Week1 (3.1-3.7): pYGRa1-mCherry plasmid construction and function validation

(A) pYGRa1-mCherry plasmid construction:

1. Amplification of vectors by PCR

Target vector fragments were obtained by PCR with the following RCR system:

PCR system	(50 µ L)	I	PCI	{	
$2 \times Mix$	25u1		98° C	5min	
mCherry-F	2u1		98° C	30s T	
mCherry-R	2u1		59° C	30s -	×25
Template	50ng		72° C	90s	
DDW	20u1		72° C	5min	
			25° C	∞	

The polymerase chain reaction product was detected by agarose gel electrophoresis and the correct length of the target fragment was obtained as 2900 bp. We got the correct target fragment.

2.(1) Amplification of Orf4 fragment by PCR

The target fragments were obtained by PCR. The RCR system was as follows:

PCR system	(50 μ L)	 PC	R	
$2 \times Mix$	25u1	 98° C	5min	
ORF4-F	2u1	98° C	30s	
ORF4-R	2u1	59° C	30s -	×25
template	50ng	72° C	35s	
DDW	20u1	72° C	5min	
		 25° C	∞	

The polymerase chain reaction product was detected by agarose gel electrophoresis and the correct length of the target fragment was obtained as 1118 bp. we got the correct target fragment.

2.(2) Amplification of PA1 fragment by PCR

The target fragments were obtained by PCR. The RCR system was as follows:

PCF	{	PCR system	(50 µ L)
98° C	5min	$2 \times Mix$	25u1
98° C	30s	PA1-F	2u1
59° C	$30s \rightarrow \times 25$	PA1-R	2u1
72° C	15s	template	50ng
72° C	5min	DDW	20u1
25° C	∞		

The polymerase chain reaction product was detected by agarose gel electrophoresis and the correct length of the target fragment was obtained as 407 bp. we got the correct target fragment and the sample was cut and recovered.

2.(3) Amplification of PA2 fragment by PCR

The target fragments were obtained by PCR, and the RCR system was as follows:

		_			
PCR system	(50 μ L)	_	PCI	2	
$2 \times Mix$	25u1	_	98° C	5min	
PA2-F	2u1		98° C	30s	
PA2-R	2u1		58° C	30s -	×25
template	50ng		72° C	10s	
DDW	20u1		72° C	5min	
		-	25° C	∞	

The polymerase chain reaction product was detected by agarose gel electrophoresis and the correct length of the target fragment was obtained as 340 bp. we got the correct target fragment.

3.Cutting glue recycling

4. Connecting the four fragments using the Gibsion assembly method

Gibson (C116) ligation was used to obtain pYGRa1-mCherry plasmid, mCherry 2900bp, Orf4 1118bp, PA1 407bp, PA2 340bp

Gibson s	system
0.02 x 2900bp	ng
0.04 x 1118bp	ng
0.04 x 407bp	ng
0.04 x 340bp	ng
$2 \times c115$ Mix	5u1
DDW	To 10u1

Gibso	n.
50° C	30min
4° C	∞

5. Chemical transformation

Using the chemical method, the ligation product was transformed into DH5 α receptor cells and then spread on LB plates containing Amp antibiotics and incubated overnight at 37°C.

6. Monoclonal inoculation

After overnight incubation at 37°C, only 1 colony grew on the plate. The single clone was inoculated into liquid medium supplemented with 1% ampicillin and incubated at 37°C for 12 hours.

7. Extract plasmid, send for sequencing

The plasmid was extracted and sequenced. pYGRa1-mCherry sequencing results were correct.

(B) Validation of pYGRa1-mCherry function

1. Conversion

The correctly sequenced plasmid was transfected into BW25113 sensory cells and then spread on LB plates containing Amp resistance and incubated at 37°C overnight. Expected resistant single colonies grew on the plates, and single colonies were subsequently selected for plasmid extraction and functional verification.

2. Pick monoclonal inoculation

The monoclone was inoculated into liquid medium supplemented with 1% ampicillin and incubated at 37°C for 12 hours.

3. Induction

Three sets of parallel experiments were done for each system, induced in a shaker at 30°C for 18h,200ul of each tube was taken into a 96-well plate, and the fluorescence and OD values were measured in an enzyme lab, the two systems were as follows:

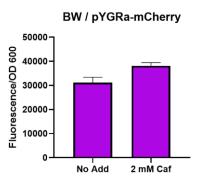
Induction of	system1
LB	5m1
Amp	2u1
Inoculum amount	5u1
IPTG	5u1

Induction of system2

-	LB	5m1
	Amp	2u1
	Inoculum amount	5u1
	IPTG	5u1
	Caffeine	2mM

4. Measure fluorescence and OD values

The results were as follows: strong fluorescence was found without caffeine, and the Orf4 prepromoter should be too weak.



Week2 (3.8-3.14): construction and functional validation of

pYGRa2-mCherry plasmid

(A) Construction of pYGRa2-mCherry plasmid:

The ORF4 front region was removed and the Pfrm promoter was added before Orf4 to substantially repress mCherry expression.

1. Amplification of vectors by PCR

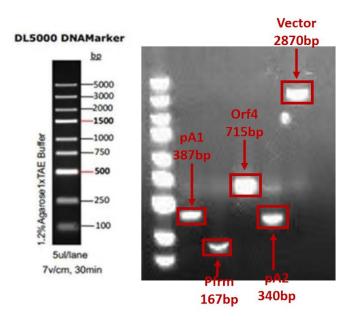
The target vector fragments were obtained by PCR, the corresponding primers and annealing temperature were changed, and the remaining operations were the same as those in the first week of the experiment.

2. Amplification of Orf4, pfrm, PA1 and PA2 fragments by PCR

The fragments were obtained by PCR, replacing the corresponding primers and annealing temperatures, and the rest of the procedure was the same as in the first week of the experiment.

3. Cutting glue recycling

The correct bands obtained were cut and recovered, and the PCR products were detected by agarose gel electrophoresis as follows:



4. Linking five fragments using the Gibsion assembly method

Gibson sys	stem	•	Gibso	n.
0.02 x 2870bp	ng	-	50° C	30min
0.04 x 715bp	ng		4° C	∞
0.04 x 167bp	ng	•		
0.04 x 387bp	ng			
0.04 x 340bp	ng			
$2 \times c116$ Mix	5u1			
DDW	to10u1	_		

The pYGRa2-mCherry plasmid was obtained using Gibson (C116) ligation mCherry 2870bp, Orf4 715bp, pfrm167bp,PA1 387bp,PA2 340bp

5. Chemical transformation

Same chemical transformation operations as in Week 1

6. Colony PCR

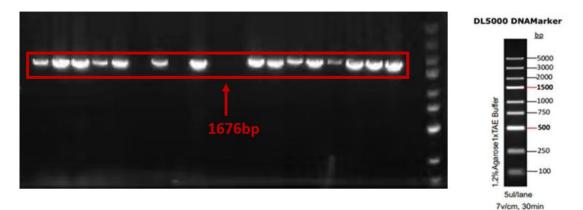
After 12 hours of incubation at 37° C, 20 colonies were selected on the plate and the colony PCR system was as follows:

PCR system (10µL)

2×Hieff 25ul	$2 \times \text{Hieff}$
CE-F 2u1	CE-F
CE-R 2u1	CE-R
DDW 4.2u1	DDW
CE-F 2u1 CE-R 2u1	CE-F CE-R

		PCR		
98°	С		5min	
98°	С		30s	
57°	С		30s	 ×25
72°	С		50s	
72°	С		5min	
25°	С		∞	

The PCR products were examined by agarose gel electrophoresis with the following results:

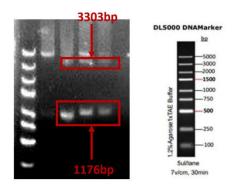


7. Inoculation with monoclonal

Each of the four monoclones was selected and inoculated into liquid medium supplemented with 1% ampicillin and incubated at 37°C for 12 hours.

8. Plasmid extraction + digestion verification

Single enzyme cleavage was verified using BgIII, and the cleavage products were detected by agarose gel electrophoresis. The target bands were 3303bp, 1176bp. the results are plotted below:



9. pYGRa2-mCherry sequencing result is correct.

(B) Functional validation of pYGRa2-mCherry

1. Transformation

The correctly sequenced pYGRa2-mCherry plasmid was transfected into BW25113 receptor cells and then spread on LB plates containing Amp antibiotics and incubated inverted at 37°C overnight. Expected resistant single colonies grew on the plates, and single colonies were subsequently selected for plasmid extraction and functional verification.

2. Pick monoclonal inoculation

Selected monoclones were inoculated into liquid medium supplemented with 1% ampicillin and incubated at 37°C for 12 hours.

3. Induction

Comparative induction was done for GRA1 and GRA2, 3 sets of parallel experiments were done for each system, and the induction was carried out in a shaker at 30°C for 18h, 200ul was taken from each tube into a 96-well plate, and fluorescence and OD values were

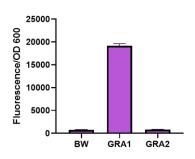
measured, the systems were as follows:

Induction of	system1
LB	5m1
Amp	2u1
Inoculum amount	5u1
IPTG	5u1
Caffeine	2mM
Induction of	system2

LB	5m1
Amp	2u1
Inoculum amount	5u1
IPTG	5u1
Caffeine	2mM

4. Measure fluorescence and OD values

RESULTS: The addition of Pfrm promoter before Orf4 significantly inhibited the expression of mCherry, but the inhibition could not be lifted even after the addition of caffeine. This may be due to the fact that Pfrm is too strong.



Week3 (3.15-3.21): construction and functional validation of

pYGRa3-mCherry, PYGRa4-mCherry plasmids

Two approaches to weaken Orf4 expression were tried separately:

(A) pYGRa3-mCherry plasmid construction

We attempted to weaken the expression of Orf4, mimic the endogenous formaldehyde transcriptional regulator in E. coli, and use the ndmA promoter to initiate Orf4 and fluorescent proteins to achieve "self-regulation".

1. Amplification of vectors by PCR

The ORF4 front region was removed and the Pfrm promoter was added before Orf4 to substantially repress mCherry expression.

The target vector fragments were obtained by PCR, the corresponding primers and annealing temperature were changed, and the remaining operations were the same as those in the first week of the experiment.

2. Amplification of Orf4, pndmA fragments by PCR

The fragments were obtained by PCR, replacing the corresponding primers and annealing temperatures, and the rest of the procedure was the same as in the first week of the experiment.

3. Cutting glue recycling

Recover the correct strip by cutting the glue.

4. Connecting three fragments using Gibsion assembly method

The pYGRa3-mCherry plasmid was obtained using Gibson (C116) ligation, vector 2979bp, Orf4 663bp, pndmA 727bp

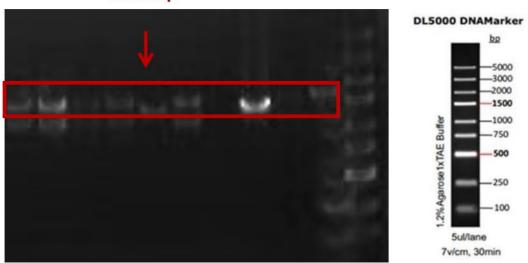
	•	-		
Gibson sy	stem		Gibso	n.
0.02 x 2979bp	ng		50° C	30min
0.04 x 663bp	ng		4° C	∞
0.04 x 727bp	ng	-		
$2 \times c116$ Mix	5u1			
DDW	to10u1	_		

5. Chemical transformation

Same chemical transformation operations as in Week 1

6. Colony PCR

Positive clones were selected and inoculated into liquid medium, and after 12 hours of incubation at 37°C, 10 colonies were selected on the plate and colony PCR was performed, and the PCR products were detected by agarose gel electrophoresis, and the results were as follows:

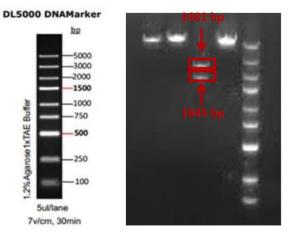


1523bp

Positive monoclones were selected and inoculated into liquid medium supplemented with 1% ampicillin and incubated at 37°C for 12 hours.

7. pYGRa3-mCherry plasmid was extracted and verified by digestion.

Single enzyme cleavage was verified using BgIII, and the cleavage products were detected by agarose gel electrophoresis. The target bands were 1845bp and 2481bp. The results are shown below:



8. pYGRa3-mCherry sequencing result is correct

(B) Construction of PYGRa4-mCherry plasmid

Orf4 was initiated using a medium-strength Anderson promoter, p101

1. Amplification of vectors by PCR

The target vector fragments were obtained by PCR, the corresponding primers and annealing temperature were changed, and the remaining operations were the same as those in the first week of the experiment.

2. Amplification of Orf4, pndmA, P101 fragments by PCR

The fragments were obtained by PCR, replacing the corresponding primers and annealing temperatures, and the rest of the procedure was the same as in the first week of the experiment.

3. Cutting glue recycling

Recover the correct strip by cutting the glue.

4. Connecting the four fragments using the Gibsion assembly method

Gibson s	ystem
0.02 x 2970bp	ng
0.04 \times 695bp	ng
0.04 $ imes$ 707bp	ng
0.04 x 124	ng
$2 \times c116$ Mix	5u1
DDW	to10u1

Orf4 695bp, pndmA 707bp, P101 124	
Gibson system	Gibson.

50° C

4° C

30min

 ∞

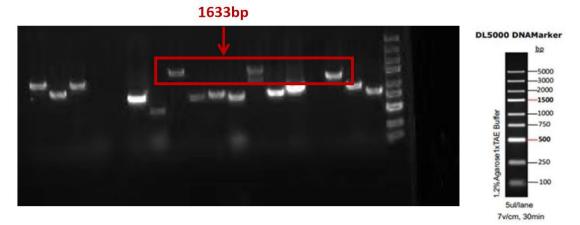
5. Chemical transformation

Same chemical transformation operations as in Week 1

6. Colony PCR

Positive clones were selected and inoculated into liquid medium, and after 12 hours of incubation at 37°C, 10 colonies were selected on the plate and colony PCR was performed, and the PCR products were detected by agarose gel electrophoresis, and the results were as follows:

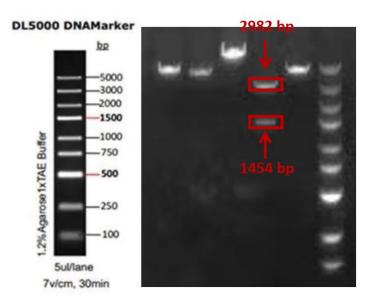
The pYGRa4-mCherry plasmid was obtained using Gibson (C116) ligation. mCherry 2970bp,



Three positive monoclones were selected and inoculated into liquid medium supplemented with 1% ampicillin and incubated at 37°C for 12 hours.

7. pYGRa4-mCherry plasmid was extracted for digestion and validation.

Single enzyme cleavage was verified using EcoRI, and the cleavage product was detected by agarose gel electrophoresis. The target band was 454bp+2982bp and the resultant graph is shown below:



8. pYGRa4-mCherry sequencing result is correct

(C) Functional validation of pYGRa3-mCherry, PYGRa4-mCherry

1. Transformation

The successfully sequenced pYGRa4-mCherry and pYGRa3-mCherry were transformed into BW25113 receptor cells, respectively, for functional validation together with the two previously constructed plasmids.

2. Induction

Comparative induction was done for GRA1, GRA2, GRA3 and GRA4, and 3 sets of parallel experiments were done for each system, which were induced in a shaker at 30°C for 18h, 200ul of each tube was taken into 96-well plates, and fluorescence and OD values were measured in an enzyme labeller, and the systems were as follows:

Induction of	system1
LB	5m1
Amp	2u1
Inoculum amount	5u1
IPTG	5u1

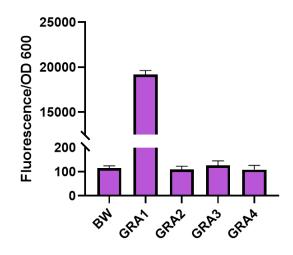
Induction of	system2
LB	5m1
Amp	2u1
Inoculum amount	5u1

IPTG	5u1
Caffeine	2mM

3. Measure fluorescence and OD values

The results were as follows: several methods used so far were ineffective, GRA1 could not repress the NdmA promoter due to its own promoter being too weak, resulting in low ORF4 expression; GRA2, GRA3, and GEA4 could not open the NdmA promoter after being repressed due to high ORF4 expression.

Afterwards it is planned to test for a weaker promoter P107.





pYGRa5-mCherry plasmid

(A) pYGRa5-mCherry plasmid construction

Orf4 was initiated using a moderately weak Anderson promoter p107

1.PCR amplification

Amplification of pYGRA4 fragments by PCR The target vector fragments were obtained by PCR with the following RCR system.

PC	R
98° C	5min
98° C	30s 7
59° C	30s – ×25
72° C	2min_
72° C	5min
25° C	∞

PCR system	(50 μ L)
$2 \times Mix$	25u1
GRA5-F	2u1
GRA5-R	2u1
pYGRa5-mCherry	50ng
DDW	20u1

The expected product size of 4456 bp was obtained, pending subsequent analysis.

2. Product purification

The PCR products were purified and analysed by electrophoresis through 1% agarose gel. The electrophoresis ran clear bands and the corresponding positions were consistent with the expected size of 4456 bp. The purified PCR product was successfully obtained. The purity and concentration of the product were as expected and the band size was correct.

3. DpnI digestion

PCR products were digested using DpnI to remove the unmutated template plasmid.

Dpn I syste	m (10µL)	_	Dpn	Ι
DNA	200ng		37° C	2h
Dpn I	1ul		55° C	15min
Cutsmart	1u1		80° C	15min
DDw	to 10u1	_	4° C	∞

The digestion reaction went smoothly and the enzyme was inactivated after treatment at 85°C. The PCR product was efficiently digested and prepared for transformation of the receptor cells in the next step.

4. Chemical transformation

The digested PCR product was transformed into DH5α receptor cells. After transformation, they were spread on LB plates containing Amp resistance and incubated at 37°C overnight. Expected resistant single colonies grew on the plates, and single colonies were subsequently selected for plasmid extraction and validation.

5. Induction

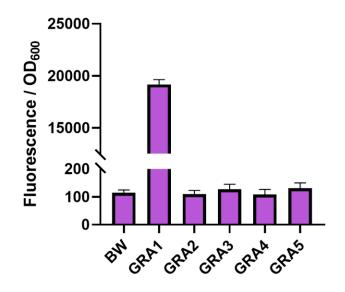
Comparative induction was done for GRA1, GRA2, GRA3, GRA4 and GRA5, and 3 sets of parallel experiments were done for each system, which were induced in a shaker at 30°C for 18h, 200ul of each tube was taken into 96-well plates, and fluorescence and OD values were measured, and the two systems were as follows:

Induction of	system1
LB	5m1
Amp	2u1
Inoculum amount	5u1
IPTG	5u1

Induction of	system2
LB	5m1
Amp	2u1
Inoculum amount	5u1
IPTG	5u1
Caffeine	2mM

6. Measure fluorescence and OD value

The results are as follows:



Several methods used so far have been ineffective, GRA1 was unable to repress the NdmA promoter due to its own promoter being too weak resulting in low ORF4 expression, GRA2, GRA3, GRA4, and GRA5 were all unable to turn on the NdmA promoter after repression due to too high ORF4 expression, and ultimately gave up on the development of the transcription factors.