SM008, SM010

Recovery medium: LB

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Cm, Kan

Transformation conditions: heat shock at 42°C for 90s

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## Construction of SM004, SM006, SM011

### Preparing bacterial glycerol stocks(operator: Adonie)

Strain name: SM004, SM006, SM011

Number in stocks: S015, S016, S017

Collection of cell pellet

Strain Name: SM004, SM006, SM011

- Centrifuge at 12000 rpm for 2 min and discard waste fluid.
- Cell pellets are stored at -20°C

### Sequencing (operator: Diana)

Sample	Primer	Requirements
<input type="checkbox"/> bacterial liquid <input type="checkbox"/> Plasmid <input type="checkbox"/> PCR purified <input checked="" type="checkbox"/> PCR unpurified	SP004-F SP004-R	<input type="checkbox"/> Forward <input type="checkbox"/> Reverse <input checked="" type="checkbox"/> Both

SM004 (1): 0 mismatches, 0 gaps/insertions. Correct.

SM004 (2): 0 mismatches, 2 gaps/insertions. Correct.

SM006 (3): Not of sufficient quality to be used for alignment.

SM006 (4): 1 mismatch, 1 gap/insertion(1 Mfp5 absent).

SM011 (5): 1 mismatch, 3 gaps/insertions(1 Mfp3 absent)

## 7.18 (Thur)

### Construction of SM015

#### Gel extraction (operator: Diana)

√Gel extraction

Liquid extraction

Elution buffer

EB

H<sub>2</sub>O

Elution volume: 40ul

Name: SM015(1), SM015(2)

Concentration(A<sub>260</sub>/280nm): 51.8ng/ul, 41.9ng/ul

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	SM015-1	2
Fragment 2	SM015-2	3
H <sub>2</sub> O		Add to 5
GA mix		5
All		10
Process		
1	50	60min
2	12	∞

#### Transformation (operator: Uranium)

Electroporation 1μL

√Chemical transformation 10μL

Host: DH5alpha, 100 μL

Plasmid to construct: SM015

Recovery medium: LB, 300 μL



Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Cm

Transformation conditions: heat shock at 42°C for 90s

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## Transformation of Sy003, SM002, SM008, SM010 into Rosetta

### Inoculation (operator: Diana)

From

Agar plates

Glycerol stock

Liquid starter

Strain	Rosetta {Sy003, SM002, SM008, SM010}
Antibiotics	Kan, Cm
Medium	LB
Volume	4mL in 15mL tube, single colony
Temperature	37
Number	4

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## Culture preservation of Sy003

### Inoculation (operator: Diana)

From

Agar plates

Glycerol stock

Liquid starter

Strain	DH5 $\alpha$ {Sy003}
Antibiotics	Kan
Medium	LB

Volume	4mL in 15mL tube, single colony
Temperature	37
Number	1

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## Culture preservation of Sy002 (Done)

### Preparing bacterial glycerol stocks(operator: Diana)

Strain name: DH5 $\alpha$  {pACYC-duet1-mTyr-CNK-7his}

Number in stocks: S018

## 7.19 (Fri)

### Construction of SM003, SM012, SM013

### Construction of SM005, SM007

### Sequencing (operator: Diana)

Sample	Primer	Requirements
<input type="checkbox"/> Bacterial liquid <input type="checkbox"/> Plasmid <input type="checkbox"/> PCR purified <input checked="" type="checkbox"/> PCR unpurified	SP004-F SP004-R	<input type="checkbox"/> Forward <input type="checkbox"/> Reverse <input checked="" type="checkbox"/> Both

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SM003: 1 mismatch, 3 gaps/insertions. Mismatch due to mutation.

SM012: Mfp5-Cfa[N] absent.

SM013: Cfa[C]-Mfp5-Cfa[N] absent.

SM007: 1 mfp3 absent.

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## **Transformation of Sy003, SM002, SM008, SM010 into Rosetta (Done)**

### **Preparing bacterial glycerol stocks(operator: XYZ)**

Strain name: Rosetta{Sy003, SM002, SM008, SM010}

Number in stocks: S016, S017, S019, S020

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## **Culture preservation of Sy003 (Done)**

### **Preparing bacterial glycerol stocks(operator: XYZ)**

Strain name: DH5 $\alpha$ {Sy003}

Number in stocks: S021

## **7.22 (Mon)**

### **Construction of SM003**

#### **Colony selection (operator: Adonie)**

From

√Agar plates

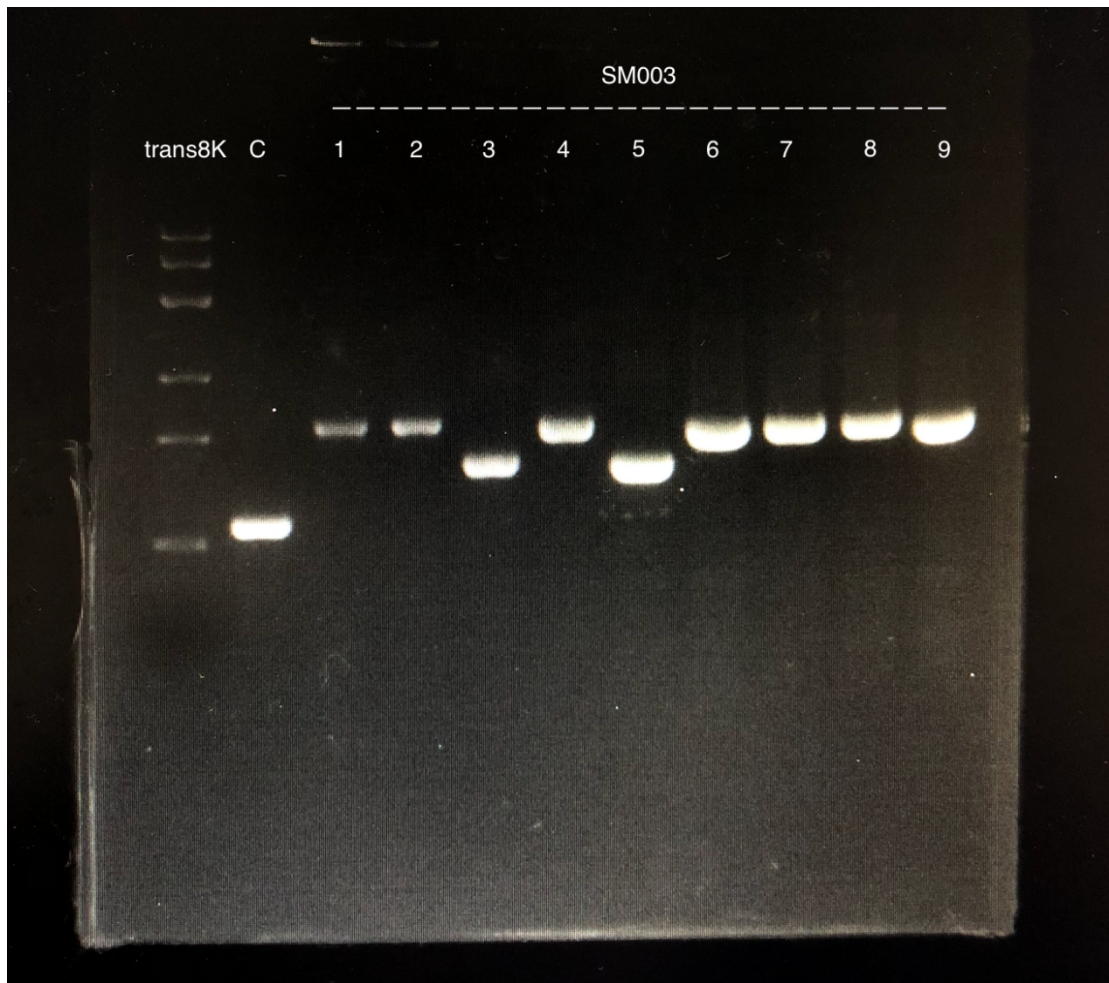
- Glycerol stock
- Liquid starter

Strain	DH5alpha{SM003}
Antibiotics resistance	Kan
Medium	LB
Volume	Inoculating single colonies on plates
Temperature	4
Number	9

**PCR Template (operator: Adonie)**

- Purified plasmid
- Purified genome
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR for verification		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	SM003 colony	
H2O		13
DNA polymerase	Taq Mix	15
All		30
		30/tubex8
Product length	852 bp	



## Transformation of SM004 into BL21(DE3) (Start)

### Plasmid extraction (operator: Adonie)

strain	DH5α {SM004}
plasmid	SM004

Elution buffer

EB

H<sub>2</sub>O

Elution volume: 50ul

Concentration: 156.3ng/ul

### Transformation (operator: Diana)

Electroporation 1μL

√Chemical transformation 0.5 $\mu$ L

Host: BL21(DE3), 100  $\mu$ L

Plasmid to construct: suicide plasmid (P194→Plux)

Recovery medium: LB, 300  $\mu$ L

Volume of bacterial liquid to spread: 400  $\mu$ L

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

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## Construction of SM012, SM013 (Start)

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	pET28b-1	0.5
Fragment 2	pET28b-2	1.5
Fragment 3	SM012 Fr3	1.7
Fragment 4	SM004 Fr4	1.3
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Backbone		
Fragment 1	pET28b-1	0.5
Fragment 2	pET28b-2	1.2
Fragment 3	SM003 Fr3	1.1
Fragment 4	SM005 Fr4	1.2
Fragment 5	SM004 Fr4	1

H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Process		
1	50	60min
2	12	∞

**Transformation (operator: Diana)**

- Electroporation 1 $\mu$ L
- Chemical transformation 10 $\mu$ L

Host: DH5alpha, 100  $\mu$ L

Plasmid to construct: SM012, SM013

Recovery medium: LB, 300  $\mu$ L

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

**7.23 (Tues)**

**Construction of SM014 (Start)**

**PCR Template (operator: Diana)**

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

rTaq

PCR		
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Primer 1	SP016	1.5
Primer 2	SP017	1.5
Template	Sy005	0.5(30ng/ul)
H2O		11.5
DNA polymerase	rTaq Mix	15
All		30
		30/tubex1
Product length	581bp	

Primer 1	SP015	1.5
Primer 2	SP005-Mfp5-R	1.5
Template	SM010	0.5(77.9ng/ul)
H2O		11.5
DNA polymerase	rTaq Mix	15
All		30
		30/tubex1
Product length	301bp	

Process		
1	94	10s
2	94	30s
3	55	30s
4	72	30s
Go to 2	35 cycle	
5	72	2min
6	12	∞

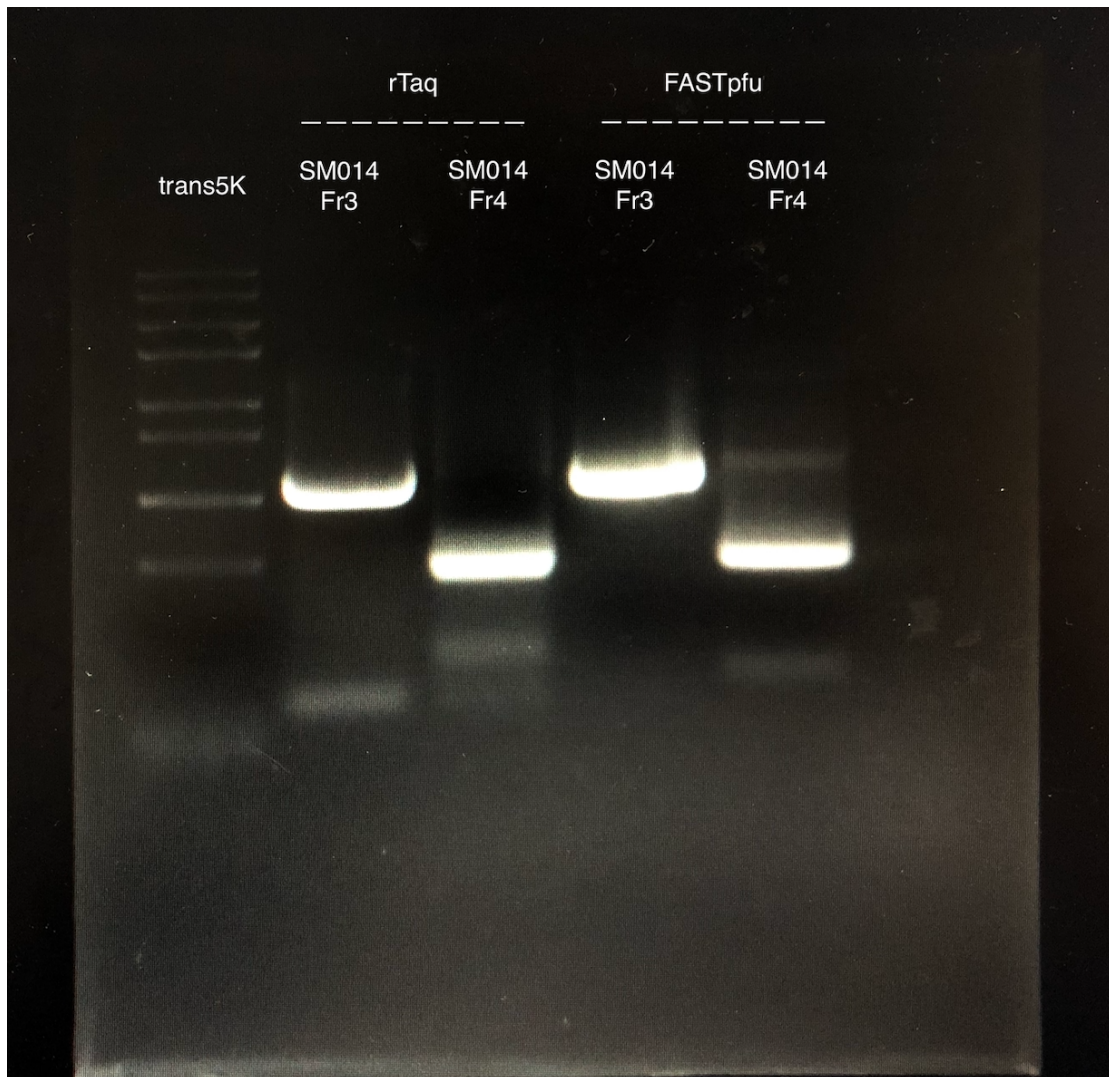
FASTpfu



PCR		
Primer 1	SP016	1
Primer 2	SP017	1
Template	Sy005	0.5(30ng/ul)
H2O		18.5
DNA polymerase	FASTpfu Mix	9
All		30
		30/tubex1
Product length	581bp	

Primer 1	SP015	1
Primer 2	SP005-Mfp5-R	1
Template	SM010	0.5(77.9ng/ul)
H2O		18.5
DNA polymerase	FASTpfuMix	9
All		30
		30/tubex1
Product length	301bp	

Process		
1	98	1min
2	98	30s
3	55	30s
4	72	30s
Go to 2	32 cycle	
5	72	2min
6	12	∞



**Gel extraction (operator: Adonie)**

- Gel extraction  
 Liquid extraction

**Elution buffer**

- EB  
 H2O

Elution volume: 30ul

Name: SM014 Fr3

Concentration(A260/280nm): 37.3ng/ul

Name: SM014 Fr4

Concentration(A260/280nm): 8.35ng/ul

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	pET28b-1	0.5
Fragment 2	pET28b-2	1.3
Fragment 3	SM014 Fr3	0.5
Fragment 4	SM014 Fr4	2.7
H <sub>2</sub> O		Add to 5
GA mix		5
All		10
Process		
1	50	60min
2	12	∞

### Transformation (operator: Uranium, Diana)

Electroporation 1  $\mu$ L

Chemical transformation 10  $\mu$ L

Host: DH5alpha

Plasmid to construct: SM014

Recovery medium: LB, 300  $\mu$ L

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

### **Construction of SM015**

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		

Fragment 1	SM015 Fr1	1.8
Fragment 2	SM015 Fr2	3.2
H <sub>2</sub> O		Add to 5
GA mix		5
All		10
Process		
1	50	60min
2	12	∞

### Transformation (operator: Uranium)

Electroporation 1 $\mu$ L

Chemical transformation 10 $\mu$ L

Host: Rosetta(DE3)

Plasmid to construct: SM015

Recovery medium: LB, 300ul

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Cm

Transformation conditions: heat shock at 42°C for 90s

### Construction of SM003

#### Sequencing (operator: Diana)

Sample	Primer	Requirements
<input type="checkbox"/> 菌液 <input type="checkbox"/> 质粒 <input type="checkbox"/> PCR纯化 <input checked="" type="checkbox"/> PCR未纯化	SP004-F SP004-R	<input type="checkbox"/> Forward <input type="checkbox"/> Reverse <input checked="" type="checkbox"/> Both

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3: 1 mismatch, 1 insertion, no mutation, correct.

4:

5: 1 mismatch due to mutation, 2 gaps, incorrect.

**Inoculation (operator: Uranium)**

From

Agar plates

Glycerol stock

Liquid starter

Strain	SM003
Antibiotics	Kan
Medium	LB
Volume	3mL in 15mL tube, single colony
Temperature	37
Number	1

**Transformation of SM004 into BL21(DE3)**

**Number of single colonies on plates (operator:Diana): 3**

**Inoculation (operator: Uranium)**

From

Agar plates

Glycerol stock

Liquid starter

Strain	SM004
Antibiotics	Kan
Medium	LB

Volume	3mL in 15mL tube, single colony
Temperature	37
Number	1

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## Construction of SM012, SM013

Number of single colonies on plates (operator:Diana): 0 on both plates

7.24 (Wed)

## Construction of SM014

Number of single colonies on plates (operator:Diana): 0

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## Construction of SM015

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	SM015 Fr1	1.8
Fragment 2	SM015 Fr2	3.2
H <sub>2</sub> O		Add to 5
GA mix		5
All		10

Process		
1	50	60min
2	12	∞

### Transformation (operator: Darcy)

Electroporation 1 $\mu$ L

Chemical transformation 10 $\mu$ L

Host: DH5alpha

Plasmid to construct: SM015

Recovery medium: LB, 300ul

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Cm

Transformation conditions: heat shock at 42°C for 90s

## Transformation of SM003 into BL21(DE3)

### Inoculation (operator: Uranium)

From

Agar plates

Glycerol stock

Liquid starter

Strain	SM003(S022)
Antibiotics	KAN
Medium	LB
Volume	3mL in 15mL tube, single colony
Temperature	37
Number	1

## Culture preservation of Se005

### Inoculation (operator: Uranium)

From

Agar plates

Glycerol stock

Liquid starter

Strain	SE005
Antibiotics	KAN
Medium	LB
Volume	3mL in 15mL tube, single colony
Temperature	37
Number	1

---

## Transformation of Se005 into rosetta

### Transformation (operator: Darcy)

Electroporation 1 $\mu$ L

Chemical transformation 10 $\mu$ L

Host: BL21(DE3) rosetta

Plasmid to construct: Se005

Recovery medium: LB, 300ul

Volume of bacterial liquid to spread: 100  $\mu$ L

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

---

## Culture preservation of SM003 (Done)

### Preparing bacterial glycerol stocks (operator: Diana)



Strain name: DH5α {SM003}

Number in stocks: S022

---

## Transformation of SM004 into BL21(DE3) (Done)

### Preparing bacterial glycerol stocks (operator: Diana)

Strain name: BL21(DE3){SM004}

Number in stocks: S023

7.25 (Thur)

## Construction of SM015

Number of single colonies on plates (operator:Diana): 0

---

## Transformation of SM003 into BL21(DE3)

### Plasmid extraction (operator: Diana, Adonie)

strain	DH5α {SM003}
plasmid	SM003

Elution buffer

TB

H2O

Elution volume: 50uL

Concentration(A260/280): 80.75ng/μL

### Transformation (operator: Diana)

Electroporation 1μL

✓Chemical transformation 10 $\mu$ L

Host: BL21(DE3)

Plasmid to construct: SM003

Recovery medium: LB, 300ul

Volume of bacterial liquid to spread: 300  $\mu$ L

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

---

## Construction of SM012, SM013

### PCR Template (operator: Diana, Adonie)

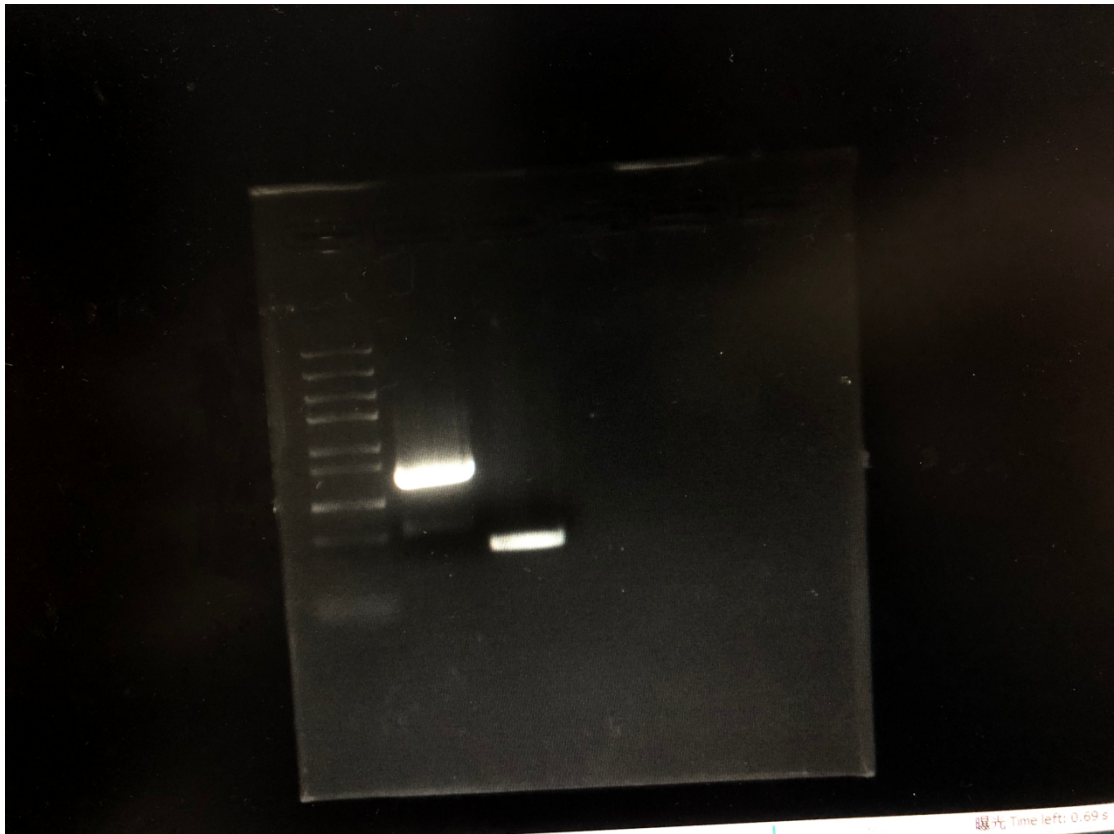
- Purified genome
- ✓ Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR		
Primer 1	SP014	1
Primer 2	SP001 CsgA-Mfp5-R	1
Template	SM010	0.5
H2O		18.5
DNA polymerase	FASTpfu	9
All		30
		30/tubex1
Product length	690bp	

Primer 1	SP005 Mfp5-CfaC-F	1
----------	-------------------	---

Primer 2	SP001 CsgA-Mfp5-R	1
Template	SM010	0.5
H2O		18.5
DNA polymerase	FASTpfu	9
All		30
		30/tubex1
Product length	228bp	

Process		
1	98	1min
2	98	30s
3	55	30s
4	72	30s
Go to 2	35 cycle	
5	72	2min
6	12	$\infty$



**Gel extraction (operator: Adonie)**

- Gel extraction
- Liquid extraction

Elution buffer:

- EB
- H2O

Elution volume: 30ul

Name: SM012 Fr3

Concentration(A260/280nm): 25.75ng/ul

<u>Gibson assembly</u>	<u>(operator: Darcy)</u>	
Backbone		
Fragment 1	PET28b-1	1
Fragment 2	PET28b-2	2.2
Fragment 3	SM012 Fr3	1
Fragment 4	SM004 Fr4	0.8

GA mix		5
All		10
Process		
1	50	60min
2	12	$\infty$

### **Transformation (operator: Diana)**

- Electroporation  $1\mu\text{L}$
- Chemical transformation  $10\mu\text{L}$

Host: DH5 alpha

Plasmid to construct: SM012

Recovery medium: LB, 300ul

Volume of bacterial liquid to spread:  $200\mu\text{L}$

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at  $42^\circ\text{C}$  for 90s

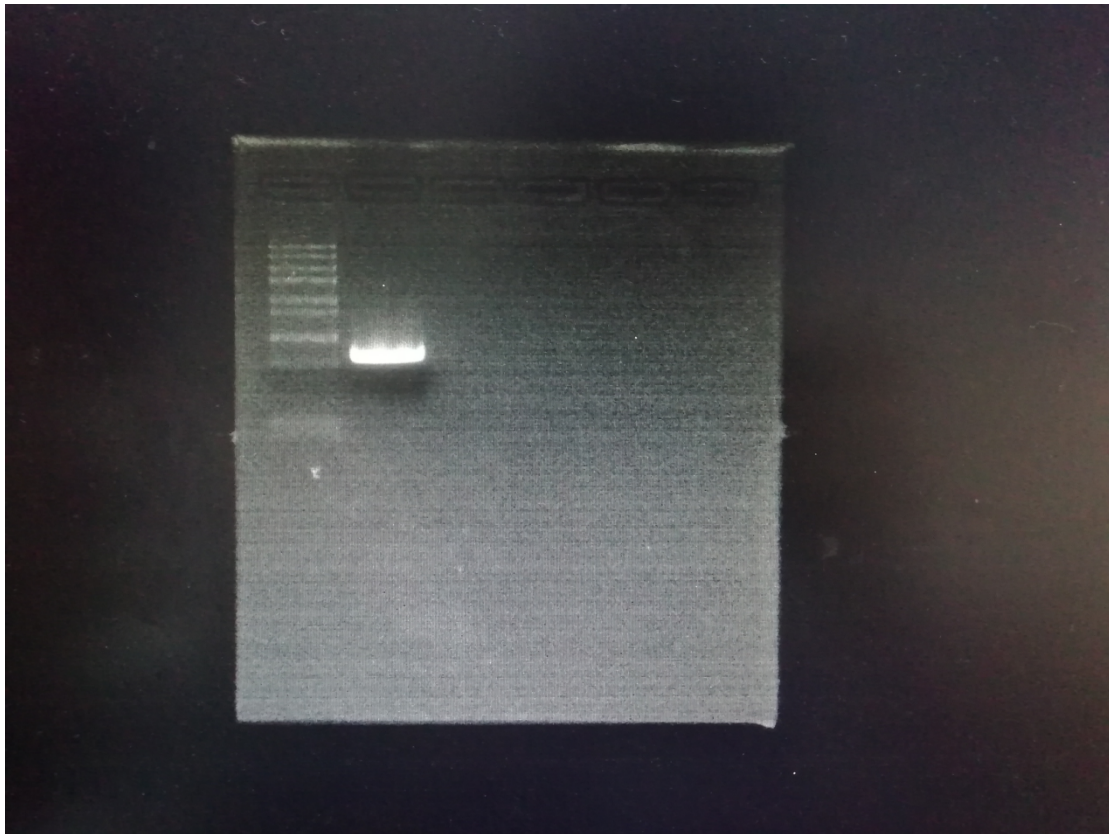
## **Construction of SM012, SM013**

### **PCR Template (operator: Diana)**

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR		
Primer 1	SP023	1
Primer 2	SP001 CsgA-Mfp5-R	1
Template	SM010	0.5

H2O		18.5
DNA polymerase	FASTpfu	9
All		30
		30/tubex1
Product length	370bp	



### **Gel extraction (operator: Adonie)**

Gel extraction

Liquid extraction

Elution buffer

EB

H2O

Elution volume: 30 ul

Name: SM013 Fr (3, 4)

Concentration(A260/280nm): 17.15ng/ul

---

## Culture preservation of Se005 (Done)

### Preparing bacterial glycerol stocks(operator: Diana)

Strain name: DH5 $\alpha$  {Se005}

Number in stocks: S024

---

## Transformation of Se005 into rosetta

Number of single colonies on plates (operator:Diana): ~100

### Inoculation (operator: Diana)

From

Agar plates

Glycerol stock

Liquid starter

Strain	BL(DE3){Se005}
Antibiotics	Cm, Kan
Medium	LB
Volume	4mL in 15mL tube, single colony
Temperature	37
Number	2

7.26 (Fri)

## Construction of SM012, SM013

Number of single colonies on plates (operator:Diana): 0

---

## Construction of SM012, SM013

<u>Gibson assembly</u>	<u>(operator: Diana)</u>	
Backbone		
Fragment 1	pET28b-1	0.7
Fragment 2	pET28b-2	1.8
Fragment 3	SM013 Fr3+4	1.5
Fragment 4	SM004 Fr4	1
H <sub>2</sub> O		Add to 5
GA mix		5
All		10
Process		
1	50	60min
2	12	∞

### Transformation (operator: Diana)

Electroporation 1  $\mu$ L

Chemical transformation 10  $\mu$ L

Host: DH5alpha

Plasmid to construct: SM013

Recovery medium: LB, 300ul

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

---



## Transformation of SM003 into BL21(DE3)

### Inoculation (operator: Diana)

From

Agar plates

Glycerol stock

Liquid starter

Strain	BL21(DE3) {SM003}
Antibiotics	Kan
Medium	LB
Volume	3mL in 15mL tube, single colony
Temperature	37
Number	1

---

## Transformation of Se005 into rosetta (Done)

### Preparing bacterial glycerol stocks(operator: Diana)

Strain name: BL21(DE3) rosetta {Se005}

Number in stocks: S025

**7.27 (Sat)**

## Transformation of SM003 into BL21(DE3) (Done)

### Preparing bacterial glycerol stocks(operator: Emma)

Strain name: BL21(DE3) {SM003}

Number in stocks: S026

## 7.29 (Mon)

### Construction of SM015

#### PCR Template (operator: Cherie, Uranium)

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR		
Primer 1	SP019	1
Primer 2	SP020	1
Template	SY002	1
H2O		30
DNA polymerase	FASTpfu	15
MgSO4		2
All		50
Product length	bp	4893



### Gel extraction (operator: Cherie)

Gel extraction

Liquid extraction

Elution buffer

EB

H2O

Elution volume: 50 ul

Name: SM015

Concentration(A260/280nm): 58.5ng/ul

6.3.Self-ligation of PCR segments (PCR片段自连)

Self-ligation (operator: Uranium)

10X T4 DNA Ligase Buffer NEB1

Fragment SM015 8

T4 DNA Ligase NEB0.5

T4 PNK NEB0.5

All 10

Process

1 25 120min

2 12 ∞

## 7.30(Tues)

Chemical transformation(化转)

### Transformation of SM015 (operator: Uranium)

Electroporation 1  $\mu$ L

Chemical transformation 10 $\mu$ L

Host: DH5a, 100  $\mu$ L

Plasmid to construct: SM015

Recovery medium: LB, 300  $\mu$ L

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Cm

Transformation conditions: heat shock at 42°C for 90s

## 7.31(Wed)

### Transformation of SM015 (operator: Uranium)

1. Inoculation (接种)

#### Inoculation (operator: Uranium)

From

Agar plates

Glycerol stock

Liquid starter

Strain SM015

Antibiotics Cm

Medium LB

Volume purify

Temperature 37

Number 1

4.1. Simple PCR

### Constuction of SM004 SM006 SM011

#### PCR Template (operator: Uranium)

Purified genome

Purified plasmid

Single colony

- Bacteria liquid
- Boiled Bacteria cell lysate

PCR

Primer 1 SP035  
Primer 2 SP034  
Template SM010  
H2O 30+2(MgSO4)  
DNA polymerase FastPfu mix15  
All 50

Product length 5960 bp

Process

1 95 1min  
2 95 20s  
3 55 20s  
4 72 210s  
Go to 2 32 cycle  
5 72 2min  
6 12 ∞

PCR

Primer 1 SP040  
Primer 2 SP034  
Template SM010  
H2O 30+2(MgSO4)  
DNA polymerase FastPfu mix15  
All 50

Product length 6251 bp

Process

1 95 1min  
2 95 20s  
3 55 20s  
4 72 210s  
Go to 2 32 cycle  
5 72 2min  
6 12 ∞

PCR SM007 Fr1

Primer 1 SP029  
Primer 2 SP028  
Template SM008  
H2O 30+2(MgSO4)  
DNA polymerase FastPfu mix15  
All 50

Product length 5870 bp

Process

1 95 1min

2 95 20s  
3 55 20s  
4 72 210s  
Go to 2 32 cycle  
5 72 2min  
6 12 ∞

PCR SM011 Fr1  
Primer 1 SP029  
Primer 2 SP028  
Template SM008  
H2O 30+2(MgSO4)  
DNA polymerase FastPfu mix15  
All 50

Product length 5870 bp

Process  
1 95 1min  
2 95 20s  
3 55 20s  
4 72 210s  
Go to 2 32 cycle  
5 72 2min  
6 12 ∞

PCR  
Primer 1 SP036  
Primer 2 SP037  
Template SM010  
H2O 32  
DNA polymerase FastPfu mix15  
All 50

Product length 256 bp

Process  
1 95 1min  
2 95 20s  
3 55 20s  
4 72 30s  
Go to 2 32 cycle  
5 72 2min  
6 12 ∞

PCR  
Primer 1 SP038  
Primer 2 SP039  
Template SM010  
H2O 32  
DNA polymerase FastPfu mix15

All 50

Product length 250 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 32 cycle

5 72 2min

6 12 ∞

PCR

Primer 1 SP036

Primer 2 SP042

Template SM004

H2O 32

DNA polymerase FastPfu mix15

All 50

Product length 256 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 32 cycle

5 72 2min

6 12 ∞

PCR

Primer 1 SP030

Primer 2 SP031

Template SM008

H2O 32

DNA polymerase FastPfu mix15

All 50

Product length 166 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 32 cycle

5 72 2min

6 12 ∞

PCR

Primer 1 SP030  
Primer 2 SP031  
Template SM008  
H2O 32  
DNA polymerase FastPfu mix15  
All 50

Product length 166 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 32 cycle

5 72 2min

6 12 ∞

PCR

Primer 1 SP032

Primer 2 SP033

Template SM008

H2O 32

DNA polymerase FastPfu mix15

All 50

Product length 160 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 32 cycle

5 72 2min

6 12 ∞

## 8.1(Thur)

Constuction of SM004 006 011

### Gel extraction (operator: Uranium)

Gel extraction

Liquid extraction

Elution buffer

EB

H2O



Elution volume: 30  $\mu$ L

Name: SM005fr1, SM006fr1, SM005fr2, SM005fr3, SM006fr2, SM007fr2,  
SM011fr2, SM011fr3

Concentration(ng/ul): 10, 40, 20, 35, 20, 10, 20, 15

PCR Template (operator: Uranium)

Purified genome

Purified plasmid

Single colony

Bacteria liquid

Boiled Bacteria cell lysate

PCR

Primer 1 SP028

Primer 2 SP029

Template SM008

H2O 30+2MgSO44

DNA polymerase Fastpfu Mix 15

All 50x2

Product length 5870 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 7 cycle

5 95 20s

6 58 20s

7 72 30s

Go to 5 25 cycle

5 72 5min

6 12  $\infty$

Colony selection (operator: )

From

Agar plates

Glycerol stock

Liquid starter

Strain SM015

Antibiotics resistance Cm

Medium LB

Volume Inoculating single colonies on plates

Temperature 37

Number 8

## 8.1

### Gel extraction (operator: Uranium)

Gel extraction

Liquid extraction

Elution buffer

EB

H2O

Elution volume: 30  $\mu$ L

Name: SM005fr1, SM006fr1, SM005fr2, SM005fr3, SM006fr2, SM007fr2, SM011fr2, SM011fr3

Concentration(ng/ul): 10, 40, 20, 35, 20, 10, 20, 15

### PCR Template (operator: Uranium)

Purified genome

Purified plasmid

Single colony

Bacteria liquid

Boiled Bacteria cell lysate

### PCR

Primer 1 SP028

Primer 2 SP029

Template SM008

H2O 30+2MgSO44

DNA polymerase Fastpfu Mix 15

All 50x2

Product length 5870 bp

Process

1 95 1min

2 95 20s

3 55 20s

4 72 30s

Go to 2 7 cycle

5 95 20s

6 58 20s

7 72 30s

Go to 5 25 cycle

5 72 5min

6 12 ∞

### Colony selection (operator: )

From

Agar plates

Glycerol stock

Liquid starter

Strain SM015

Antibiotics resistance Cm

Medium LB

Volume Inoculating single colonies on plates

Temperature 37

Number 8

## 8.2

# Golden Gate for SM005, SM006

Backbone p Pet28b

Fragment 1 MFP5(both)

Fragment 2 MFP5(SM005)

H2O 1

T4 Ligase Buffer NEB 2

Restriction enzyme BbsI (NEB) 1

T4 Ligase NEB 1

All 20

Process

1 37 5min

2 16 10min

Go to 2 10 cycle

3 37 10min

4 50 5min

5 80 5min

6 12 ∞

**Transformation (operator: Uranium)**

Electroporation 1  $\mu\text{L}$

Chemical transformation 10  $\mu\text{L}$

Host: DH5a, 50  $\mu\text{L}$  each

Plasmid to construct: sm005, sm006

Recovery medium: LB, 300  $\mu\text{L}$

Volume of bacterial liquid to spread: 150  $\mu\text{L}$

Agar plate and antibiotic to use: LB, Kan

Transformation conditions: heat shock at 42°C for 90s

**PCR Template (operator: Cherie)**

Purified genome

Purified plasmid

Single colony

Bacteria liquid

Boiled Bacteria cell lysate

PCR		
Primer 1	SP028	0.6

Primer 2	SP029	0.6
Template	SM008	0.6
H2O		18
DNA polymerase	fastpfu mix	9
All		30
MgSO4		1.2
Product length	5870bp	

PCR		
Primer 1	SP034	0.6
Primer 2	SP035	0.6
Template	SM010	0.6
H2O		18
DNA polymerase	fastpfu mix	9
All		30
MgSO4		1.2
Product length	5960bp	

## 8.3

### PCR Template (operator: Uranium)

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR

Primer 1 SP004 F 1.5

Primer 2 SP004 R 1.5

Template single colony of SM006

H2O 12

DNA polymerase Q5 Mix 13

25/tubex2

Product length 1681bp

Process

1 94 10min

2 94 30s

3 55 30s

4 72 30s

Go to 2 35cycle

5 72 2min

6 12 ∞

## 8.6

### Preparing bacterial glycerol stocks (operator: Adonie)

Strain name: S027

Number in stocks: 2

## 8.7

### PCR Template (operator: Adonie)

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR	Name	Volume
Primer 1	SP004	1.5
Primer 2	SP004	1.5
Template	SM005 & SM011 colony	/
H2O		12
DNA polymerase	2x Starmix	15
All		30
Product length	1617bp	

Process	Temperature	Time
1	95	1min
2	95	30s
3	55	30s
4	72	1min 30s
Go to	32 cycle	
5	72	2 min
6	12	∞

Gel Picture

## 8.8

Preparing bacterial glycerol stocks (operator: Ginny Amy)

Strain name: S028,S029,S030,S031,S032,S033

Number in stocks: 2 for each\

**PCR Template (operator: Ginny Amy)**

- Purified genome
- Purified plasmid
- √ Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

**PCR x8u**

Primer 1 sp004 F 1.5

Primer 2 sp004 R 1.5

Template SM014

H2O 12

DNA polymerase R-taq 15

All 30

Product length 1400 bp

## 8.9

**Sequencing Results Analysis of SM005, SM006 and SM014**

Primers: SP004 F/R

SM005:

- Wrong insertion of mfp3 appears in sequencing results
- 8 mismatches

Conclusion: PCR template may be faulty

SM006:

- 4 gaps/insertions



- pET28b isn't loaded (空载)

Conclusion: ???

SM014

- 1 gap in cp19k that could lead to frame shifting (移码)
- Otherwise ok

Conclusion: Specimen number 19 should be sent for sequencing again

## 8.12

### Inoculation (operator: Uranium )

From

Agar plates

√Glycerol stock

Liquid starter

Strain S006, S010

Antibiotics Kan

Medium LB

Volume 3mL in 15mL tube, single colony

Temperature 37

Number 1

## 8.13

### Preparing bacterial glycerol stocks (operator: Adonie)

Strain name: S034

Number in stocks: 2

### Plasmid extraction (operator: Adonie)

strain	pET28b-rBalcp19k-linker-mfp5-7*His (SM014)
--------	--

plasmid	SM014
---------	-------

strain	pET28b-CsgA-linker-mfp3-7*his (SM008)
plasmid	SM008

strain	pET28b-csgA-linker-mfp5-7*His (SM010)
plasmid	SM010

Elution buffer

TB

H2O

Elution volume: 50 $\mu$ l

Concentration(A260/280):

SM014(1): 80.7 ng/ul

SM014(2): 96.15 ng/ul

SM010: 132.2 ng/ul

SM008: 122 ng/ul

## 8.14

### Sequencing results analysis (operator: XYZ)

Sample	sequencing result	next plan
SM006-4/8	empty vector	stop
SM011-8(csgA-mfp3-mfp3-mfp3)	the sequence is the same as SM010 (csgA-mfp5) resulting from contaminated PCR templates	stop

SM014-4/6/8/11/13/15	all correct, therefore the SM014-19/20 have been sent to sequence several days ago probably be right.	preparing glycerol stocks
----------------------	---	---------------------------

**PCR Template (operator: Adonie)**

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

SM005

Fr1

PCR	Name	Volume
Primer 1	SP035	1.5
Primer 2	SP034	1.5
Template	SM010	1
H2O		16
DNA polymerase	FASTPFU	9
MgSO4		1
All		30
Product length	5460bp	

Fr2

PCR	Name	Volume
Primer 1	SP036	1.5
Primer 2	SP037	1.5

Template	SM010	1
H2O		17
DNA polymerase	FASTPFU	9
All		30
Product length	256bp	

### Fr3

PCR	Name	Volume
Primer 1	SP038	1.5
Primer 2	SP039	1.5
Template	SM010	1
H2O		17
DNA polymerase	FASTPFU	9
All		30
Product length	250bp	

### SM016

#### Fr1

PCR	Name	Volume
Primer 1	SP034	1.5
Primer 2	SP035	1.5
Template	SM010	1
H2O		16

DNA polymerase	FASTPFU	9
MgSO4		1
All		30
Product length	5960	

## Fr2

PCR	Name	Volume
Primer 1	SP034	1.5
Primer 2	SP035	1.5
Template	SM010	1
H2O		16
DNA polymerase	FASTPFU	9
All		30
Product length	256bp	

## SM017

PCR	Name	Volume
Primer 1	SP049	1.5
Primer 2	SP003_R	1.5
Template	SM010	1
H2O		16
DNA polymerase	FASTPFU	9
MgSO4		1
All		30

Product length	5521bp	
----------------	--------	--

## SM018

PCR	Name	Volume
Primer 1	SP050	1.5
Primer 2	SP003_R	1.5
Template	SM010	1
H2O		16
DNA polymerase	FASTPFU	9
MgSO4	u	1
All		30
Product length	5431bp	

## Process

Process	Temperature	Time
1	98	1min
2	98	30sec
3	62	30sec, -0.5°C/cyc
4	72	3min
Go to 2	12 cycle	
5	98	30sec
6	55	30sec
7	72	3min
Go to 6	22 cycle	
7	72	7min

8	12	forever
---	----	---------

Process		
1	98	10min
2	98	30s
3	55	30s
4	72	30s
Go to 2	32 cycle	
5	72	2min
6	12	$\infty$

**Gel Picture:**

SM005 Fr1 will not undergo gel extraction because it's band did not display the right results. Replace this sample with SM016 Fr1.

**Gel Extraction (operator: Diana, Adonie)**

√Gel extraction

Liquid extraction

**Elution buffer**

EB

H2O

Elution volume: 30ul

Name: SM005 Fr2

Concentration(A260/280nm): 7.9ng/ul

Name: SM005 Fr3

Concentration(A260/280nm): 8ng/ul

Name: SM016 Fr1

Concentration(A260/280nm): 15.1ng/ul

Name: SM016 Fr2

Concentration(A260/280nm): 8.45ng/ul

Name: SM017 Fr1

Concentration(A260/280nm): 27.15ng/ul

Name: SM018 Fr1

Concentration(A260/280nm): 8.25ng/ul

The concentrations of gel extraction are too low for GoldenGate Assembly.  
Another round of PCR and gel extraction will be undertaken.

Transformation (operator: Diana)

Electroporation 1 $\mu$ L

Chemical transformation 0.5 $\mu$ L

1. Host: BL21(DE3)

Plasmid to construct: SM014, SM015

2. Host: BL21(DE3) Rosetta

Plasmid to construct: SM014

Recovery medium: LB

Volume of bacterial liquid to spread: 200  $\mu$ L

Agar plate and antibiotic to use: LB, Cm + Kan

Transformation conditions: heat shock at 42°C for 90s

## 8.15



## PCR Template (operator: Adonie)

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

SM005

Fr1

PCR	Name	Volume
Primer 1	SP035	1.5
Primer 2	SP034	1.5
Template	SM010	1
H2O		16
DNA polymerase	FASTPFU	9
MgSO4		1
All		30
Product length	5460bp	

Fr2

PCR	Name	Volume
Primer 1	SP036	1.5
Primer 2	SP037	1.5
Template	SM010	1
H2O		17
DNA polymerase	FASTPFU	9

All		30
Product length	256bp	

### Fr3

PCR	Name	Volume
Primer 1	SP038	1.5
Primer 2	SP039	1.5
Template	SM010	1
H2O		17
DNA polymerase	FASTPFU	9
All		30
Product length	250bp	

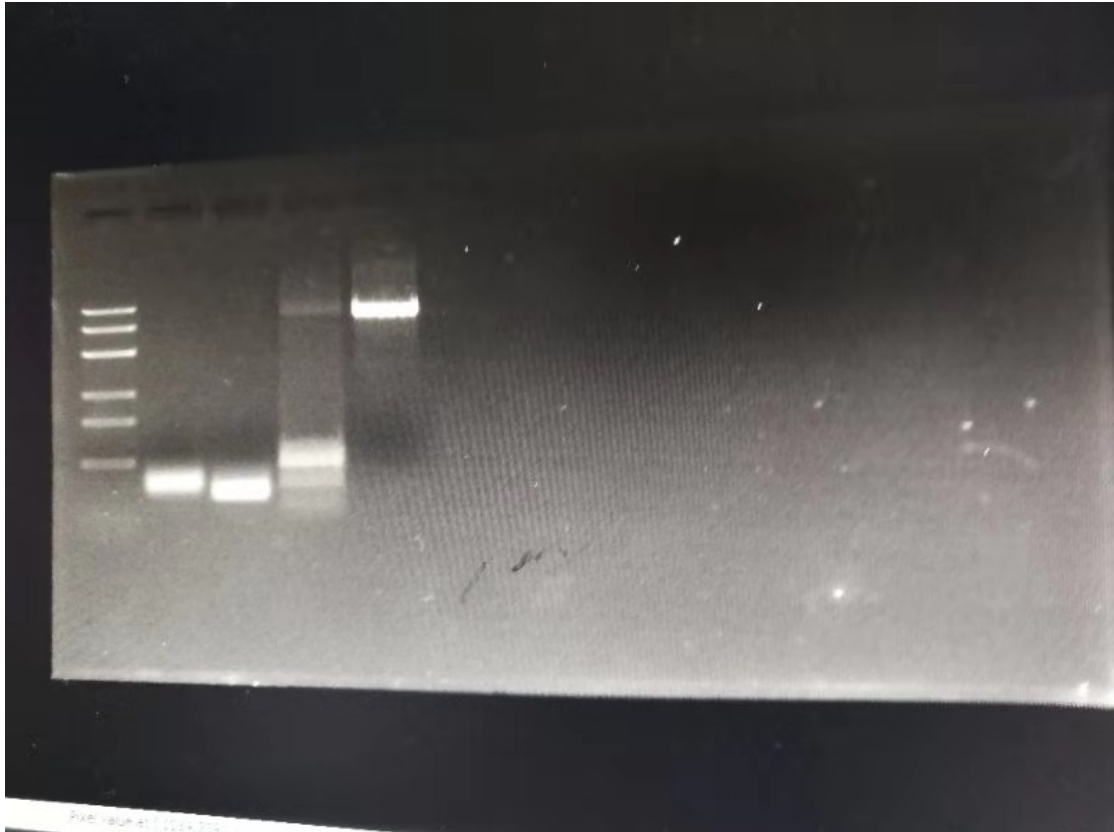
### SM018

PCR	Name	Volume
Primer 1	SP050	1.5
Primer 2	SP003_R	1.5
Template	SM010	1
H2O		16
DNA polymerase	FASTPFU	9
MgSO4		1
All		30
Product length	5431bp	

Process	Temperature	Time
1	98	1min
2	98	30sec
3	62	30sec, -0.5°C/cyc
4	72	3min
Go to 2	12 cycle	
5	98	30sec
6	55	30sec
7	72	3min
Go to 6	22 cycle	
7	72	7min
8	12	forever

Process		
1	98	10min
2	98	30s
3	55	30s
4	72	30s
Go to 2	32 cycle	
5	72	2min
6	12	∞

Gel Picture:



## Gel extraction (operator: Uranium)

Gel extraction

Liquid extraction

Elution buffer

EB

H<sub>2</sub>O

Elution volume: 30  $\mu$ L

Name: SM005Fr2,Fr3,Fr1,SM018 Fr1

Concentration(A260/280nm): 14.2 ng/ $\mu$ L

# 8.17

## PCR of all constructed vector (Operator: Adonie)

### PCR of Sy003 and SM015 again

- Purified genome
- Purified plasmid
- Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

#### Sy003

PCR	Name	Volume
Primer 1	SP004-F	1.5
Primer 2	SP004-R	1.5
Template	Sy003	1
H2O		10
DNA polymerase	Starmix 2x	10
All		20
Product length	1140bp	

#### SM015

PCR	Name	Volume
Primer 1	SP024	1.5
Primer 2	SP025	1.5
Template	Sy003	1
H2O		10

DNA polymerase	Starmix 2x	10
All		20
Product length	425bp	

Process	Temperature	Time
1	95	1min
2	95	30s
3	58	45s
4	72	45s
Go to	32 cycle	
5	72	5 min
6	12	∞

---

## Construction of SM005/016/017/018

### Clony PCR

- Purified genome
- Purified plasmid
- √ Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR	Name	Volume
Primer 1	SP004-F	1.5
Primer 2	SP004-R	1.5
Template	SM005/016/017/018 colonies	/
H2O		12
DNA polymerase	Starmix 2x	15
All		30
Product length	SM005 - 1617 bp SM016 - 1329 bp SM017 - 738 bp SM018 - 648 bp	

Number of colonies:

SM005 - 6

SM016 - 3

SM017 - 7

SM018 - 8

Process	Temperature	Time
1	95	1min
2	95	30s
3	55	30s
4	72	1min 30s
Go to	32 cycle	
5	72	2 min
6	12	∞

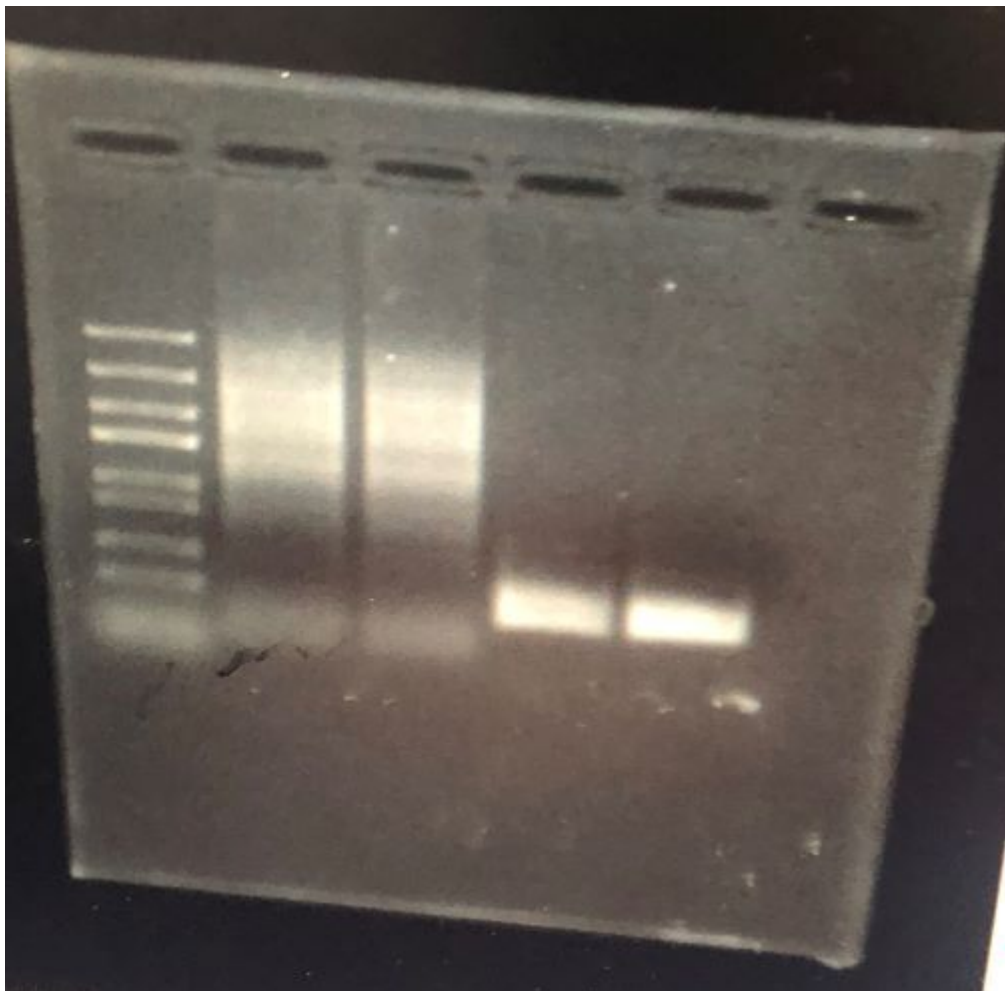
Gel picture

---

## Construction of SM019

### Dnpi digestion and gel electrophoresis

Gel Picture:



M SM019 Fr2 SM019 Fr2 SM019 Fr1 SM019 Fr1

Analysis:

SM019 Fr2: smear bands. PCR again.



SM019 Fr2: Correct size. Cut and store in -20 in E&A box

## 8.18

### Construction of SM005/016/017/018

#### Gel electrophoresis

Results analysis

Bands of SM016/017/018 are right while SM005 are smaller than predicted.

Lane	samples	predicted size	band size	send for sequencing
1	Trans 5K			
2-7	SM005	1617	1200	1
8-10	SM016	1329	1300	1, 2
11-17	SM017	738	700	3, 5
18-25	SM018	648	600	7, 8

#### To be Sent for sequencing

in 4 refrigerator

#### Colony inoculation on agar plates

Colonies sent for sequencing were inoculated on Kan plates.

---

# PCR of all constructed vector

## Inoculation of DH5α/Sy003

From

- Agar plates
- √Glycerol stock
- Liquid starter

Strain	S021 DH5α/Sy003
Antibiotics	Kan
Medium	LB
Temperature	37
Number	1

## 8.19

# PCR of all constructed vector

## Colony PCR of sy003(S021)

### PCR Template

- Purified genome
- Purified plasmid
- √Single colony
- Bacteria liquid
- Boiled Bacteria cell lysate

PCR		
Primer 1	SP004-F	1
Primer 2	SP004-R	1
Template	S021 colonies	/
H2O		8
DNA polymerase	2XStarmix	10
All		20
		20/tubex3
Product length	1140 bp	
Process		
1	95	10min
2	95	30s
3	55	30s
4	72	45s
Go to 2	32 cycle	
5	72	5min
6	12	$\infty$

pET28b empty vector was set as control.

## Gel electrophoresis

Analysis: Nonspecific bands(~600bp) still present.

## 8.21

# Construction of SM005/016/017/018

## Sequencing

Sample	Primer	Requirements
<input type="checkbox"/> Bacteria liquid <input type="checkbox"/> plasmid <input type="checkbox"/> purified PCR <input checked="" type="checkbox"/> unpurified PCR SM005 No.1 SM017 No.1&2 SM018 No.3&5 SM016 No.7&8	SP004-F SP004-R	<input checked="" type="checkbox"/> Forward <input type="checkbox"/> Reverse <input checked="" type="checkbox"/> Both

Results analysis:

SM005: empty vectors. Next plan: PCR by using gel extracted samples as templates.

SM017: No.5 is correct while No.3 has 3 insertions. Next plan: preparing glycerol stocks of No.5.

SM018: Both No.7 and No.8 are correct. Next plan: preparing glycerol stocks of No.7 and No.8.

SM016: empty vectors. Next plan: PCR by using gel extracted samples as templates.

## 8.24

### PCR Template (operator: Adonie)

- Purified genome
- Purified plasmid
- Single colony

- Bacteria liquid
- Boiled Bacteria cell lysate

SM005/SM016

Fr1

PCR	Name	Volume
Primer 1	SP035	1.5
Primer 2	SP034	1.5
Template	SM016 Fr1 (Gel extraction)	1
H2O		30
DNA polymerase	FASTPFU	15
MgSO4		1
All		50
Product length	5460bp	

SM005 Fr2

PCR	Name	Volume
Primer 1	SP036	1.5
Primer 2	SP037	1.5
Template	SM005 Fr2 (Gel extraction)	1
H2O		31
DNA polymerase	FASTPFU	15
All		50
Product length	256bp	

### SM005 Fr3

PCR	Name	Volume
Primer 1	SP038	1.5
Primer 2	SP039	1.5
Template	SM005 Fr3 (Gel extraction)	1
H2O		30
DNA polymerase	FASTPFU	15
All		50
Product length	250bp	

### SM016 Fr2

PCR	Name	Volume
Primer 1	SP044	1.5
Primer 2	SP043	1.5
Template	SM016 Fr2 (Gel extraction)	1
H2O		30
DNA polymerase	FASTPFU	15
All		50
Product length	198bp	

### SM019 Fr1

PCR	Name	Volume
Primer 1	SP047	1.5

Primer 2	SP048	1.5
Template	SM008	1
H2O		30
DNA polymerase	FASTPFU	15
MgSO4		1
All		50
Product length	5483bp	

### Process:

Process	Temperature	Time
1	98	1min
2	98	30sec
3	62	30sec, -0.5°C/cyc
4	72	3min
Go to 2	12 cycle	
5	98	30sec
6	55	30sec
7	72	3min
Go to 6	22 cycle	
7	72	7min
8	12	forever

Process		
1	98	10min

2	98	30s
3	55	30s
4	72	30s
Go to 2	32 cycle	
5	72	2min
6	12	$\infty$

Gel extraction (operator: Uranium)

Gel extraction

Liquid extraction

Elution buffer

EB

H2O

Elution volume: 30  $\mu$ L

Name: SM005 fr2, SM005fr3, SM016fr2, SM018fr1

Concentration(A260/280nm): 20 ng/ $\mu$ L, 9ng/ul, 8ng/ul, 12.5ng/ul