

## Experiment 0928

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- ◆ Pick 2 colonies from each of the transformation plates and inoculate in 5mL
- ◆ LB medium + Chloramphenicol. Grow the cells overnight (14 hours) at 37°C and 220 rpm.

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### ◆ Cell growth, sampling, and measure

1. Make a 1:10 dilution of each overnight culture in LB + Chloramphenicol (0.5 mL of culture into 4.5mL of LB + Chlor)
2. Measure Abs600 of these 1:10 diluted cultures
3. Record the data in notebook
- 4.

Abs600 :

	1	2	3	4	5	6	N	P	LB
Colony1	0.155	0.151	0.096	0.111	0.102	0.111	0.122	0.110	0.048
Colony2	0.174	0.117	0.140	0.126	0.116	0.155	0.133	0.111	0.047

5. Dilute the cultures further to a target Abs600 of 0.02 in a final volume of 12 ml LB medium + Chloramphenicol in 50 mL Falcon tube (amber, or covered with foil to block light).

Culture (ml) :

	1	2	3	4	5	6	N	P
Colony1	2.24	2.33	5.00	3.80	4.44	3.81	3.24	3.87
Colony2	1.90	3.48	2.60	3.08	3.53	2.24	2.82	3.81

LB+Cm :

	1	2	3	4	5	6	N	P
Colony1	9.76	9.67	7.00	8.20	7.56	8.19	8.76	8.13
Colony2	10.10	8.52	9.40	8.92	8.47	9.76	9.18	8.19

6. Take 500 µl samples of the diluted cultures into 1.5 ml Eppendorf tubes at 0 hour, prior to incubation. (Take a sample from each of

the 8 devices, two colonies per device, for a total of 16 Eppendorf tubes with 500 µl samples at 0 hour and 6 hours, 32 samples total). Place the samples on ice.

## 7. Measure samples (Abs600 and Fluorescence measurement)

### Abs600 : (0hr)

Hour 0:	Neg. Contr	Pos. Contr	Device 1	Device 2	Device 3	Device 4	Device 5	Device 6	LB + Chlor (blank)
Colony 1, Replicate 1	0.074	0.202	0.065	0.267	0.090	0.099	0.083	0.242	0.115
Colony 1, Replicate 2	0.321	0.087	0.158	0.094	0.056	0.070	0.157	0.113	0.132
Colony 1, Replicate 3	0.108	0.073	0.296	0.076	0.070	0.197	0.241	0.089	0.107
Colony 1, Replicate 4	0.079	0.272	0.100	0.084	0.125	0.117	0.082	0.219	0.062
Colony 2, Replicate 1	0.074	0.063	0.046	0.055	0.043	0.039	0.109	0.080	0.188
Colony 2, Replicate 2	0.083	0.333	0.071	0.092	0.062	0.097	0.275	0.078	0.322
Colony 2, Replicate 3	0.102	0.285	0.060	0.164	0.077	0.327	0.077	0.091	0.328
Colony 2, Replicate 4	0.070	0.066	0.055	0.119	0.124	0.057	0.076	0.081	0.108

### Abs600 : (6hr)

Hour 6:	Neg. Contr	Pos. Contr	Device 1	Device 2	Device 3	Device 4	Device 5	Device 6	LB + Chlor (blank)
Colony 1, Replicate 1	0.701	0.518	0.170	0.620	0.666	0.823	0.382	0.447	0.065
Colony 1, Replicate 2	0.696	0.539	0.289	0.586	0.398	0.589	0.490	0.548	0.065
Colony 1, Replicate 3	0.651	0.634	0.160	0.456	0.513	0.471	0.443	0.439	0.065
Colony 1, Replicate 4	0.712	0.646	0.335	0.455	0.420	0.952	0.489	0.427	0.066
Colony 2, Replicate 1	0.735	0.534	0.385	0.354	0.632	0.883	0.493	0.696	0.065
Colony 2, Replicate 2	0.728	0.665	0.147	0.572	0.723	0.532	0.613	0.718	0.066
Colony 2, Replicate 3	0.687	0.589	0.270	0.508	0.640	0.669	0.495	0.732	0.063
Colony 2, Replicate 4	0.675	0.635	0.134	0.397	0.656	0.667	0.355	0.498	0.070

## ◆Digest pGAPZ A(X3)/ pUCIDT\_Lac1/ pUCIDT\_Px16/ pUCIDT\_Px18 with Agel, EcoRI

### Step 1

Set up reaction for Agel as follows:

	DNA(2µg)	Agel	NEBuffer1.1	ddH <sub>2</sub> O
pGAPZ A(X3)	5.6µl(X3)	1µl (X3)	1.5µl (X3)	6.9µl (X3)
pUCIDT_Lac1	6µl	1µl	1.5µl	6.5µl
pUCIDT_Px16	9µl	1µl	1.5µl	3.5µl
pUCIDT_Px18	11.4µl	1µl	1.5µl	1.1

Incubate at 37°C/ 4hr

### Step 2

Heat inactive 65°C/ 20min

### Step 3

Add the following for EcoRI reaction:

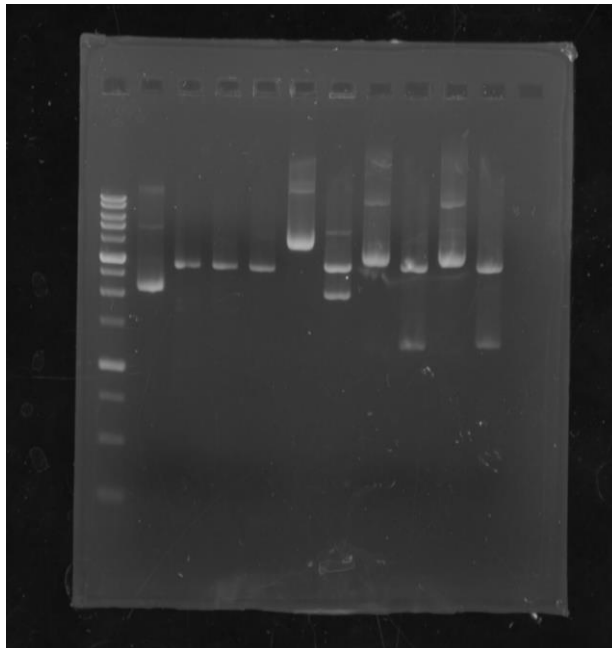
	1M NaCl	EcoRI	ddH <sub>2</sub> O
pGAPZ A(X3)	1µl (X3)	0.5µl	3.5µl
pUCIDT_Lac1	1µl	0.5µl	3.5µl

pUCIDT_Px16	1µl	0.5µl	3.5µl
pUCIDT_Px18	1µl	0.5µl	3.5µl

Incubate at 37°C/ 1.5hr

Step 4

Check with gel electrophoresis



- 1-1 marker
- 1-2 pGAPZ A uncut
- 1-3 pGAPZ A(EcoR I , Age I ) digest
- 1-4 pGAPZ A(EcoR I , Age I ) digest
- 1-5 pGAPZ A(EcoR I , Age I ) digest
- 1-6 pUCIDT\_Lac1 uncut
- 1-7 pUCIDT\_Lac1(EcoR I , Age I )  
digest
- 1-8 pUCIDT\_Px16 uncut
- 1-9 pUCIDT\_Px16 (EcoR I , Age I )  
digest
- 1-10 pUCIDT\_Px18 uncut
- 1-11 pUCIDT\_Px18 (EcoR I , Age I )  
digest

Step 5

Heat inactivate 65°C/ 20min

◆NanoDrop

	OD260/280	ng/µl
pGAPZ A cut_(1)	1.52	12.5
pGAPZ A cut_(2)	1.69	12.1
pGAPZ A cut_(3)	1.77	19.1
Lac1 cut	1.56	16.6
Px16 cut	1.76	18.2
Px18 cut	1.69	15.1

## 1. Fluorescence measurement

Fluorescence : (0hr)

Hour 0:	Neg. Contr	Pos. Contr	Device 1	Device 2	Device 3	Device 4	Device 5	Device 6	LB + Chlor (
Colony 1, Replicate 1	155	233	294	274	150	348	166	253	19
Colony 1, Replicate 2	160	165	292	256	154	333	164	179	20
Colony 1, Replicate 3	161	166	296	259	148	326	167	189	19
Colony 1, Replicate 4	151	165	340	258	170	355	254	194	19
Colony 2, Replicate 1	154	167	297	195	159	232	186	164	18
Colony 2, Replicate 2	159	173	349	212	159	289	201	157	18
Colony 2, Replicate 3	154	178	361	198	181	268	247	181	20
Colony 2, Replicate 4	158	172	323	288	176	267	175	173	19

Fluorescence : (6hr)

Hour 6:	Neg. Contr	Pos. Contr	Device 1	Device 2	Device 3	Device 4	Device 5	Device 6	LB + Chlor (
Colony 1, Replicate 1	202	875	1229	2564	218	5431	547	1014	105
Colony 1, Replicate 2	209	860	1268	2616	160	5376	509	869	117
Colony 1, Replicate 3	206	908	1256	2424	226	3951	511	918	99
Colony 1, Replicate 4	206	807	1276	2660	227	7369	508	872	117
Colony 2, Replicate 1	197	902	1394	1606	230	4239	563	682	108
Colony 2, Replicate 2	193	860	1348	1929	231	2939	583	677	164
Colony 2, Replicate 3	194	746	1384	1580	236	2798	644	686	141
Colony 2, Replicate 4	201	839	1364	1693	205	2772	666	673	160

## ◆Ligation

Set up the following reaction in a microcentrifuge tube on ice. (Insert: Vector = 3:1)

	T4 Ligase	Ligase buffer	Insert	Vector 50ng	ddH2O
pGAPZ A_Lac1	1µl	2µl	6.6µl 110ng	2.6µl	2.8µl
pGAPZ A_Px16	1µl	2µl	3.8µl 70ng	2.6µl	5.6µl
pGAPZ A_Px18	1µl	2µl	4.6µl 70ng	2.6µl	4.8µl

Incubate at 16°C/ 15hr

Pick 2 colonies from each of the transformation plates and inoculate in 5mL LB medium + Chloramphenicol. Grow the cells overnight (16 hours) at 37°C and 220 rpm.

### ◆Transformation

1. Add all pGAPZ A\_Lac1/ pGAPZ A\_Px16/ pGAPZ A\_Px18 (15µl) into 20µl ECOS™ 101 Competent Cells [DH5α]
2. Incubate on ice 5min
3. Heat shock at 42°C 45sec
4. Incubate on ice 5min
5. Add 140µl LB
6. Incubate at 37°C 1hr
7. Spread on LB+ Zeocin plate

### Ligation

Set up the following reaction in a microcentrifuge tube on ice. (Insert: Vector = 3: 1)

	T4 Ligase	Ligase buffer	Insert	Vector	ddH2O
pGAPZ A_Lac1	1µl	2.5µl	13.2µl 170ng	5.2µl 75ng	3.1µl
pGAPZ A_Px16	1µl	2µl	7.6µl 144ng	5.2µl 100ng	4.2µl
pGAPZ A_Px18	1µl	2µl	9.2µl 144ng	5.2µl 100ng	2.6µl

And incubate at 16°C/ 15hr

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### ◆Cell growth, sampling, and measure

1. Make a 1:10 dilution of each overnight culture in LB + Chloramphenicol (0.5mL of culture into 4.5mL of LB + Chlor)
2. Measure Abs600 of these 1:10 diluted cultures
3. Record the data in notebook

Abs600 :

	1	2	3	4	5	6	N	P	LB
Colony1	0.1494	0.1235	0.1169	0.1196	0.1275	0.1123	0.1088	0.1181	0.0445

Colony2	0.1282	0.1306	0.1333	0.1216	0.1438	0.1145	0.1312	0.1175	0.0432
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- Dilute the cultures to 0.02(Abs600) in a final volume of 12 ml LB medium + Chloramphenicol in 50 mL Falcon tube (amber, or covered with foil to block light).

Culture (ml) :

	1	2	3	4	5	6	N	P
Colony1	2.29	3.00	2.78	2.75	2.70	3.43	3.24	3.36
Colony2	2.87	2.79	2.70	3.11	2.42	3.43	2.77	3.29

LB+Cm :

	1	2	3	4	5	6	N	P
Colony1	9.71	9.00	9.22	9.25	9.30	8.57	8.76	8.64
Colony2	9.13	9.21	9.3	8.89	9.58	8.57	9.23	8.71

- Take 500  $\mu$ L samples of the diluted cultures at 0 hours into 1.5 ml Eppendorf tubes, prior to incubation. (At 0 hour and 6 hours, you will take a sample from each of the 8 devices, two colonies per device, for a total of 16 Eppendorf tubes with 500  $\mu$ L samples per time point, 32 samples total). Place the samples on ice.
- Measure samples (Abs600 and Fluorescence measurement)

Abs600 : (0hr)

Hour 0:	Neg. Contrc	Pos. Contrc	Device 1	Device 2	Device 3	Device 4	Device 5	Device 6	LB + Chlor (blank)
Colony 1, Replicate 1	0.346	0.060	0.223	0.360	0.277	0.242	0.285	0.355	0.106
Colony 1, Replicate 2	0.358	0.387	0.242	0.379	0.386	0.354	0.302	0.316	0.351
Colony 1, Replicate 3	0.291	0.374	0.154	0.378	0.246	0.348	0.388	0.054	0.346
Colony 1, Replicate 4	0.372	0.402	0.274	0.284	0.344	0.291	0.191	0.398	0.257
Colony 2, Replicate 1	0.332	0.404	0.400	0.375	0.336	0.386	0.170	0.358	0.333
Colony 2, Replicate 2	0.310	0.227	0.124	0.375	0.349	0.339	0.185	0.239	0.268
Colony 2, Replicate 3	0.181	0.367	0.381	0.406	0.385	0.298	0.417	0.281	0.229
Colony 2, Replicate 4	0.377	0.307	0.090	0.176	0.375	0.272	0.194	0.305	0.223

Abs600 : (6hr)

Hour 6:	Neg. Contrc	Pos. Contrc	Device 1	Device 2	Device 3	Device 4	Device 5	Device 6	LB + Chlor (blank)
Colony 1, Replicate 1	0.521	0.509	0.183	0.574	0.688	0.464	0.114	0.552	0.045
Colony 1, Replicate 2	0.531	0.668	0.181	0.618	0.717	0.458	0.115	0.593	0.033
Colony 1, Replicate 3	0.546	0.556	0.378	0.642	0.821	0.542	0.131	0.587	0.032
Colony 1, Replicate 4	0.522	0.535	0.311	0.685	0.770	0.515	0.189	0.619	0.046
Colony 2, Replicate 1	0.684	0.591	0.109	0.682	0.598	0.614	0.144	0.670	0.035
Colony 2, Replicate 2	0.746	0.556	0.236	0.700	0.426	0.609	0.131	0.638	0.218
Colony 2, Replicate 3	0.706	0.678	0.306	0.733	0.537	0.516	0.137	0.557	0.113
Colony 2, Replicate 4	0.597	0.552	0.148	0.646	0.418	0.506	0.092	0.555	0.030

### ◆Transformation

1. Add all pGAPZ A\_Lac1/ pGAPZ A\_Px16/ pGAPZ A\_Px18 (20µl) into 20µl Competent Cells DH5a and pGAPZ A 1µl+19µl ddH<sub>2</sub>O into 10µl Competent Cells DH5a as positive control.
2. Incubate on ice 15min
3. Heat shock at 42°C 45sec
4. Incubate on ice 5min
5. Add 140µl LB
6. Incubate at 37°C 1hr
7. Spread on LB+ Zeocin plate

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### ◆Digest pUCIDT\_Lac1/ pUCIDT\_Px16/ pUCIDT\_Px18 with AgeI, EcoRI

#### Step 1

Set up reaction for AgeI as follows:

	DNA(5µg)	AgeI	NEBuffer1.1	ddH <sub>2</sub> O
pUCIDT_Lac1	15.0µl	1.0µl	2.0µl	2.0µl
pUCIDT_Px16	24.2µl	2.0µl	3.0µl	1.8µl
pUCIDT_Px18	23.0µl	1.0µl	2.6µl	0

Incubate at 37°C/ 6hr

#### Step 2

Heat inactivate 65°C/ 20min

#### Step 3

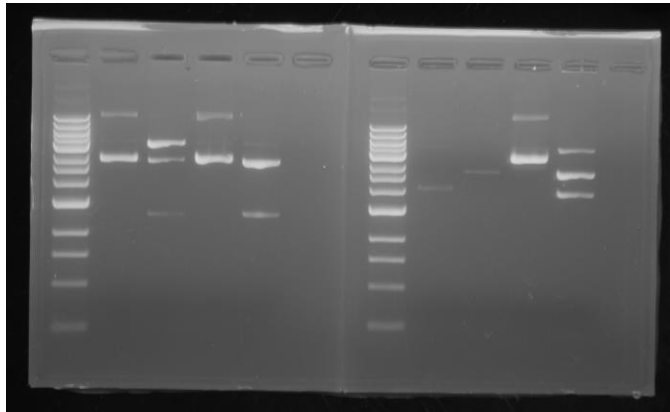
Add the following for EcoRI reaction:

	1M NaCl	EcoRI	ddH <sub>2</sub> O
pUCIDT_Lac1	1.0µl	0.5µl	3.5µl
pUCIDT_Px16	1.0µl	0.5µl	3.5µl
pUCIDT_Px18	1.0µl	0.5µl	3.5µl

Incubate at 37°C/ 2.5hr

**Step 4**

Check with gel electrophoresis



- 1-1 2-1marker
- 1-2 pUCIDT\_Px16 uncut
- 1-3 pUCIDT\_Px16 cut
- 1-4 pUCIDT\_Px18 uncut
- 1-5 pUCIDT\_Px18 cut
- 2-2 pGAPZ A uncut
- 2-3 pGAPZ A cut
- 2-4 pUCIDT\_Lac1 uncut
- 2-5 pUCIDT\_Lac1 cut

**Step 5**

Heat inactivate 65°C/ 20min

**◆NanoDrop**

	OD260/280	ng/μl
Lac1 cut	1.72	33.8
Px16 cut	1.68	19.7
Px18 cut	1.81	30.4

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**◆Ligation**

Set up the following reaction in a microcentrifuge tube on ice. (Insert: Vector = 5: 1)

	T4 Ligase	Ligase buffer	Insert	Vector	ddH2O
pGAPZ A_Lac1	1.0μl	1.5μl	5.0μl 170ng	5.2μl 80ng	2.3μl
pGAPZ A_Px16	1.0μl	1.5μl	7.3μl 144ng	5.2μl 80ng	0μl
pGAPZ	1.0μl	1.5μl	4.8μl	5.2μl 80ng	2.5μl



A_Px18			144ng		
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Incubate at 16°C/ 18hr

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◆Transformation

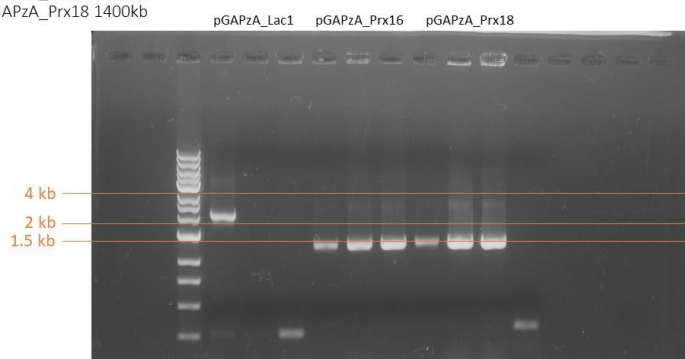
pGAPZ A\_Lac1/ pGAPZ A\_Px16/ pGAPZ A\_Px18

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◆pGAPZ A\_Lac1-pGAPZ A\_Px16-pGAPZ A\_Px18-clone

20180813 PCR (pGAPZ A\_Lac1, pGAPZ A\_Px16, pGAPZ-A\_Px18)

20189813 21:00 colony PCR Result (45° C)  
 pGAPzA\_Lac1 2072kb  
 pGAPzA\_Prx16 1300kb  
 pGAPzA\_Prx18 1400kb



Colony PCR

Result

pGAPZ A\_Lac1

2072kb

pGAPZ A\_Px16

1300kb

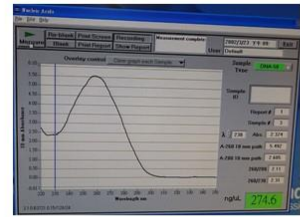
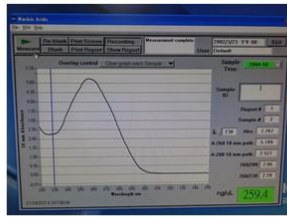
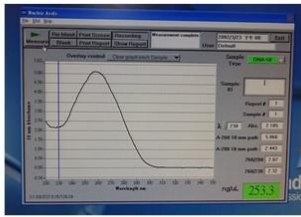
pGAPZ A\_Px18

1400kb

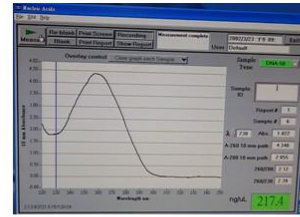
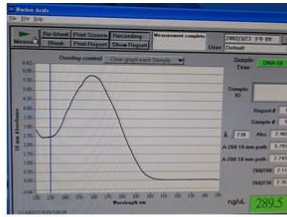
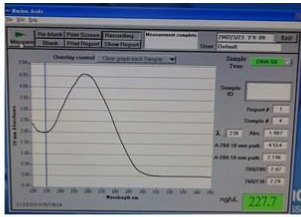
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◆Miniprep

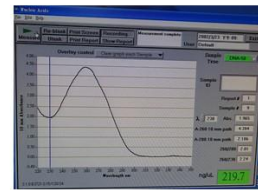
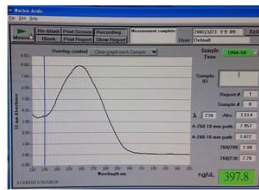
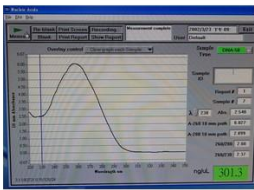
Result:



Lac1(1) OD<sub>260/280</sub>=2.07   Lac1(2) OD<sub>260/280</sub>=2.05   Lac1(3) OD<sub>260/280</sub>=2.11



Prx16(1) OD<sub>260/280</sub>=2.07   Prx16(2) OD<sub>260/280</sub>=2.11   Prx16(3) OD<sub>260/280</sub>=2.12



Prx18(1) OD<sub>260/280</sub>=2.37   Prx18(2) OD<sub>260/280</sub>=2.08   Prx18(3) OD<sub>260/280</sub>=2.01

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◆Digest (use Agel, EcoRI)

Protocol reference: <https://nebcloner.neb.com/#/protocol/re/sequential-heat/Agel,EcoRI>

	(1ng)	EcoRI	Agel	10Xbuffer	ddH <sub>2</sub> O	Total V
Lac1 1	4μl	0.5μl	0.4μl	2μl	13.1	20μl
2	4μl				13.1	
3	4μl				13.1	
Prx16 1	4.5μl				12.6	
2	4μl				13.1	
3	5μl				12.1	
Prx18 1	3.5μl	13.6				
2	3μl	14.1				

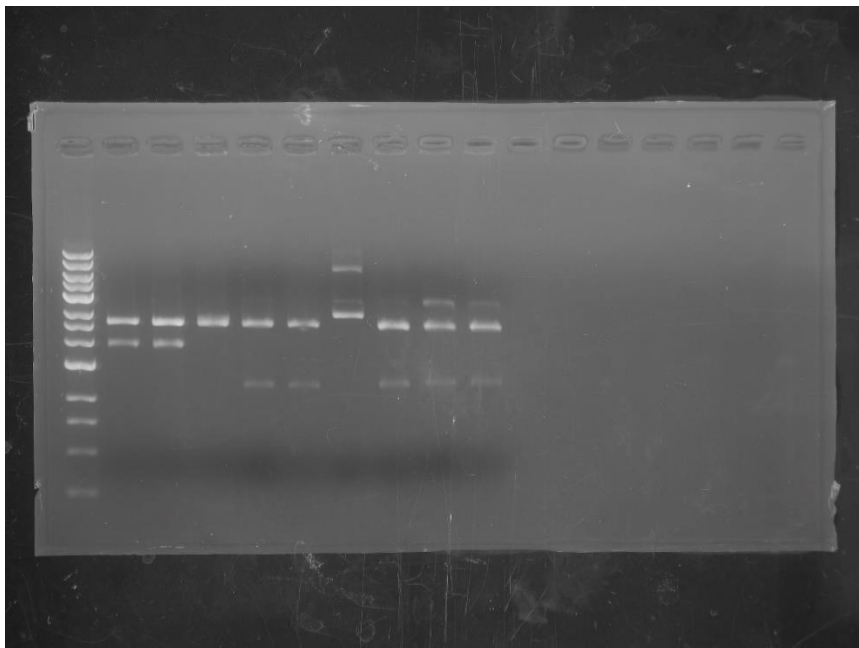
3	5 $\mu$ l				12.1	
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Set up reaction for first restriction enzyme\* as follows:

COMPONENT	20 $\mu$ l REACTION
DNA	1 ng
10X NEBuffer 1.1	2 $\mu$ l (1X)
Agel	0.4 $\mu$ l (or 10 units)
Nuclease-free Water	to 20 $\mu$ l

1. Incubate at 37°C for 2 hours.
2. Heat inactivate by incubating at 65°C for 20 minutes.
3. Add 50 mM NaCl(2 $\mu$ l) to bring the salt concentration of NEBuffer EcoRI.  
Add 0.5 $\mu$ l of EcoRI.
4. Incubate at 37°C for 2hr

◆Digest result

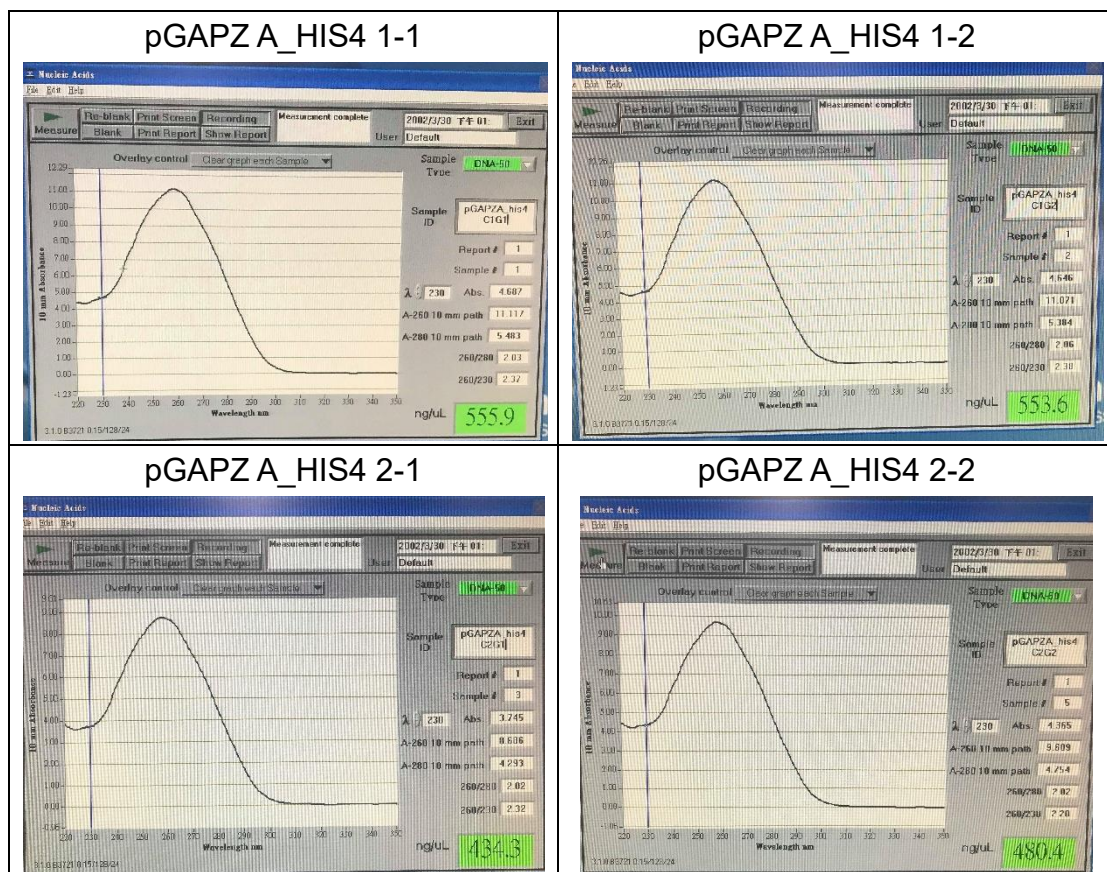


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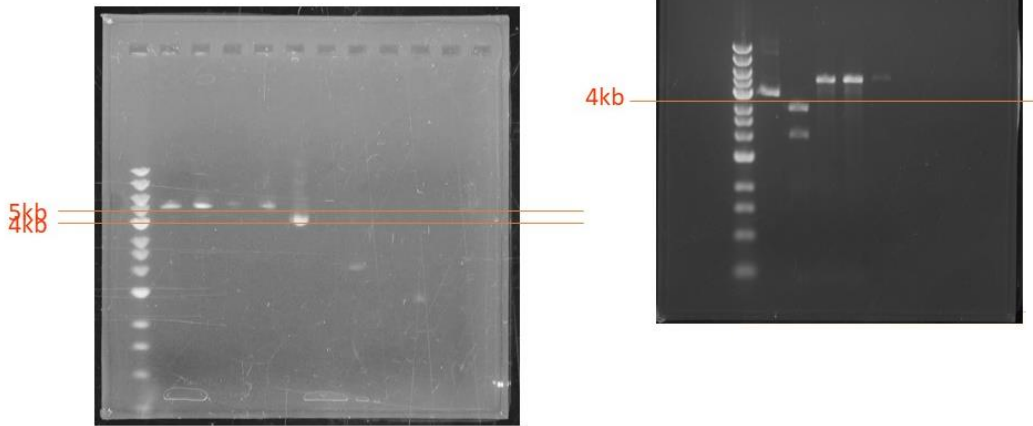
◆ Digestion (pGAPZ A\_HIS4 cut Agel, EcoR)

	DNA(3μl)	Agel (5unit/μl)	EcoRI (20unit/μl)	10Xbuffer1.1	ddH2O	Total V
1-1-1	5.4μl	0.6μl	0.4μl	2 μl	11.6 μl	20 μl
1-1-2	5.4μl					
1-2-1	5.4μl					
1-2-2	5.4μl					

◆ DNA quality

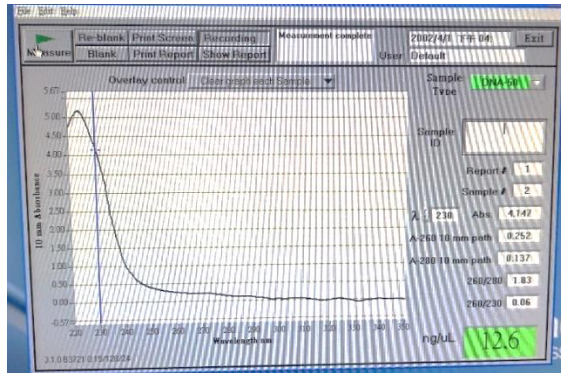
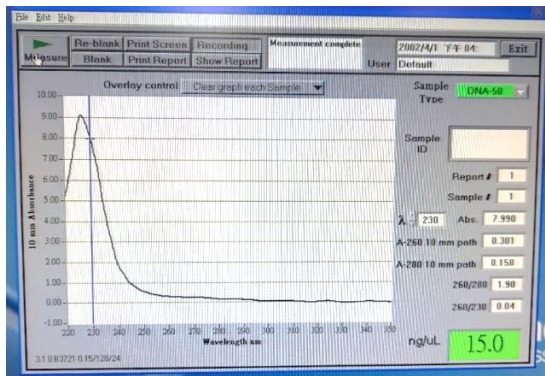


pGAPzA\_His4 5183bp  
 EcoRI\_AgeI 193bp  
 5183-193= 4990bp



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◆Result of gel extract



◆Ligation

insert	pGAPZA_HIS4 (vector)	(insert)	ligase	10Xligationn buffer	ddH <sub>2</sub> O	Total V
Lac1	5.0μl	5μl	0.5μl	2.0μl	7.5μl	20.0μl
Prx16	5.0μl	5.5μl	0.5μl	2.0μl	7.0μl	
Prx18	5.0μl	3.5μl	0.5μl	2.0μl	9.0μl	

8/24



◆ Transform over night

8/27

Digestion – pGAPZ A+HIS4 cut with Agel and EcoRI (x3)	Ligation - vector: insert = 1:3																								
<table border="1"> <tr><td>DNA 3µg</td><td>5.5 µl</td></tr> <tr><td>Agel (5 units/µl)</td><td>1 µl</td></tr> <tr><td>NEbuffer 1.1</td><td>1.5 µl</td></tr> <tr><td>ddH<sub>2</sub>O</td><td>7 µl</td></tr> <tr><td>Total</td><td>15 µl</td></tr> <tr><td>T : 37 , 8hr</td><td></td></tr> </table>	DNA 3µg	5.5 µl	Agel (5 units/µl)	1 µl	NEbuffer 1.1	1.5 µl	ddH <sub>2</sub> O	7 µl	Total	15 µl	T : 37 , 8hr		<table border="1"> <tr><td>Px18 106.1ng(30.4ng/µl)</td><td>3.5 µl</td></tr> <tr><td>DNA vector 75ng (15ng/µl)</td><td>5 µl</td></tr> <tr><td>Ligase</td><td>0.5 µl</td></tr> <tr><td>Ligation buffer</td><td>2 µl</td></tr> <tr><td>ddH<sub>2</sub>O</td><td>9 µl</td></tr> </table>	Px18 106.1ng(30.4ng/µl)	3.5 µl	DNA vector 75ng (15ng/µl)	5 µl	Ligase	0.5 µl	Ligation buffer	2 µl	ddH <sub>2</sub> O	9 µl		
DNA 3µg	5.5 µl																								
Agel (5 units/µl)	1 µl																								
NEbuffer 1.1	1.5 µl																								
ddH <sub>2</sub> O	7 µl																								
Total	15 µl																								
T : 37 , 8hr																									
Px18 106.1ng(30.4ng/µl)	3.5 µl																								
DNA vector 75ng (15ng/µl)	5 µl																								
Ligase	0.5 µl																								
Ligation buffer	2 µl																								
ddH <sub>2</sub> O	9 µl																								
<table border="1"> <tr><td>NaCl</td><td>1 µl</td></tr> <tr><td>EcoRI (20units/µl)</td><td>0.25 µl</td></tr> <tr><td>DNA</td><td>15 µl</td></tr> <tr><td>NEbuffer EcoRI</td><td>2 µl</td></tr> <tr><td>ddH<sub>2</sub>O</td><td>2 µl</td></tr> <tr><td>Total</td><td>20 µl</td></tr> <tr><td>T :37 , 4hr</td><td></td></tr> </table>	NaCl	1 µl	EcoRI (20units/µl)	0.25 µl	DNA	15 µl	NEbuffer EcoRI	2 µl	ddH <sub>2</sub> O	2 µl	Total	20 µl	T :37 , 4hr		<table border="1"> <tr><td>Px16 107.3ng (19.7ng/µl)</td><td>5.5 µl</td></tr> <tr><td>DNA vector 75ng (15ng/µl)</td><td>5 µl</td></tr> <tr><td>Ligase</td><td>0.5 µl</td></tr> <tr><td>Ligation buffer</td><td>2 µl</td></tr> <tr><td>ddH<sub>2</sub>O</td><td>7 µl</td></tr> </table>	Px16 107.3ng (19.7ng/µl)	5.5 µl	DNA vector 75ng (15ng/µl)	5 µl	Ligase	0.5 µl	Ligation buffer	2 µl	ddH <sub>2</sub> O	7 µl
NaCl	1 µl																								
EcoRI (20units/µl)	0.25 µl																								
DNA	15 µl																								
NEbuffer EcoRI	2 µl																								
ddH <sub>2</sub> O	2 µl																								
Total	20 µl																								
T :37 , 4hr																									
Px16 107.3ng (19.7ng/µl)	5.5 µl																								
DNA vector 75ng (15ng/µl)	5 µl																								
Ligase	0.5 µl																								
Ligation buffer	2 µl																								
ddH <sub>2</sub> O	7 µl																								
	<table border="1"> <tr><td>Lac1 169.8ng (33.8ng/µl)</td><td>5 µl</td></tr> <tr><td>DNA vector 75ng (15ng/µl)</td><td>5 µl</td></tr> <tr><td>Ligase</td><td>0.5 µl</td></tr> <tr><td>Ligation buffer</td><td>2 µl</td></tr> <tr><td>ddH<sub>2</sub>O</td><td>7.5 µl</td></tr> </table>	Lac1 169.8ng (33.8ng/µl)	5 µl	DNA vector 75ng (15ng/µl)	5 µl	Ligase	0.5 µl	Ligation buffer	2 µl	ddH <sub>2</sub> O	7.5 µl														
Lac1 169.8ng (33.8ng/µl)	5 µl																								
DNA vector 75ng (15ng/µl)	5 µl																								
Ligase	0.5 µl																								
Ligation buffer	2 µl																								
ddH <sub>2</sub> O	7.5 µl																								

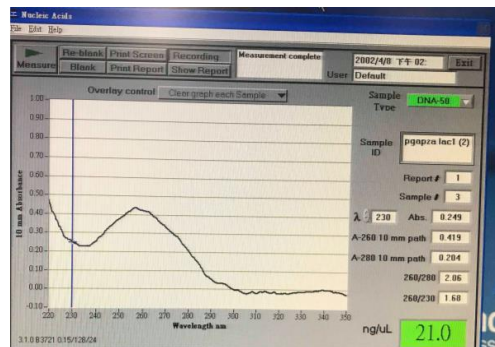
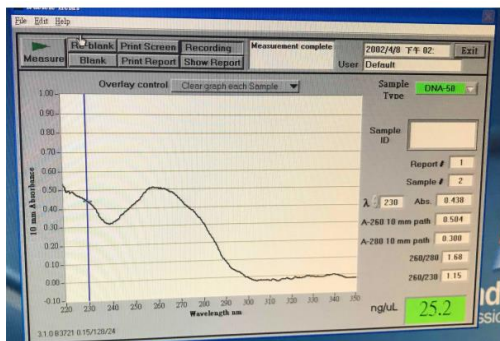
8/28

◆ Transform (LB + zeocin)

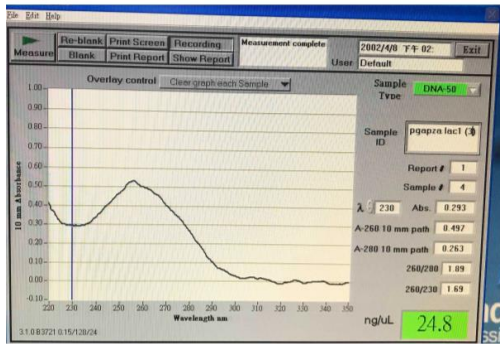
1. pGAPZ A\_HIS4\_Px16
2. pGAPZ A\_HIS4\_Px18
3. pGAPZ A\_HIS4\_Lac1

8/29

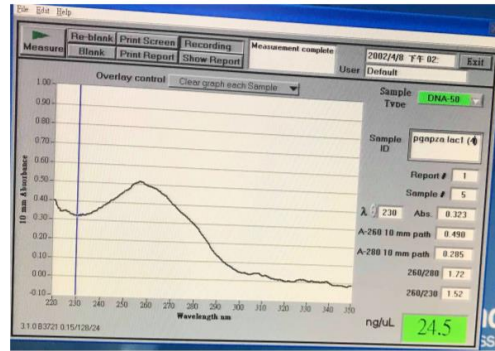
◆ Nanodrop



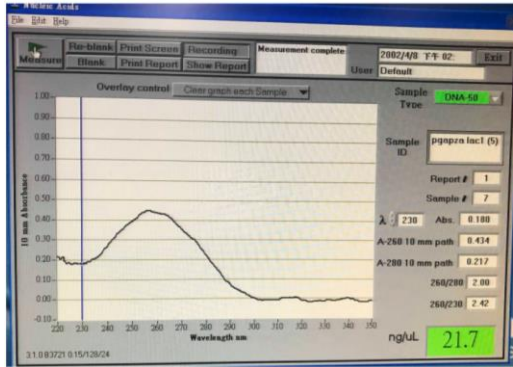
pGAPZ A + Lac1 (1)



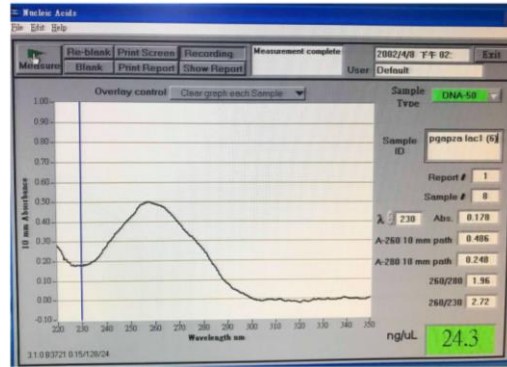
pGAPZ A + Lac1 (2)



pGAPZ A + Lac1 (3)



pGAPZ A + Lac1 (4)



pGAPZ A + Lac1 (5)

8/30

◆Check by Agel

pGAPZ A + HIS4

9/1

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

6x his	SOUP/PELLET	LOAD	ZEOCIN
1:20000	SOUP	LMH	o

◆Digest pGAPZ A HIS4(X2) with Agel, EcoRI

Step 1

Set up reaction for Agel as follows:

	DNA(3µg)	AgeI	NEBuffer1.1	ddH <sub>2</sub> O
pGAPZ A(X2)	6.5µl(X3)	1.0µl (X3)	1.5µl (X3)	7.0µl (X3)

incubate at 37°C/ 6.5hr

## Step 2

Add following for EcoRI reaction:

	1M NaCl	EcoRI	ddH <sub>2</sub> O
pGAPZ A(X2)	1.0µl (X2)	0.25µl	2.0µl

incubate at 37°C/ 8hr

## ◆Ligation

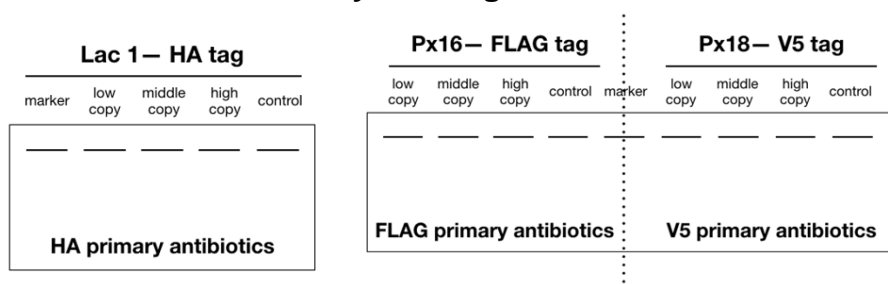
Set up the following reaction in a microcentrifuge tube on ice. (Insert: Vector = 3: 1)

	T4 Ligase	Ligase buffer	Insert	Vector 75ng	ddH <sub>2</sub> O
pGAPZ A HIS4_Lac1	1.0µl	2.0µl	6.0µl	6.0µl	5.0µl
pGAPZ A HIS4_Px16	1.0µl	2.0µl	6.0µl	6.0µl	5.0µl
pGAPZ A HIS4_Px18	1.0µl	2.0µl	6.0µl	6.0µl	5.0µl

incubate at 16°C/ 18hr

9/2

## ◆Western blot: Antibody staining → Chemiluminescence detection



## ◆Transformation pGAPZ A HIS4\_Lac1/ pGAPZ A HIS4\_Px16/ pGAPZ A HIS4\_Px18 (15µl) into 15µl ECOS™ 101 Competent Cells



◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining

6x his	SOUP/PELLET	LOAD	ZEOCIN
1:10000	SOUP	LMH	x

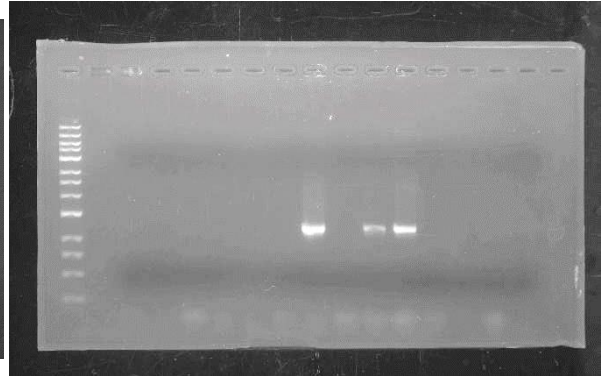
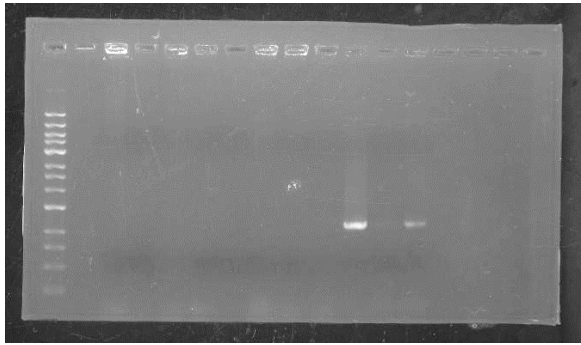
◆1.PCR—pGAPZ A\_HIS4 + Px16, Px18, Lac1

Component	Final(250µl)
10X Standard <i>Taq</i> Reaction Buffer	25µl
10 mM dNTPs	5µl
0.2 µM Primer suffix	5µl
0.2 µM Primer prefix	5µl
Template DNA	pGAPZ A_HIS4_Px16 pGAPZ A_HIS4_Px18 pGAPZ A_HIS4_Lac1
5 units/µl <i>Taq</i> DNA Polymerase	3.8µl
Nuclease-free water	207.5µl

Distribute to 25 eppendorf tubes, each of the tubes contains 10µl

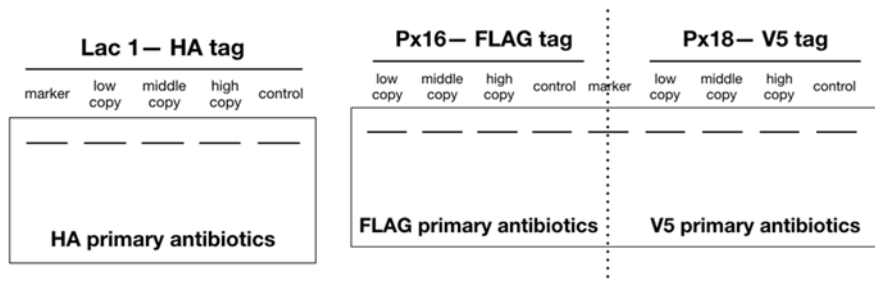
STEP	TEMP	TIME
Initial Denaturation	95°C	30 seconds
30 Cycles	95°C	30 seconds
	45°C	30 seconds
	72°C	2.5minutes
Final Extension	72°C	10 minutes
Hold	4°C	

## 2.PCR Check



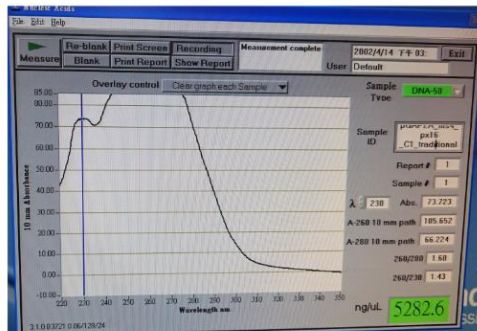
9/4

### ◆Western blot: Antibody staining → Chemiluminescence detection

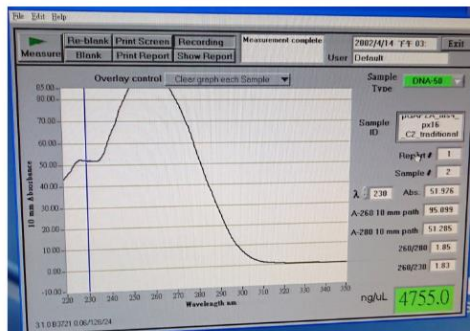


miniprep – pGAPZ A\_HIS4 + Px16, Px18

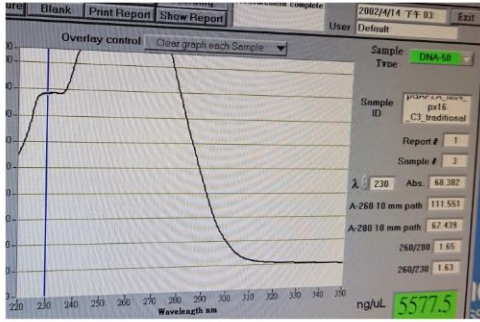
Traditional:



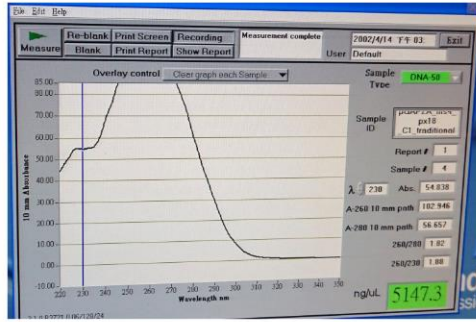
pGAPZ A\_HIS4\_Px16 (1)



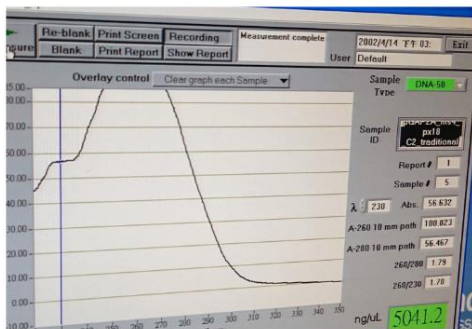
pGAPZ A-1\_HIS4\_Px16 (2)



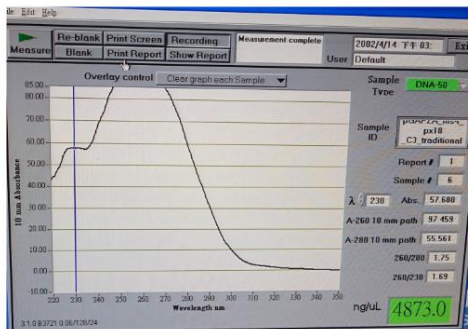
pGAPZ A\_HIS4\_Px16 (3)



pGAPZ A\_HIS4\_Px18 (1)

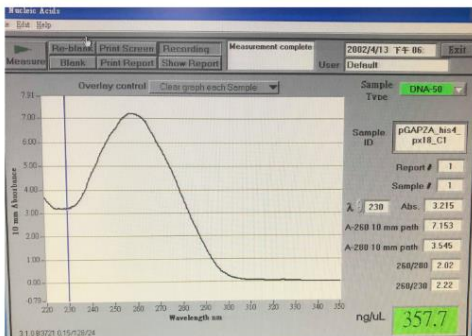


pGAPZ A\_HIS4\_Px18 (2)

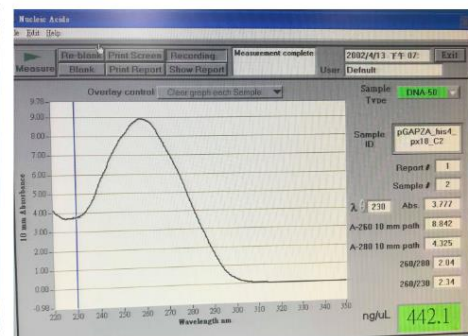


pGAPZ A\_HIS4\_Px18 (3)

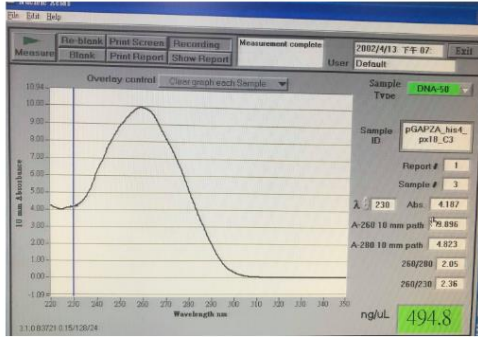
Miniprep:



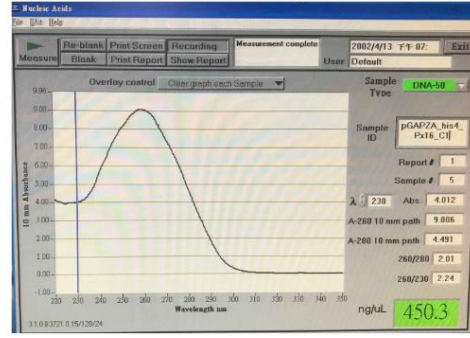
pGAPZ A\_HIS4\_Px18 (1)



pGAPZ A1\_HIS4\_Px18 (2)



pGAPZ A\_HIS4\_Px18 (3)



pGAPZ A\_HIS4\_Px16

9/5

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:5000	1:5000	1:5000	SOUP	LMH	x

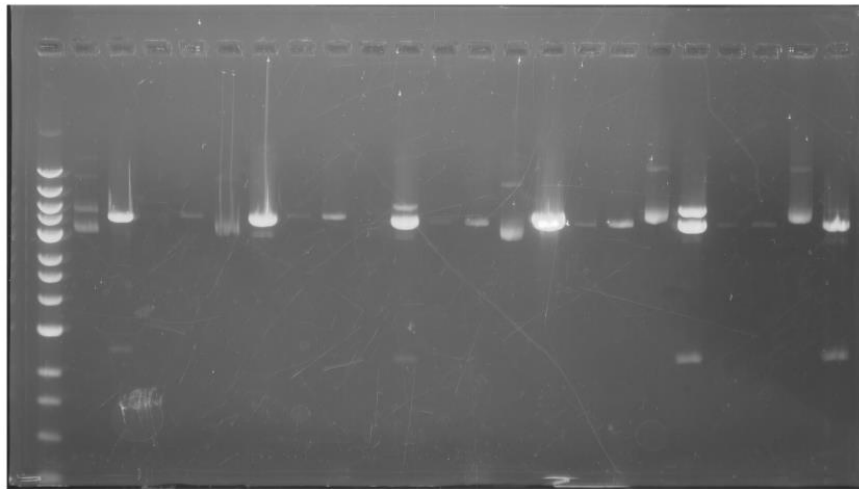
◆Transform – pGAPZ A\_HIS4\_Lac1(20μl competent cell, 20μl DNA)

DNA 3μg	5.5 μl
Agel	0.6 μl
NEbuffer 1.1	1.5 μl
ddH <sub>2</sub> O	7.4 μl
Total	15 μl
T : 37, 6hr	

NaCl	1 μl
EcoRI	0.25 μl
DNA	15 μl
ddH <sub>2</sub> O	2 μl
NEbuffer EcoRI	2 μl
Total	20 μl
T : 37, 10hr	

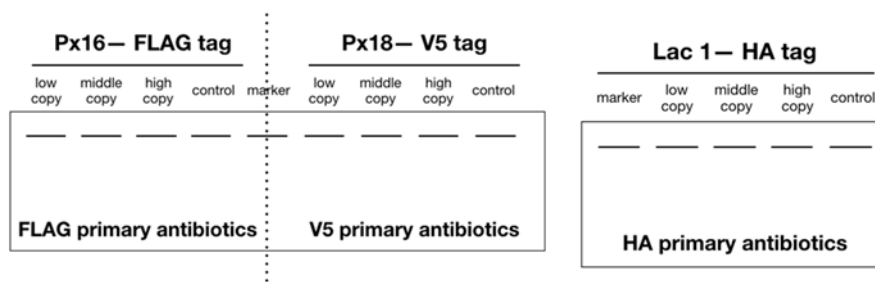
9/6

◆ Digestion – pGAPZ A\_HIS4\_lac1



- 1-12. pGAPZ A\_HIS4\_Px16
- 1.5.9 miniprep uncut
- 2.6.10 miniprep cut
- 3.7.11 traditional uncut
- 4.8.12 traditional cut
- 13-22. pGAPZ A\_HIS4\_Px18
- 13.17.21 miniprep uncut
- 14.18.22 miniprep cut
- 15.19 traditional uncut
- 16.20 traditional cut

◆ Western blot: Antibody staining → Chemiluminescence detection



9/7

◆ Prepare yeast

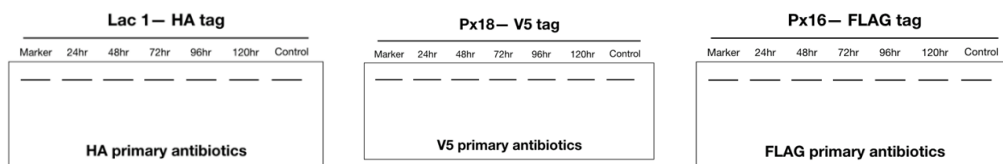
Pick up yeast(SD1168) from a colony + 50 ml YPD in erlenmeyer flask,  
Breeding in incubator in 30°C for days 2.5

◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:2000	1:2000	1:2000	SOUP	Time point	x

9/8

◆Western blot: Antibody staining → Chemiluminescence detection



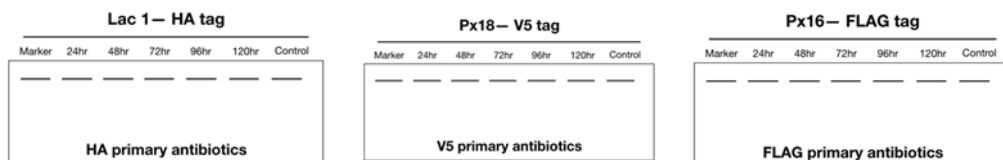
9/9

◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:1000	1:1000	PELLET	Time point	x

9/10

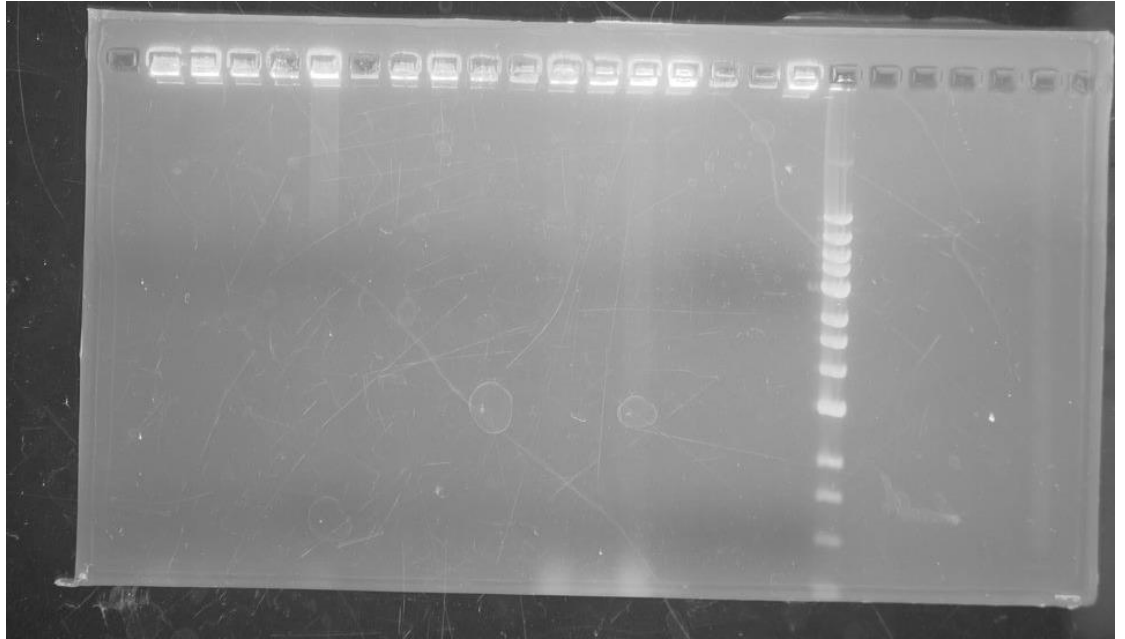
◆Western blot: Antibody staining → Chemiluminescence detection



1.Transform target genes into yeast → spread on SD plate

Plates:

1. Pichia pastoris
2. Pichia pastoris+ pGAPZ A\_his4
3. Pichia pastoris+ pGAPZ A\_his4\_px18



>> FAIL 😞, cuz we pick too much bacteria

## 2.Minimal culturing of lac1

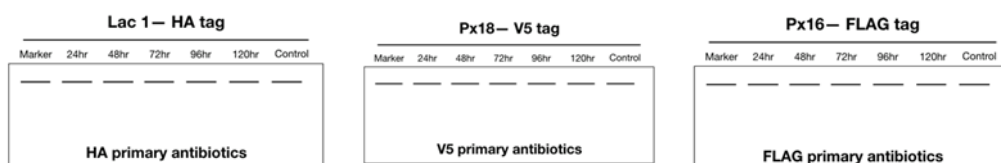
9/11

◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:5000	1:5000	SOUP	Time point	o

9/12

◆Western blot: Antibody staining → Chemiluminescence detection



9/13

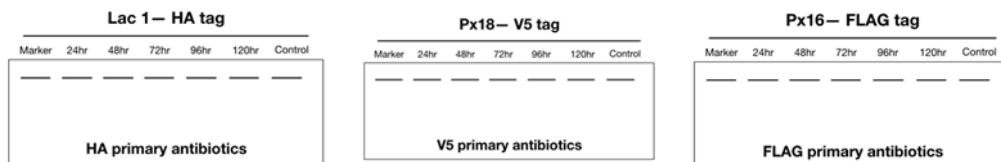
◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining



FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	SOUP	Time point	o

9/14

◆Western blot: Antibody staining → Chemiluminescence detection



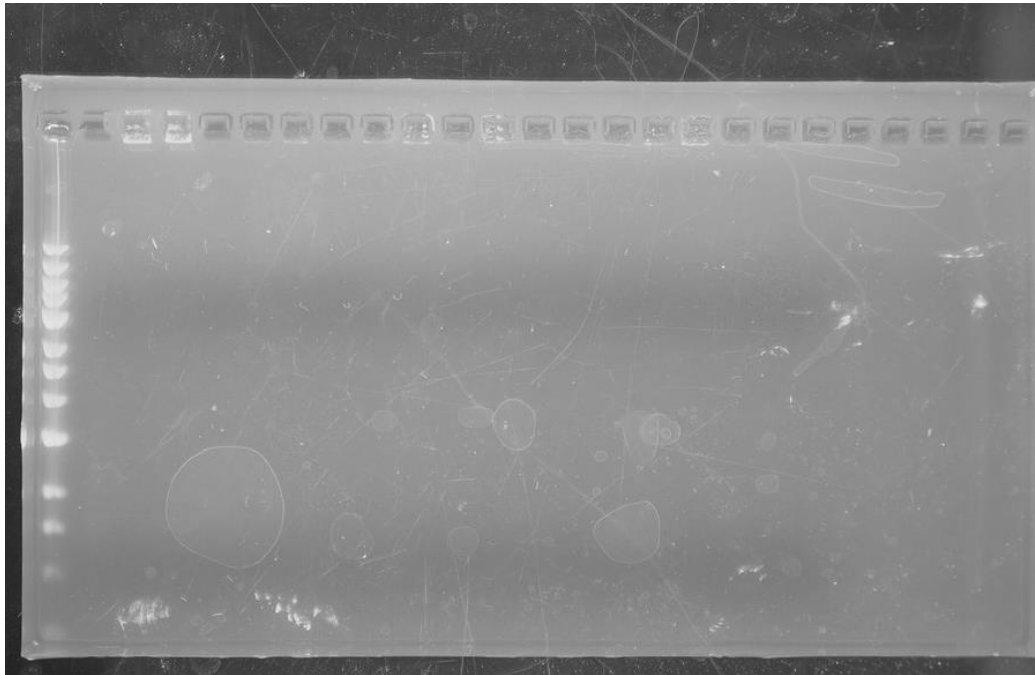
1. Check Px16, Px18

	DNA(3µl)	Agel (5unit/µl)	10Xbuffer1.1	ddH <sub>2</sub> O	Total V	time
Px16-1	2.5µl	0.40µl	2.00 µl	11.60 µl	15.00 µl	6hr
Px16-2	2.5µl					
Px18-1	2.0µl					
Px18-2	2.0µl					

		EcoRI (20unit/ µl)	10Xbuffer EcoR I	ddH <sub>2</sub> O	Total V	time
Px16-1	DNA(15µl )	0.25µl	2.0 µl	2.0 µl	20.0 µl	4hr
Px16-2						
Px18-1						
Px18-2						



## gel electrophoresis



Well 2 : miniprep cut px16

Well 3 : miniprep uncut px16

Well 4 : miniprep cut px18

Well 5 : miniprep uncut px18

Well 6 : traditon cut px18

Well 7 : traditon cut px18

Well 1 :

marker

2.

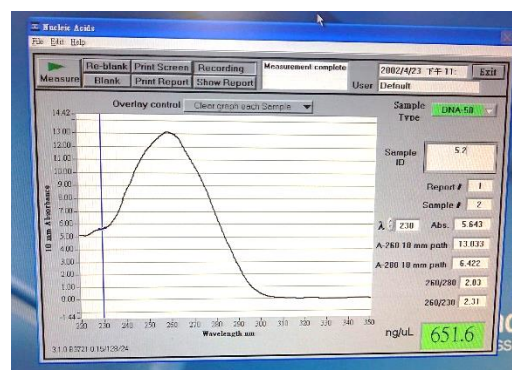
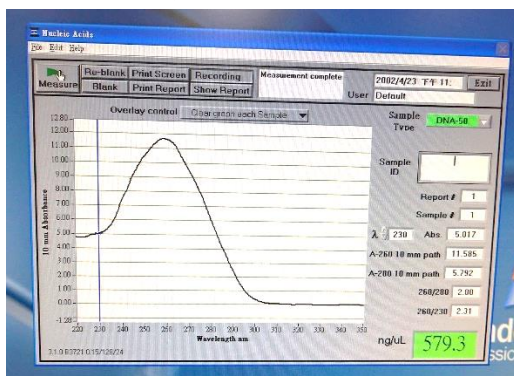
### ◆Miniprep Lac1 → store in -20°C

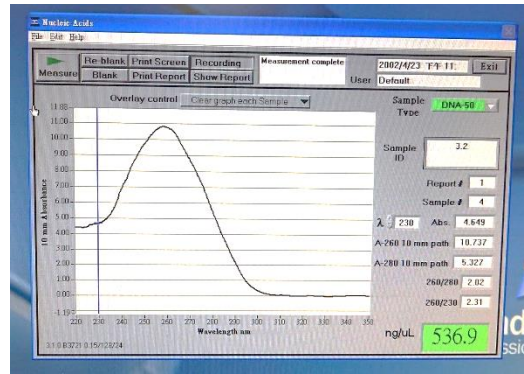
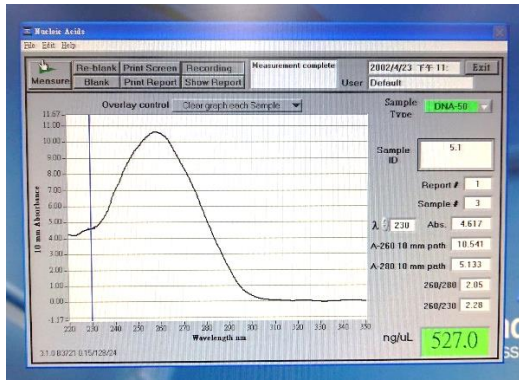
9/15

### ◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	SOUP	Time point	o

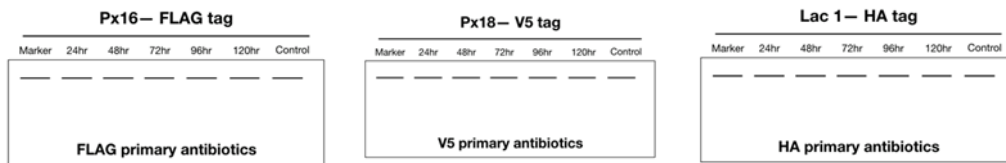
### ◆Nanodrop miniprep Lac



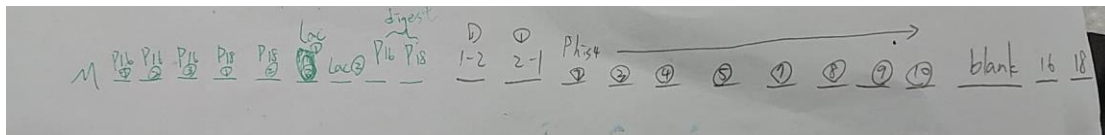


9/16

◆ Western blot: Antibody staining → Chemiluminescence detection



◆ Gel electrophoresis



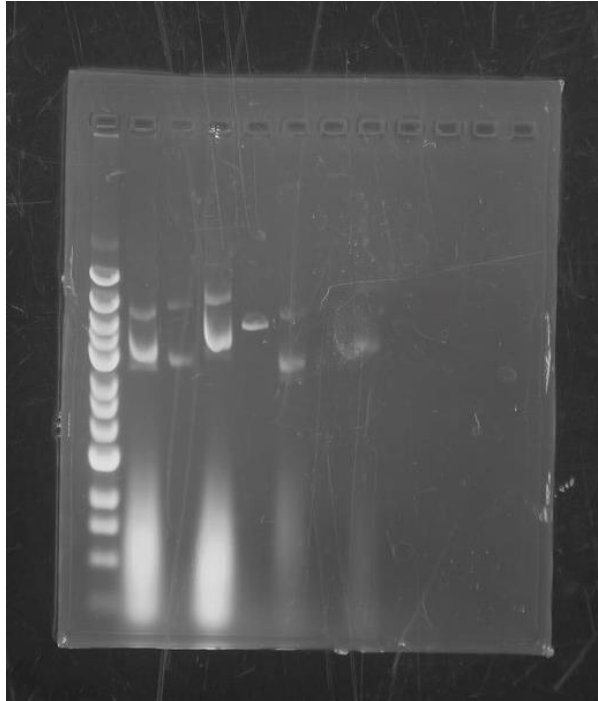
9/17

◆ Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
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1:1000	1:2000	1:5000	PELLET	Time point	o
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#### ◆Gel electrophoresis



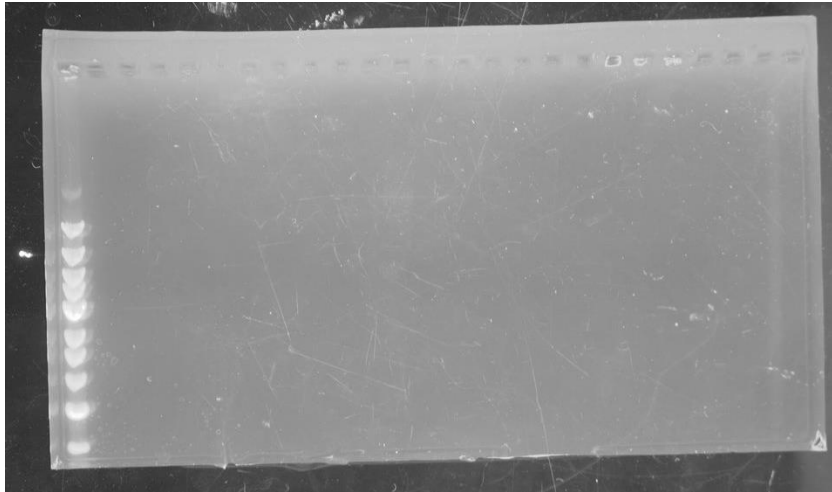
Well 1 : marker
Well 2 : uncut px16
Well 3 : cut px16
Well 4 : uncut px18
Well 5 : cut px18
Well 6 : uncut px16
Well 7 : cut px16
Well 8 : uncut px18
Well 9 : cut px18

#### ◆PCR

Component	Final(250µl)
10X Standard <i>Taq</i> Reaction Buffer	25µl
10 mM dNTPs	5µl
0.2 µM Primer suffix	5µl
0.2 µM Primer prefix	5µl
Template DNA	pGAPZ A_HIS4_Px16 pGAPZ A_HIS4_Px18 pGAPZ A_HIS4_Lac1
5 units/µl <i>Taq</i> DNA Polymerase	3.8µl
Nuclease-free water	207.5µl

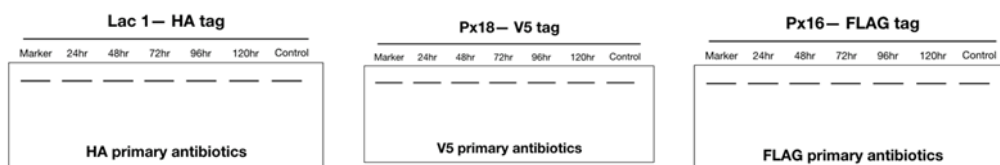
**Distribute to 25 eppendorf tubes, each of the tubes contains 10µl**

STEP	TEMP	TIME
Initial Denaturation	95°C	30 seconds
30 Cycles	95°C	30 seconds
	45°C	30 seconds
	72°C	2.5minutes
Final Extension	72°C	10 minutes
Hold	4°C	



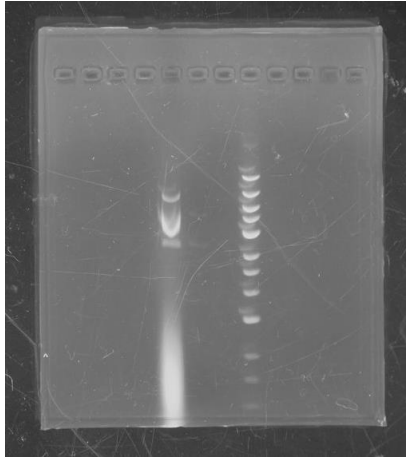
9/18

◆Western blot: Antibody staining → Chemiluminescence detection



Cut to linear for yeast transform

	DNA(10µl)	Sal I (100unit/µl)	10Xbuffer3.1	ddH <sub>2</sub> O	Total V	time
Px18	2.0µl	0.20µl	2.00 µl	16.00 µl	20.00 µl	6hr



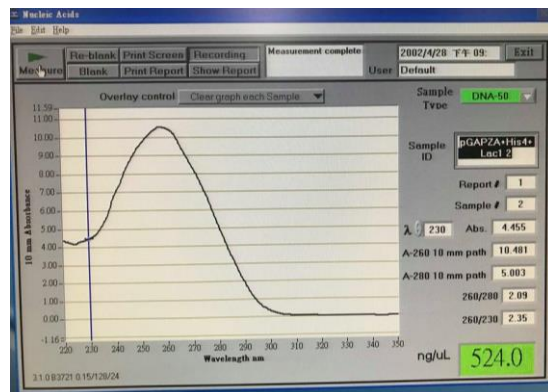
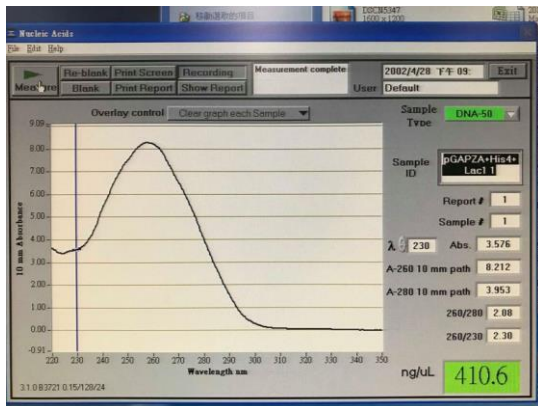
2. Miniprep Lac1 → store in -20°C

9/19

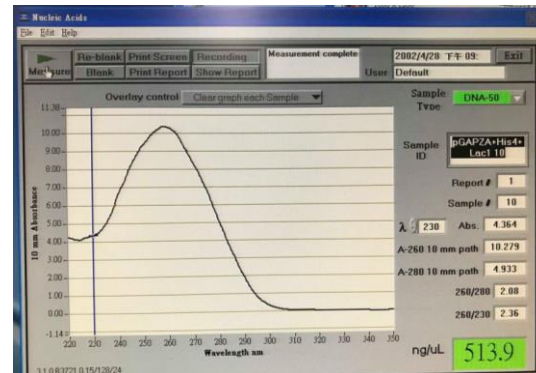
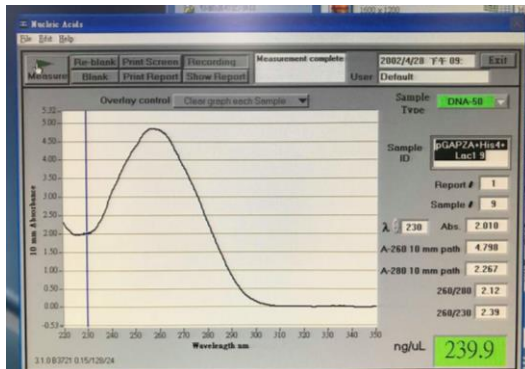
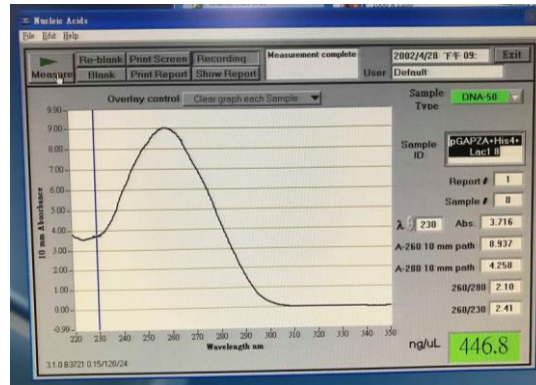
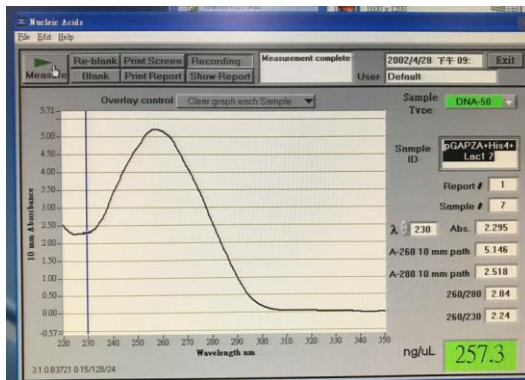
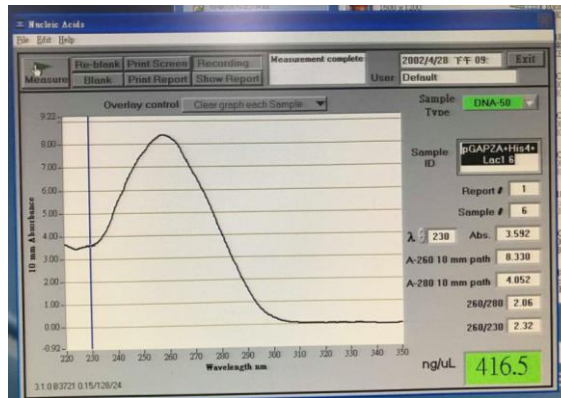
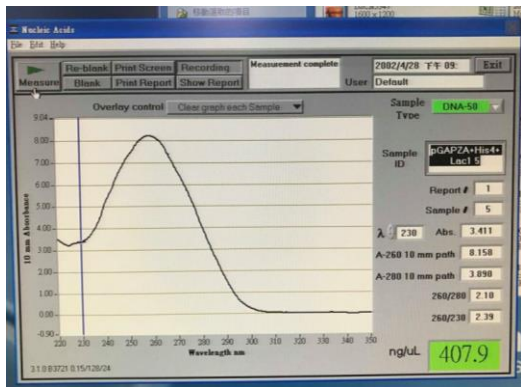
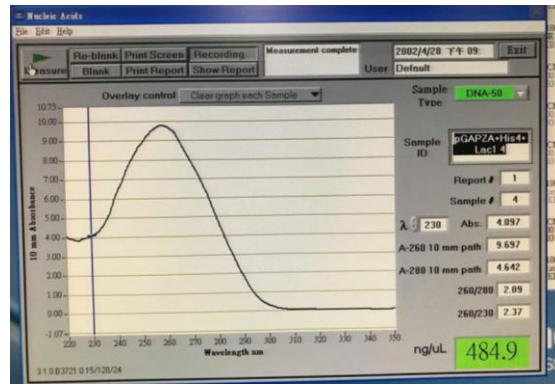
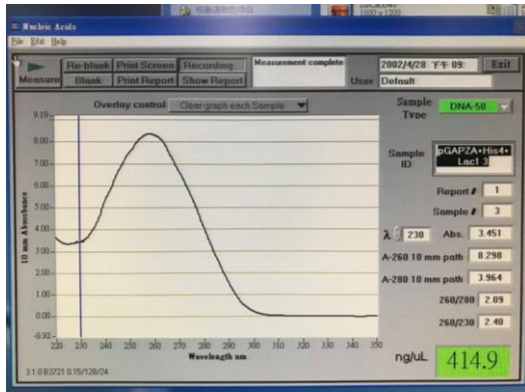
◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	PELLET	Time point	o

◆Nanodrop : Miniprep Lac1



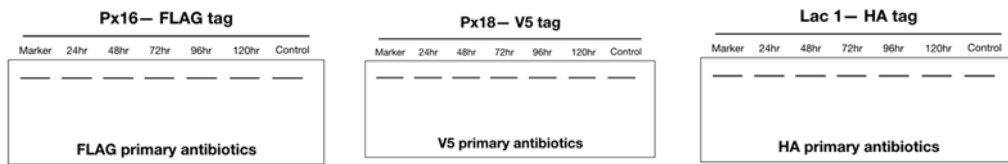




## 2.Minimal culturing of lacI

9/20

◆Western blot: Antibody staining → Chemiluminescence detection



1.Tradition miniprep Lac1 Day1

2.Prepare SD plate

9/21

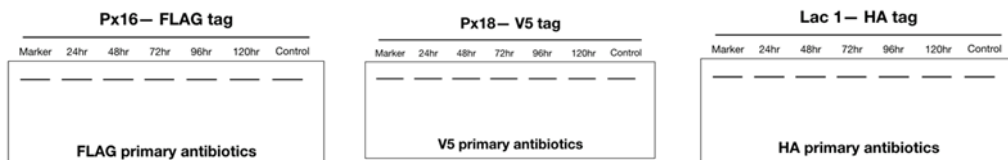
◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	PELLET	Time point	o

◆Tradition miniprep Lac1 Day1

9/22

◆Western blot: Antibody staining → Chemiluminescence detection



◆1.Prepare yeast

Pick up yeast(SD1168) from a colony + 50 ml YPD in erlenmeyer flask,  
Breeding in incubator in 30°C for days 2.5

2.Tradition miniprep Lac1 Day2

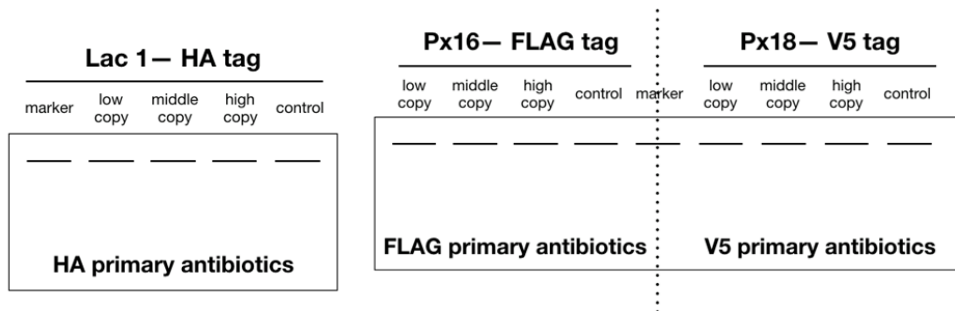
9/23

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

6x his	SOUP/PELLET	LOAD	ZEOCIN
1:20000	SOUP	LMH	x

9/24

◆Western blot: Antibody staining → Chemiluminescence detection



- ◆1.Minimal culturing of lac1
- 2.Store yeast in 4°c

9/25

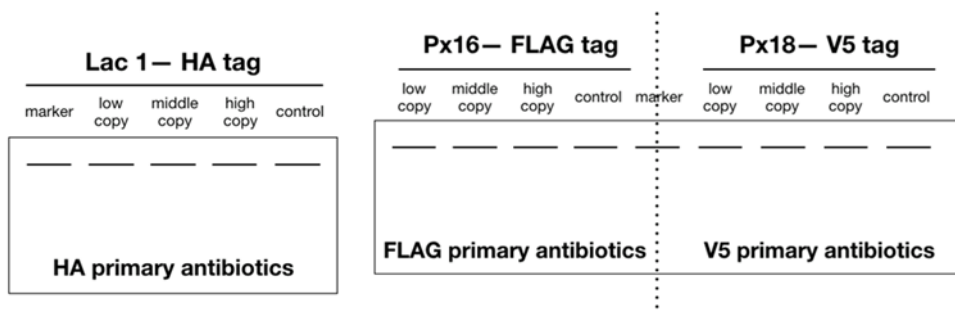
◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

6x his	SOUP/PELLET	LOAD	ZEOCIN
1:10000	SOUP	LMH	X

◆Tradition miniprep Lac1 Day1

9/26

◆Western blot: Antibody staining → Chemiluminescence detection



◆1.Tradition miniprep Lac1 Day2

- 2.Transform px16&px18 into yeast(SD1168)



9/27

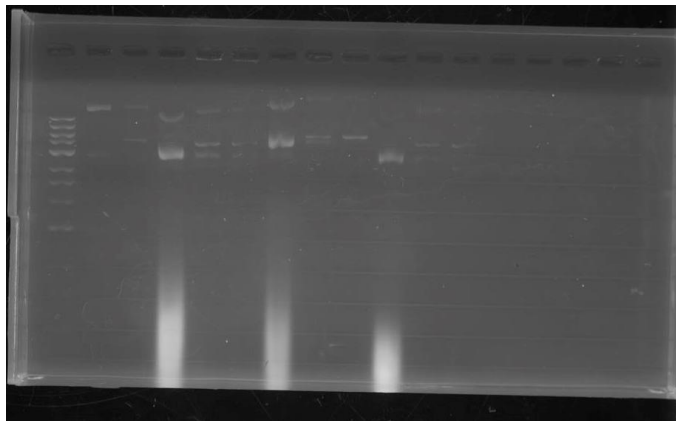
◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:5000	1:5000	1:5000	SOUP	LMH	X

◆Cut to linear for yeast transform

	DNA(10µl)	Sal I (100unit/µl)	10Xbuffer3.1	ddH <sub>2</sub> O	Total V	time
Px18	2.0µl	0.20µl	2.00 µl	16.00 µl	20.00 µl	6hr
Px16						

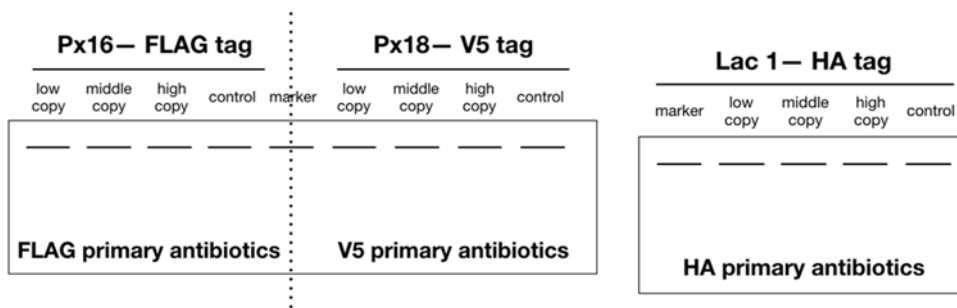
◆Gel electrophoresis



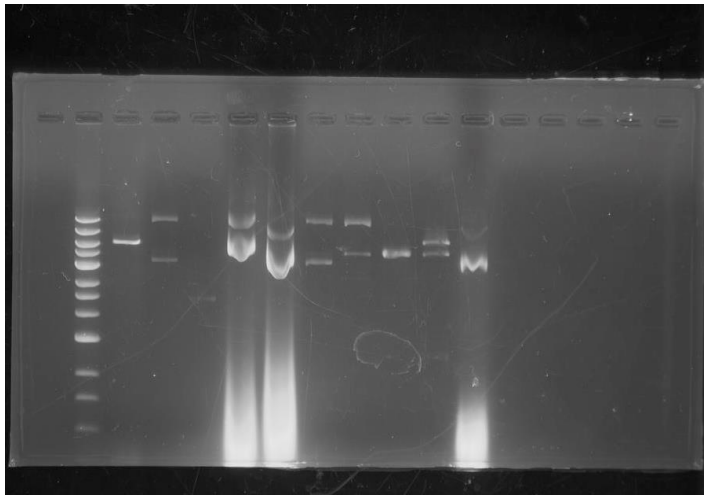
Well 1 :marker  
 Well 2 :pGAPZ A\_his4 uncut  
 Well 3 : pGAPZ A\_his4 cut  
 Well 4 : px16\_his4 uncut  
 Well 5 : px16\_his4 cut 1  
 Well 6 : px16\_his4 cut 2  
 Well 7 : px18\_his4 uncut  
 Well 8 : px18\_his4 cut 1  
 Well 9 : px18\_his4 cut 2  
 Well 10 : lac\_his4 uncut  
 Well 11 : lac\_his4 cut 1  
 Well 12 : lac\_his4 cut 2

9/28

◆Western blot: Antibody staining → Chemiluminescence detection



1. Transform px16&px18 into yeast(SD1168)
2. Gel electrophoresis



Well 1.2 :marker  
 Well 3 : px18\_his4 cut (sal I)  
 Well 4 : pGAPZ A\_his4 2-2  
 Well 5 : ?  
 Well 6 : px18\_his4 tradition uncut  
 Well 7 : px16\_his4 tradition uncut  
 Well 8 : pGAPZ A\_his4 1-2  
 Well 9 : 1-2  
 Well 10.11.12: ?

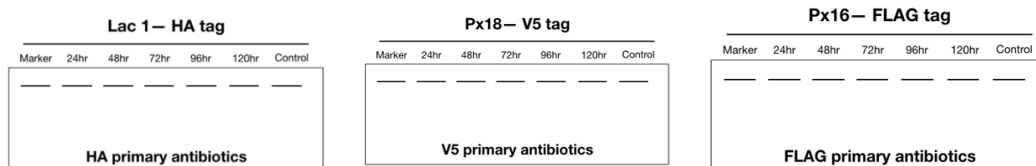
9/29

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:2000	1:2000	1:2000	SOUP	TIME POINT	X

9/30

◆Western blot: Antibody staining → Chemiluminescence detection

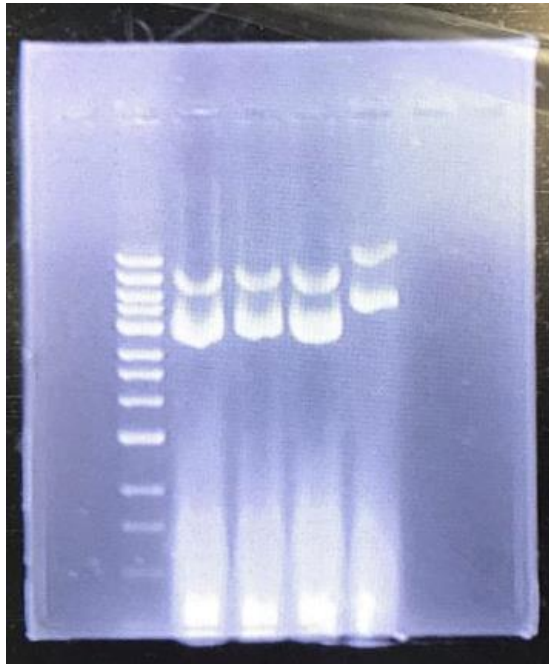


10/1

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:1000	1:1000	PELLET	TIME POINT	X

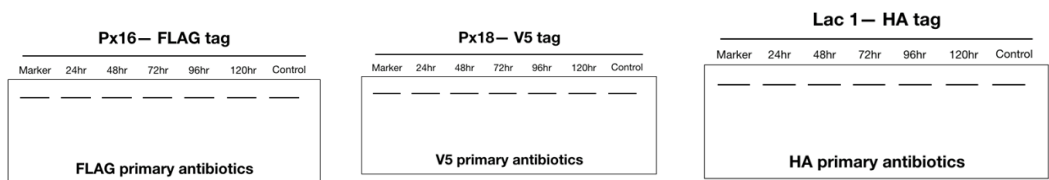
- ◆ 1. Minimal culturing of lac1
- 2. Gel electrophoresis



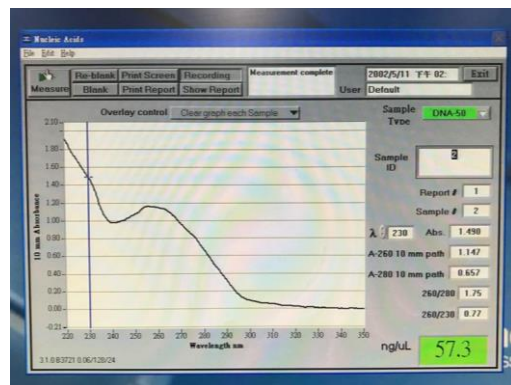
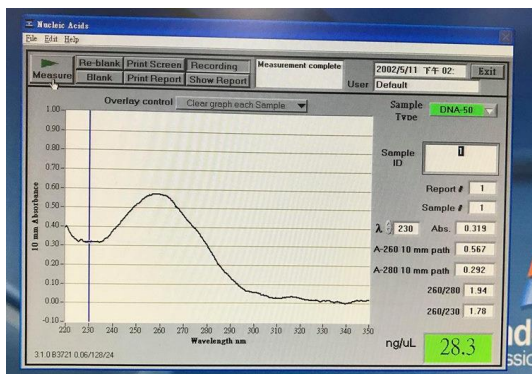
Well 1 : marker  
 Well 2.3.4 : px16\_his4 cut (sal I)

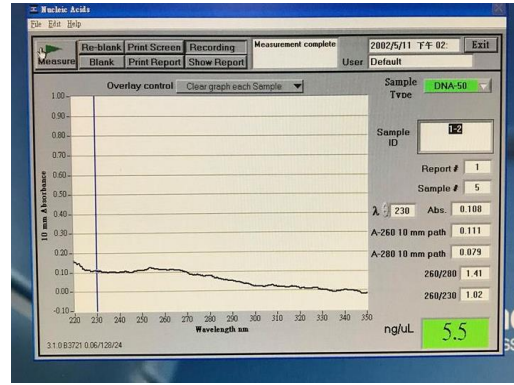
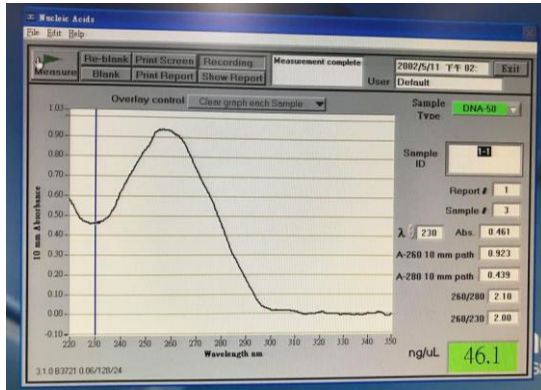
10/2

- ◆ Western blot: Antibody staining → Chemiluminescence detection



- ◆ 1. Nanodrop lac1(miniprep)





>> vector is out of the gene sequence ☹️

2. Transform px18 into yeast(SD1168)
3. Former SD plates are contaminated
4. Tradition DAY1 (Lac x1, Px16 x1)

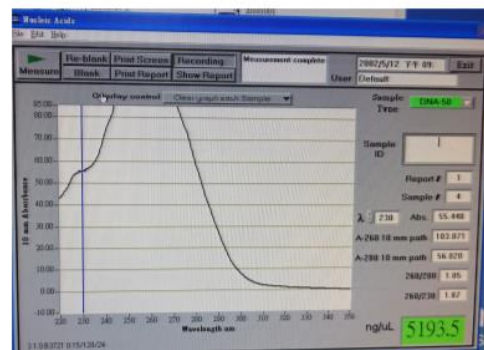
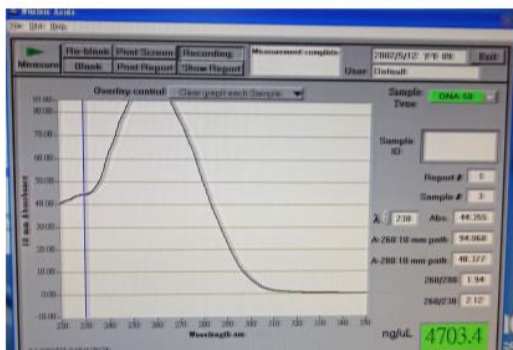
10/3

◆Western blot: Sample preparation→ SDS-PAGE→ Transfer→ Blocking→ Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:5000	1:5000	SOUP	TIME POINT	O

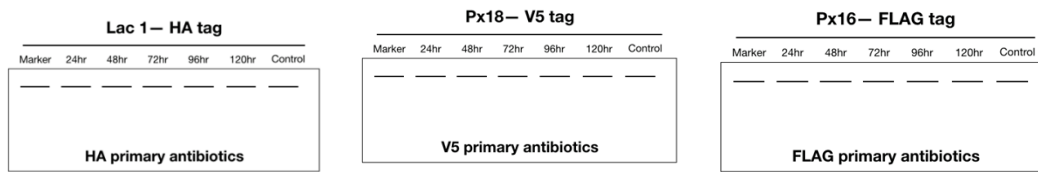
◆1.Safety part x3 (30°C 、 50°C 、 70°C)

- 2.Tradition DAY2 (Lac x1, Px16 x1)
3. Colony PCR (Lac x6, Px16 x6)



10/4

◆Western blot: Antibody staining → Chemiluminescence detection



◆Prepare new SD plate( the former are contaminated)

Yeast Nitrogen Base 6.7g + 100 ml H<sub>2</sub>O

Agar 20g + 750 ml H<sub>2</sub>O

50% Glucose 40 ml

Amino acids without Histidine 100 ml

10/5

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	SOUP	TIME POINT	O

1. Gel electrophoresis (Tradition miniprep Lac x3 & PCR: Lac x6, Px16 x6)

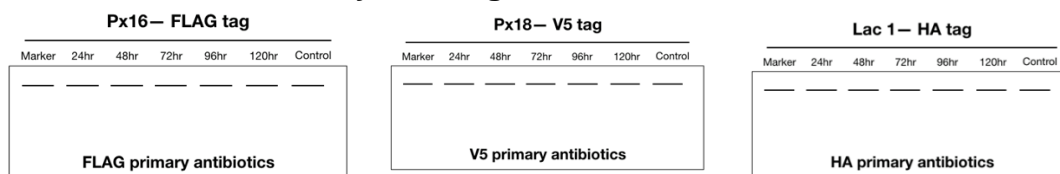




### 3. Transform px16&px18&pGAPZ A\_his4 into yeast(SD1168)

10/6

#### ◆Western blot: Antibody staining → Chemiluminescence detection



10/7

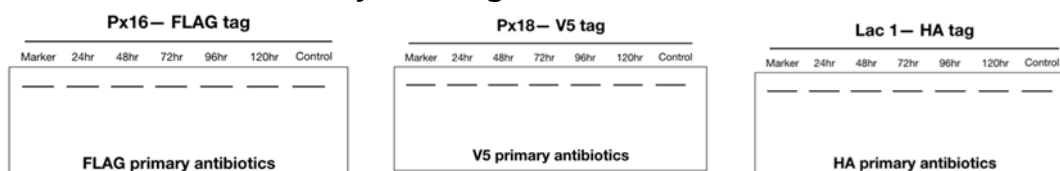
#### ◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	SOUP	TIME POINT	O

#### ◆Minimal culturing of px16

10/8

#### ◆Western blot: Antibody staining → Chemiluminescence detection



#### ◆1.PCR Lac1 >>fail

#### 2. Tradition Miniprep px16 day1

10/9

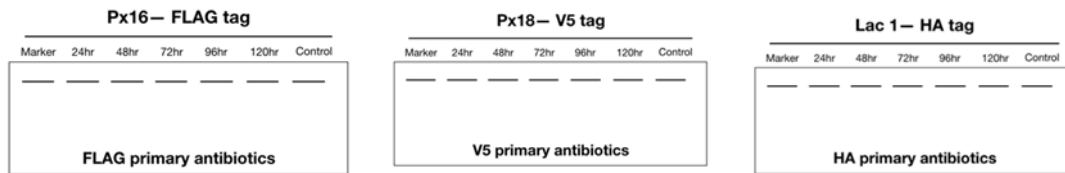
#### ◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	PELLET	TIME POINT	O

1. Miniprep px16
2. Tradition Miniprep px16 day2

10/10

◆Western blot: Antibody staining → Chemiluminescence detection



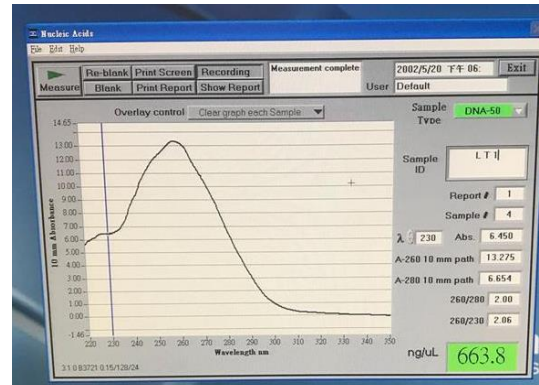
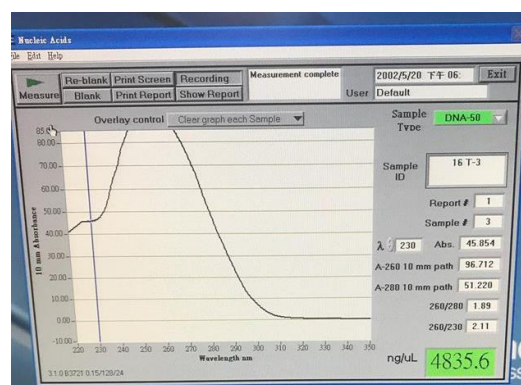
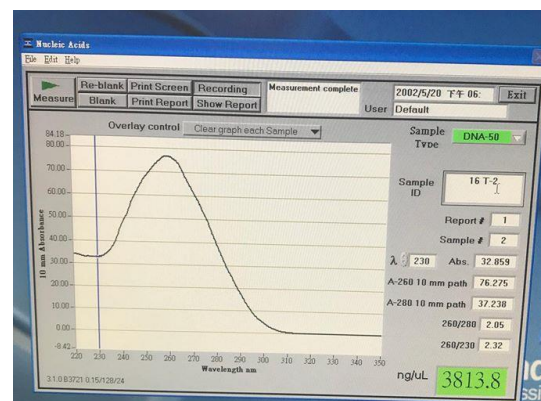
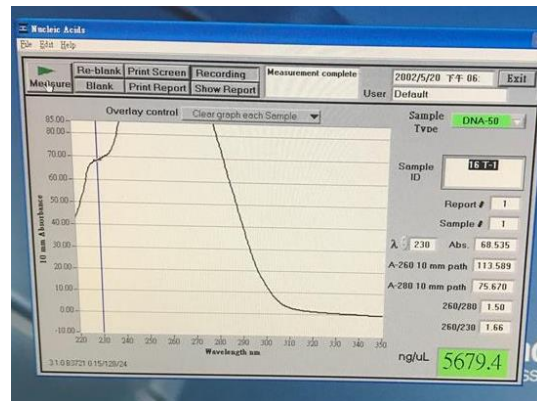
◆Tradition miniprep Lac1

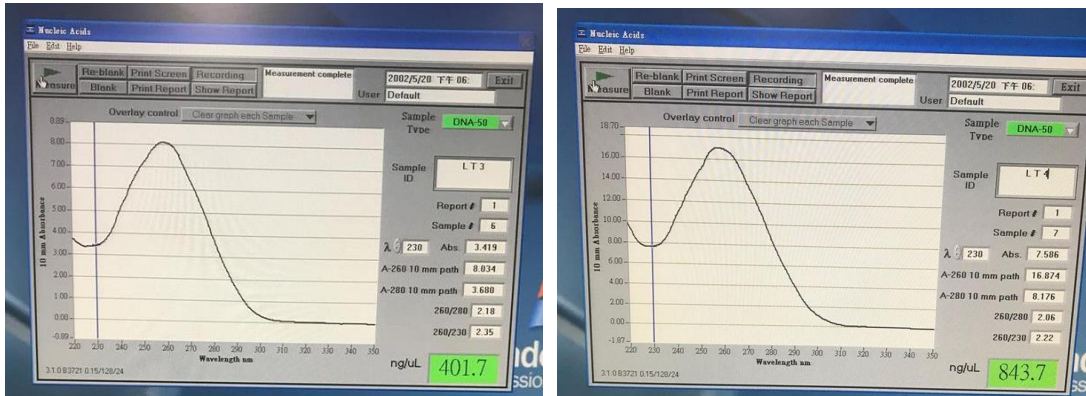
10/11

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	PELLET	TIME POINT	O

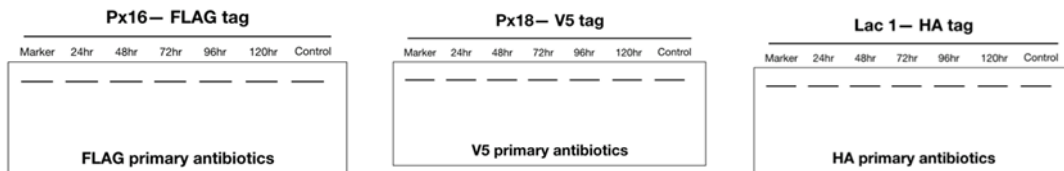
◆Nanodrop



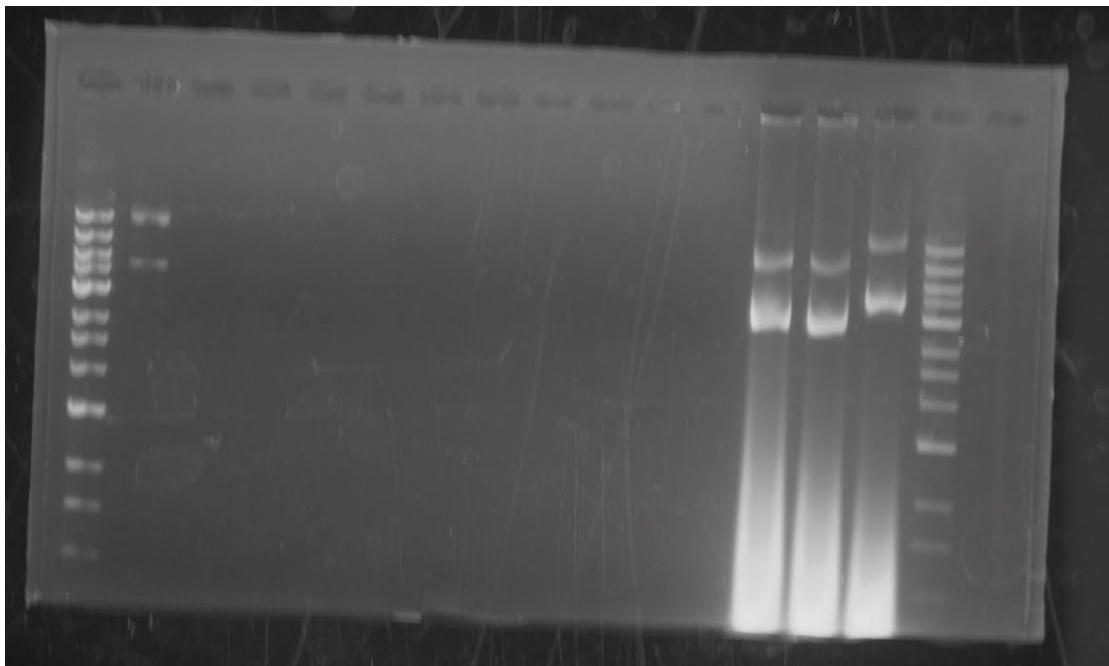


10/12

◆ Western blot: Antibody staining → Chemiluminescence detection



◆ Gel electrophoresis



marker/16-2 mini/ 16-3 cut/ 16-2cut/16-2cut/16-3 mini/16-3cut/lac1 mini/ lac1 cut/lac2 mini/lac2cut/lac2cut/lac3 mini/lac3cut/lac3cut/lac4 mini/lac4 cut

>> forget to dilute.

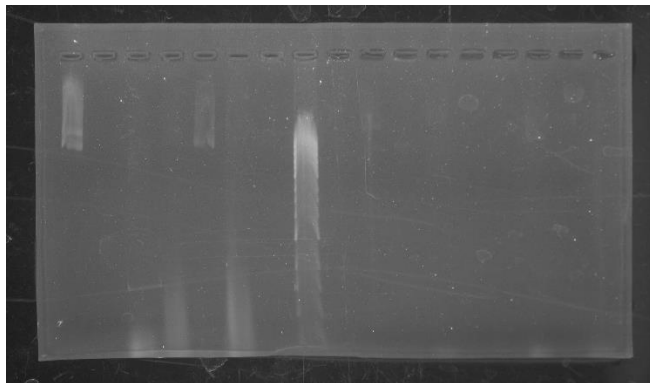
10/13



◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

FLAG	HA	V5	SOUP/PELLET	LOAD	ZEOCIN
1:1000	1:2000	1:5000	PELLET	TIME POINT	O

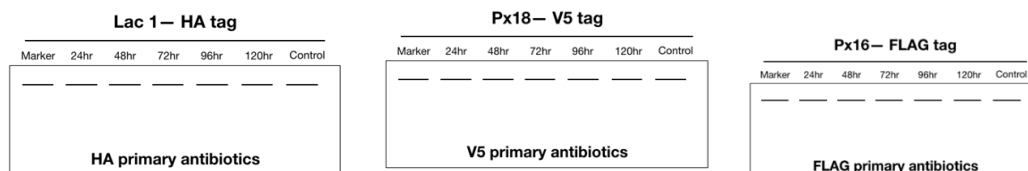
◆Gel electrophoresis



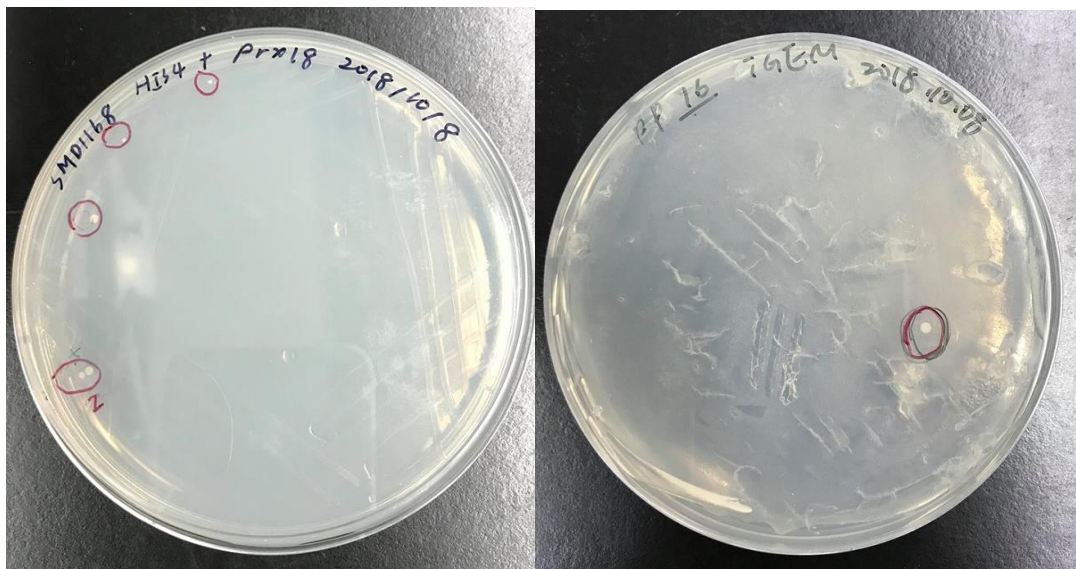
16-2 mini/ 16-3 cut/ 16-2cut/16-2cut/16-3 mini/16-3cut/lac1 mini/marker/lac1 cut/lac2 mini/lac2cut/lac2cut/lac3 mini/lac3cut/lac3cut/lac4 mini/lac4 cut

10/14

◆Western blot: Antibody staining → Chemiluminescence detection



◆PLATE

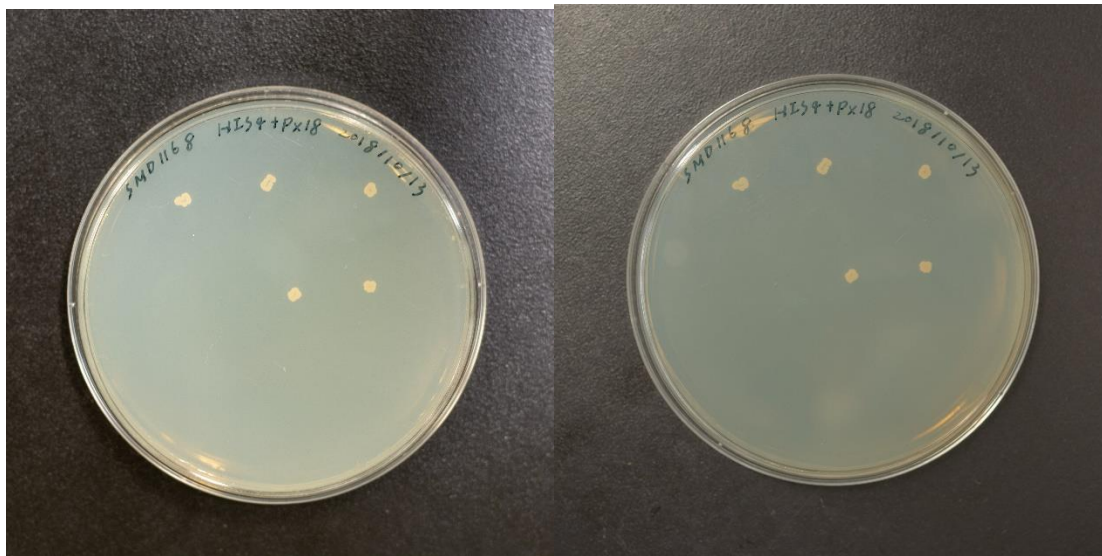


10/15

◆Western blot: Sample preparation → SDS-PAGE → Transfer → Blocking → Antibody staining

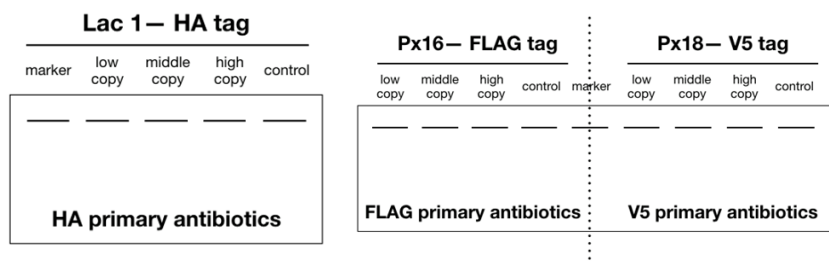
6x his	SOUP/PELLET	LOAD	ZEOCIN
1:20000	SOUP	LMH	X

◆Stick pellet to culture (SD)

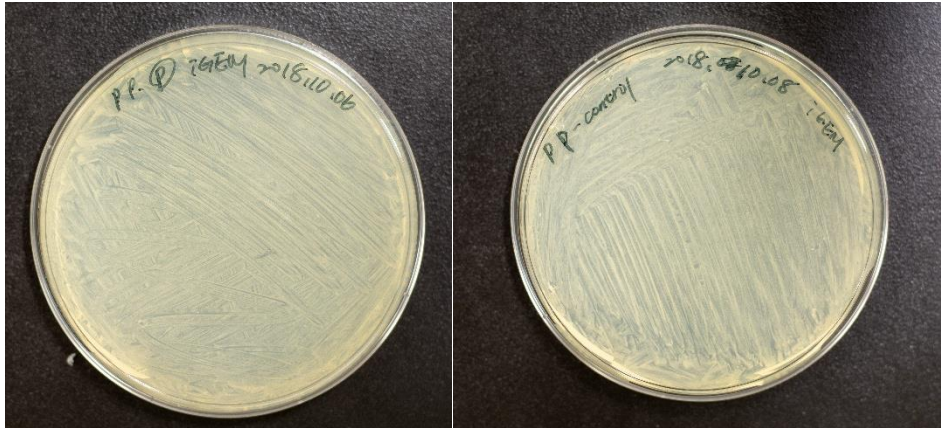


10/16

◆Western blot: Antibody staining → Chemiluminescence detection



## Control



## PLATE

