

An example for users

# Plasmid library

Add

Delete

**2. Add a new one.**

id	Plasmid		Type	Part-only		Backbone		date	Conservation date
	Name	Location		Sequence	Length	Name	Length		
1	AI		rbs-tetr-rr	agaaagaggaga	904	psb1a2	2079	2013/9/25 19:24:43	2013/9/25 19:24:49

**1. Click here to see the plasmid library.**

## Plasmid library

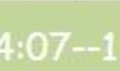
Add

Delete

id	Plasmid		Type	Part-only		Backbone		date	Conservation date
	Name	Location		Sequence	Length	Name	Length		
1	AI		rbs-tetr-rr	agaaagaggaga	904	psb1a2	2079	2013/9/25 19:24:43	2013/9/25 19:24:49



**Fill the form and now you can use the data.**



54:07--1





+

New



New

Try

2013 / 22 12:54:07--1

Try

**Click here to use the data.**





+  
New  
Try

New  
Try



### Add Formwork

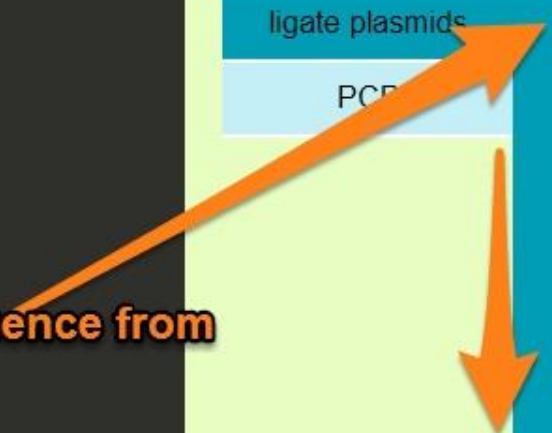
ligate plasmids

PCR

- Choose Plasmids
- Cultivation
- Plasmid Extraction
- Digestion
- Gel Extraction
- Ligation

exit

**Choose them in sequence from the top to the end.**





New



New

Try

2013/9/25 21:06:27--1

Try

### Start to Ligate Plasmids

#### choose plasmid (A)

plasmid	name	
	location	
type	Promoter	
part-only	sequence	
	length	<input type="text"/>
backbone	name	
	length	<input type="text"/>

#### choose plasmid (B)

plasmid	name	
	location	
type	Promoter	
part-only	sequence	
	length	<input type="text"/>
backbone	name	
	length	<input type="text"/>

if you need another plasmid (C) , double click here

**Choose the plasmid.**

2013/9/25  
21:06:56







New



New

Try

2013/9/25 22:55:11--1

Try

### Start to Ligate Plasmids

#### choose plasmid (A)

#### choose plasmid (B)

2013/9/25  
22:55:18

plasmid	name	
	location	
type		
part-only	sequence	
	length	<input type="text"/>
backbone	name	
	length	<input type="text"/>

plasmid	name	
	location	
type		
part-only	sequence	
	length	<input type="text"/>
backbone	name	
	length	<input type="text"/>

**Double click here to get another plasmid.**



if you need another plasmid (C) , double click here

### Cultivation

Plasmid (A )		Plasmid (B )	
Strain		Strain	





New



New

Try

2013/9/25 21:06:27--1

Try

### Start to Ligate Plasmids

#### choose plasmid (A)

#### choose plasmid (B)

#### choose plasmid (C)

plasmid	name		plasmid	name		plasmid	name	
	location			location			location	
type			type			type		
part-only	sequence		part-only	sequence		part-only	sequence	
	length	<input type="text"/>		length	<input type="text"/>		length	<input type="text"/>
backbone	name		backbone	name		backbone	name	
	length	<input type="text"/>		length	<input type="text"/>		length	<input type="text"/>



**That is the plasmid C.**

2013/9/25  
21:06:56

if you don't need the plasmid (C) , double click here

### Cultivation

Plasmid (A )		Plasmid (B )		Plasmid (C )	
Strain		Strain		Strain	



+ New

New

Try

New



background length background length

if you need another plasmid (C) , double click here

**Cultivation**

**Cultivation settings.**

2013/9/25  
22:55:20

Plasmid (A )		Plasmid (B )	
Strain		Strain	
antibiotics	none none none	antibiotics	none none none
Medium		Medium	
Volume	20 μL	Volume	20 μL
Temperature	37 °C	Temperature	37 °C
Speed	200 rpm	Speed	200 rpm

2013/9/25  
22:55:22

Plasmid Extraction	
	A <sub>260/280</sub> ng/μL
(A)	
(B)	



+ New

New

Try

New



22:55:20

Medium		
Volume	20	μL
Temperature	37	°C
Speed	200	rpm

Medium		
Volume	20	μL
Temperature	37	°C
Speed	200	rpm

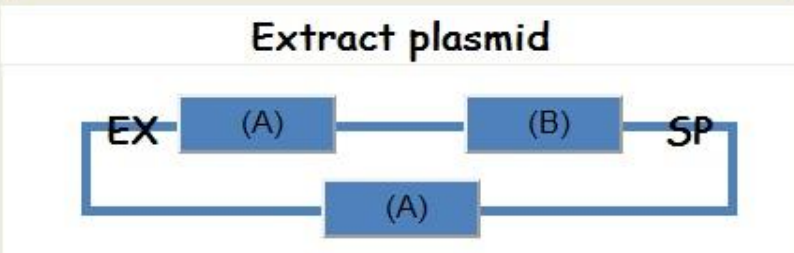


**This help you evaluate plasmid.**

2013/9/25  
22:55:22

Plasmid Extraction	
A <sub>260/280</sub>	ng/μL
(A)	
(B)	

Evaluate



100	μL reaction	(A)	(B)
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2013/9/25

Plasmid	62	62
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+

New



New

Try

22:55:20

Medium		
Volume	20	μL
Temperature	37	°C
Speed	200	rpm

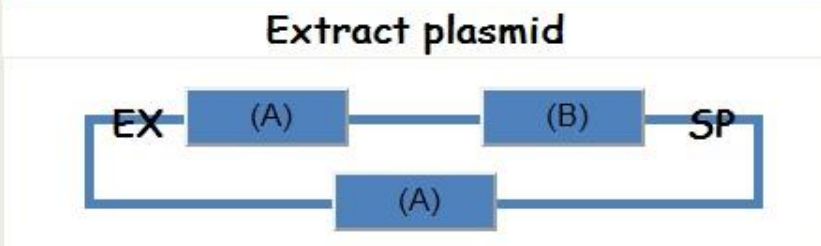
Medium		
Volume	20	μL
Temperature	37	°C
Speed	200	rpm

Extracte plasmid that is used.

2013/9/25  
22:55:22

Plasmid Extraction	
A <sub>260/280</sub>	ng/μL
(A)	
(B)	

Evaluate



100	μL reaction	(A)	(B)
	Plasmid	62	62

2013/9/25





+ New

New

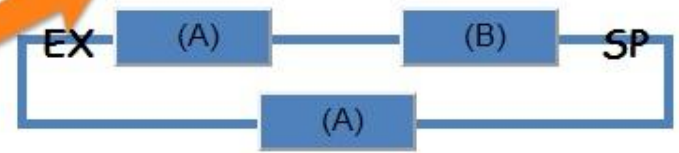
Try

New



Evaluate

Extract plasmid



2013/9/25  
22:55:26

Choose the system.

100	μL reaction	(A)	(B)
	Plasmid	62	62
10	*H Buffer	10	10
	<i>EcoR</i> I / μL	0	0
	<i>Xba</i> I / μL	0	5
	<i>Spe</i> I / μL	5	0
	<i>Pst</i> I / μL	5	5
	H <sub>2</sub> O / μL	18	18

electrophoretogram and gel extraction

Analyse Electrophoretogram

2013/9/25



New



New

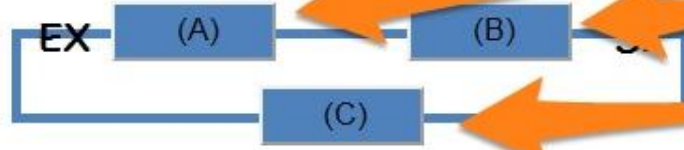
Try

2013/9/25  
21:08:40

	$A_{260/280}$	ng/ $\mu$ L
(A)		
(B)		
(C)		

Evaluate

Extract plasmid



Click on them to change the sequence.

2013/9/25  
21:09:15

100	$\mu$ L reaction	(A)	(B)	(C)
	Plasmid	62	62	62
10	*H Buffer	10	10	10
	<i>EcoR</i> I / $\mu$ L	5	0	5
	<i>Xba</i> I / $\mu$ L	0	5	0
	<i>Spe</i> I / $\mu$ L	5	0	0
	<i>Pst</i> I / $\mu$ L	0	5	5





+ New

New

Try

New



<i>Spe</i> I / $\mu$ L	5	<input type="text" value="0"/>
<i>Pst</i> I / $\mu$ L	5	<input type="text" value="5"/>
H <sub>2</sub> O / $\mu$ L	18	<input type="text" value="18"/>

electrophoretogram and gel extraction

Analyse Electrophoretogram

2013/9/25  
22:55:28

	A <sub>260/280</sub>	ng/ $\mu$ L	bp	
(A)	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	Correct
(B)	<input type="text" value="0"/>	<input type="text" value="0"/>	<input type="text" value="0"/>	

Here you can record the data and use it to analyse for the next step.

2013/9/25  
22:55:30

Ligation

10 <input type="text"/>	$\mu$ L reaction	Insert/Vector=	3 <input type="text"/>	mol/mol
		Recommend		Actual use
	A (vector) / $\mu$ L	<input type="text"/>	<input type="text"/>	<input type="text"/>
	B (insert) / $\mu$ L	<input type="text"/>	<input type="text"/>	<input type="text"/>
10 <input type="text"/>	*Buffer / $\mu$ L	1	<input type="text"/>	<input type="text"/>
	ligase / $\mu$ L	1	<input type="text"/>	<input type="text"/>



New



New

Try

2013/9/25  
21:10:33

### Electrophoretogram and recover

#### Analyse Electrophoretogram

	A <sub>260/280</sub>	ng/ $\mu$ L	bp	
(A)		<input type="text" value="0"/>	<input type="text" value=""/>	Correct
(B)		<input type="text" value="0"/>	<input type="text" value=""/>	

2013/9/25  
21:12:01

choose the photo you want to submit

**Upload the graph you want to show to others.**



选择文件 未选择文件

submit

+

New



New

Try

<i>Spe</i> I / $\mu$ L	5	0
<i>Pst</i> I / $\mu$ L	5	5
H <sub>2</sub> O / $\mu$ L	18	18

electrophoretogram and gel extraction

Analyse Electrophoretogram

2013/9/25  
22:55:28

	A <sub>260/280</sub>	ng/ $\mu$ L	bp	Correct
(A)		0		
(B)		0		

Ligation is set to help you record your data.



Ligation

2013/9/25  
22:55:30

10 $\mu$ L reaction	Insert/Vector= 3 mol/mol										
	<table border="1"> <thead> <tr> <th>Recommend</th> <th>Actual use</th> </tr> </thead> <tbody> <tr> <td>A (vector) / <math>\mu</math>L</td> <td></td> </tr> <tr> <td>B (insert) / <math>\mu</math>L</td> <td></td> </tr> <tr> <td>10 *Buffer / <math>\mu</math>L</td> <td>1</td> </tr> <tr> <td>ligase / <math>\mu</math>L</td> <td>1</td> </tr> </tbody> </table>	Recommend	Actual use	A (vector) / $\mu$ L		B (insert) / $\mu$ L		10 *Buffer / $\mu$ L	1	ligase / $\mu$ L	1
Recommend	Actual use										
A (vector) / $\mu$ L											
B (insert) / $\mu$ L											
10 *Buffer / $\mu$ L	1										
ligase / $\mu$ L	1										



New



New

Try

2013/9/25 22:55:11--1

Try

**Ok! Now you have written a note. Try to design a model by yourself!**

2013/9/25  
22:55:18

### Start to Ligate Plasmids

#### choose plasmid (A)

plasmid	name	
	location	
type		
part-only	sequence	
	length	<input type="text"/>
backbone	name	
	length	<input type="text"/>

#### choose plasmid (B)

plasmid	name	
	location	
type		
part-only	sequence	
	length	<input type="text"/>
backbone	name	
	length	<input type="text"/>

if you need another plasmid (C) , double click here

### Cultivation

Plasmid (A )		Plasmid (B )	
Strain		Strain	

Thank you for  
watching.