

10ng/ul       $\frac{1000\text{ng}}{100\text{ul}} = \text{leg DNA}$

Digest      31/7/2017

- 150 ml solution  
in microcentrifuge tube + aliquot into 3 PCR tubes.
- running on a gel

Gel 1/8/2017

1 LAA
1 Ladder
1 Das
1 ladder
1 Ø
1 LAA
1 Ladder
1 Das
1 <del>#</del> Ladder (1μg)
1 <del>ladder</del> Ø

15 μl of DNA, 3 loading dyn

1<sup>st</sup> time  
- no bands

2<sup>nd</sup> time  
- purple spots  
- no bands

order  
more  
TSPurple

purple bands are just  
degr. The gel had  
no DNA.

with  
standby

→ ask to test other tubes

## Transformation 2/6/2017

pλR lacI GFP LAA AK3 on kan plate.

↳ colonies grew, no fluorescence

PLG1 DAS CIPXP CI 1:100 on plain plate

↳ colonies grew, no fluorescence

PLG1 CIPXP CI 1:100 on plain plate

↳ colonies grew, no fluorescence

R11 CIPXP CI 1K3 on kan plate

↳ colonies grew, SUCCESSFUL

### Next steps:

- miniprep R11 CIPXP CI 1K3

- transform pλR lacI, GFP DAS, and GFP LAA to miniprep and obtain more DNA

Transformation 2/19/17P<sub>2</sub>R LacI in 3T5

↳ colonies grew → fluorescence (not expected)

↳ liquid cultures (x3)

GFP DAS in IC3

↳ colonies grew → no fluorescence (was expected: GFP)

↳ streak to singles

GFP LAA in IC3

↳ no growth

next: colony PCR + gel P<sub>2</sub>R LacI + GFP DASorder GFP LAAsend P<sub>2</sub>R LacI + GFP DAS

2/23/17

## Colony PCR of transformations

Completed in triplicates

→ colonies pulled from plates

→ P<sub>2</sub>R LacI in 3TS

→ GFP DAS in 1C3

## In Thermocycler

① ② ③ ④ ⑤ ← P<sub>2</sub>R LacI 3TS cPCR 2/22  
JAL

⑥ ⑦ ⑧ ⑨ ⑩ ← GFP DAS 1C3 cPCR 2/22  
JAL

Next: run a gel

into gel      S → 10 μL of DNA  
Z → 2 μL of Loading Dye

10 | 1      10 μL of 2 log ladder into ~~loading~~ gel

9 | P<sub>2</sub>R LacI 1

8 | P<sub>2</sub>R LacI 2 → ~~not a lot~~

7 | P<sub>2</sub>R LacI 3

6 | 2 Log Ladder

5 | GFP DAS 1 → less

4 | GFP DAS 2 → less

3 | GFP DAS 3

# Notes - From Google Call 2/24 3:45

- HSV or HSL values
- Pellet from liquid culture
- Make sure to account for # of cells + time of cells +  
multiple measures of same sample
- What would you expect graphs to look like?

Bar Graph

$\rho$ LAC - synthetic leakiness has been eliminated

(T7 promoter) interesting but not on topic

- Varying Cpx Cpx amino acid - Probably NOT Good idea - difficult
- Varying tags to other known working tags
  - Use the sequences in Registry
  - Feedback times

Slide 10 - Don't mutate tags

K10 strain Question - Not essential to address

#9 = Absolutely

\*8 - Yes!! + also background motivation or

3/23/2017 Transformation

\* used NEB protocol  
↳ includes heat block

Parts:

- ① RII C1pXP 3TS  
3/23 transf
- ② RII C1pXP 3TS  
33 C1pP 3TS  
3/23 transf
- ③ 33 C1pX AK3  
33 C1pP AK3  
3/23 transf
- ④ p2R 33 LacI 3TS  
3/23 transf
- ⑤ B33 CI IC3  
3/23 transf
- ⑥ p2R LacI RII RFP 3KS  
3/23 transf

{ 50uL cells  
10uL plasmid (diluted)  
green microcentrifuge tubes  
  
3 Tet plates  
2 Kan plates  
1 Cam plate  
  
plated 250uL onto each plate

Results:

- ① RII C1pXP 3TS - growth → liquid culture in tet resistance
- ② RII C1pXP 3TS - no growth → replating on tet  
33 C1pP
- ③ 33 C1pX AK3 - growth → liquid culture in kan resistance  
33 C1pP
- ④ p2R 33 LacI 3TS - no growth → replating on tet
- ⑤ B33 CI IC3 - growth → liquid culture in cam resistance
- ⑥ p2R LacI RII RFP 3KS - growth → liquid culture in kan resistance

3/29/17 Transformation \*used NEB protocol\*  
↳ includes heat block

Parts/Results:

- ① P<sub>E</sub> LacI [3T5] - growth on plain  
- no growth on tet

3/30/17 Colony PCR

Parts:

- P<sub>E</sub> LacI [3T5]  
① ② ③

Results:

4/14/17: colony PCR

8:00: made gel

made liquid cultures

- R0011 CIPXP (3T5)
- CI (IC3)

3rd gel: colony PCR

- R0011 CIPXP (3T5)
- CI (IC3)

JL & AF

4/14/17 = Part Lengths

PART	LENGTH
CIPP	624 bp
CIPX	1275 bp
R0011	222
CI	775 bp

CL and CP

4/17/17: miniprep of LC<sub>a</sub> from 4/14/17

· R0011 CIPXP (3T5) A<sub>260</sub>: 8.21, A<sub>260</sub>/A<sub>280</sub>: 1.35, 41.2 ng/μL

· CI (IC3) A<sub>260</sub>: 1.059, A<sub>260</sub>/A<sub>280</sub>: 1.42, 52.9 ng/μL

JL & AF

re-miniprep & for better results

4/18/19 miniprep (redo)

JL & AF

• R0011 CIPXP(3T5)  
• CI (IK3)

R0011 CIPXP

A<sub>260</sub>: .479 / .49  
A<sub>260</sub>/A<sub>280</sub>: 1.85 / 1.57  
24 ng/μL / 24.6

CI

A<sub>260</sub>: .422  
A<sub>260</sub>/A<sub>280</sub>: 1.81  
21.1 ng/μL

4/18/2017 Digest of RII CIPXP CI IK3

92.8 ng/μL

$$\frac{92.8 \text{ ng}}{1 \mu\text{L}} \left| \begin{array}{c} 1 \text{ ng} \\ 1000 \text{ ng} \end{array} \right. = \frac{0.0928 \text{ ng}}{1 \mu\text{L}} = \frac{1 \text{ ng}}{x \mu\text{L}} \quad x = 10.78 \mu\text{L}$$

2 μL × (10X Re-Mix)

1 μL P (standard enzyme)

10.8 μL DNA

6.2 μL dH<sub>2</sub>O

GEL

used 100 bp (only access) so could not tell

5/02/2017 Re-Digest

↳ RII CIPXP CI IK3

$$\hookrightarrow 92.8 \text{ ng/μL} \quad \frac{92.8 \text{ ng}}{1 \mu\text{L}} \left| \begin{array}{c} 1 \text{ ng} \\ 1000 \text{ ng} \end{array} \right. = \frac{0.0928 \text{ ng}}{1 \mu\text{L}} = \frac{1 \text{ ng}}{x \mu\text{L}} \quad x \approx 10.78 \mu\text{L}$$

2 μL × (10X Re-mix)

1 μL P (standard enzyme)

10.8 μL DNA

6.2 μL dH<sub>2</sub>O

5/07/2017 miniprep

R0011 C1PXP CI in 1K3

A<sub>260</sub> (10 nm): 0.687

A<sub>260</sub>/A<sub>280</sub>: 1.83

34.4 ng/ $\mu$ l

as this is the 2nd or 3rd  
miniprep with low concentration  
we think that the generation  
of R11 C1PXP CI is straining for  
the cells to generate, resulting  
in low cell growth

Colony PCR

12.5  $\mu$ l Q5  
1.25  $\mu$ l VF2  
1.25  $\mu$ l VR  
6  $\mu$ l DNA  
4  $\mu$ l H<sub>2</sub>O

using junior's in  
a 1K3 also

10ng/ul       $\frac{1000\text{ng}}{100\text{ul}} = \text{leg DNA}$

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in microcentrifuge tube + aliquot into 3 PCR tubes.
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15 μl of DNA, 3 loading dyn

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more  
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purple bands are just  
degr. The gel had  
no DNA.

with  
standby

→ ask to test other tubes

Ligation Calculation 1/8/2017

vect:

Part 1: TS Purple

(waiting on p2r LacI backbone)

Backbone Digest 1/8/2017

1C3

1A3

1T3

1K3

(all with EP)

→ should we  
run on  
gel?  
ask standby

can't  
→ truth

Transformation 8/1/2017

miniprep media  
w/ kanamycin

P2R LacT<sup>+</sup>K<sup>3</sup> (plating on all 4 resistant plates) ~  
(\*) miniprep 3/22

C1pX P Carb Miniprep AK<sup>3</sup>

miniprep 9/28

out of Kan

C1pX D CI 3K5 → cell growth on plain

miniprep 4/28

streak onto Kan

→ all grew on Kan

P2R LacI 1K<sup>3</sup>

plain - ✓ growth → streak onto kan

can - ✗ no growth

carb - ✗ no growth

tet - ✗ no growth

kan - ✓ growth → from streaks

## Miniprep

P2R LacI (1K<sup>3</sup>)

liquid culture made 8/4/2017

failed

August 4<sup>th</sup>

Gel Electro.

- LAA
  - Ladder
  - DAS
  - ladder
  - φ
- 
- LAA
  - Ladder
  - DAS
  - ladder 2log
  - φ

results: ladders were clear and visible, however no DNA ~~is in~~ in the wells w/ our constructs.  
L. reordered parts <sup>→ 1D purple</sup> 8/4/17  
- JLG EG

8/8/17 - replacing PCR lysis with water place  
JL and LH transformed from DNA in freezer  
water place - transformed  
R0011 ApXPC1 -  
tspinylers - ordered  
estimated shipping 8/11  
estimated arrival 8/17

reminder: you are all wonderful,  
lovely people. we got this!

Wednesday, August 9, 2017

\* Result: Growth of weak pLAC AK3 on Kan plate from transformation on 8/8/2017

\* Lab: Liquid cultures (3 5mL LC tubes)  $\Rightarrow$  inoculation

Thursday, August 10, 2017

\* Liquid cultures  $\Rightarrow$  did not grow (used AF Kan LB 10-14-14)

$\hookrightarrow$  new liquid cultures: 3 new Kan LB, 1 plain LB, 1 from 8/9

\* Digest of tsPurple, tsPurple LAA, tsPurple DAS after hydration

$\hookrightarrow$  Protocol for Hydration: Pipette 10 $\mu$ L milliQ H<sub>2</sub>O to make 100 ng/ $\mu$ L

$\rightarrow$  15.5  $\mu$ L H<sub>2</sub>O + 2.5  $\mu$ L DNA + 2 mL E + 1 mL P

Friday, August 11, 2017

\* Ligate tsPurple into TC3 (in box  $\Rightarrow$  ligase box M13)

8/11 miniprep (pLacI) JL, LH, EG

1:	18.1 ng/ $\mu$ L	A <sub>260</sub> : 0.362	A <sub>260</sub> /A <sub>280</sub> : 1.98
2:	11.7 ng/ $\mu$ L	A <sub>260</sub> : 0.235	A <sub>260</sub> /A <sub>280</sub> : 1.99
3:	19.5 ng/ $\mu$ L	A <sub>260</sub> : 0.391	A <sub>260</sub> /A <sub>280</sub> : 1.57
4:	11.5 ng/ $\mu$ L	A <sub>260</sub> : 0.231	A <sub>260</sub> /A <sub>280</sub> : 2.07

Next miniprep: open new kit

R0011 ClpXP CI backbone: 1K3 (8/9/2017)

- ↳ miniprep → stored as
- ↳ digest with  $\lambda + P_3'$  ask standby
- ↳ 21 - 25 of august

Friday, August 10

Transformation of TS Purple ( $\emptyset$ , DAS, LAA)

Tuesday, August 15

Ligation of TS Purple and AK3 ( $\emptyset$ , DAS, LAA)  
plated on K (all 3 grew + control  $\rightarrow$  10 beta, grew)

Wednesday, August 16

Made Amp & Kan LB

Liquid culture of TS Purple ( $\emptyset$ , DAS, LAA) AK3  
Liquid culture of pAR LacI Tet LB

Friday, August 18 (miniprep, AF)

nanodrop (5L)

$\emptyset \Rightarrow 0.6 \text{ ng}/\mu\text{L}, A_{260}/A_{280} = 1.31$

DAS  $\Rightarrow 1.3 \text{ ng}/\mu\text{L}, A_{260}/A_{280} = 1.47$

LAA  $\Rightarrow 1.1 \text{ ng}/\mu\text{L}, A_{260}/A_{280} = 1.37$

1.6 ng/ $\mu\text{L}, A_{260}/A_{280} = 1.61$

2.9 ng/ $\mu\text{L}, A_{260}/A_{280} = 1.56$

1.1 ng/ $\mu\text{L}, A_{260}/A_{280} = 1.26$

p $\lambda$ R LacI 3TS from 3/22 : 2.3 ng/ $\mu$ L A<sub>260</sub>/A<sub>280</sub>: 1.86

p $\lambda$ R LacI 1T3 from 3/22 : 7.0 ng/ $\mu$ L A<sub>260</sub>/A<sub>280</sub>: 1.77

p $\lambda$ R LacI 3TS from 3/22 : 4.0 ng/ $\mu$ L A<sub>260</sub>/A<sub>280</sub>: 1.71

p $\lambda$ RLacI 3TS from 8/17/17 : 4.3 ng/ $\mu$ L A<sub>260</sub>/A<sub>280</sub>: 2.00

p $\lambda$ RLacI 3TS from 8/17/17 : 16.8 mg/ $\mu$ L A<sub>260</sub>/A<sub>280</sub>: 1.54 (18.5 ng/ $\mu$ L original)

pODAI(Cl) $_2$ PCl from 5/15/17 41.2 mg/ $\mu$ L A<sub>260</sub>/A<sub>280</sub>: 1.81

8/22 remini<sup>ss</sup> from 8/16

trypsin 5.4

Dar 11.8

Lac 6.3

Wednesday 8/23

PCR:

- ① P<sub>ZR</sub> LacI 3TS (6.0 ng/μL)
- ② P<sub>ZR</sub> LacI 3TS (18.5 ng/μL)
- ③ R0011 C1pxP CT

Thursday 8/24



- 1 -
- 2 -
- 3 - P<sub>ZR</sub> LacI 3TS (6.0 ng/μL)
- 4 - P<sub>ZR</sub> LacI 3TS (18.5 ng/μL)
- 5 - 2'-Log-Ladder
- 6 - R0011 C1pxP CT

8/29/17

sent for sequencing:

AHA772 → P<sub>0011</sub>C<sub>1</sub>pXPC1 with VF<sub>2</sub> (5NL DNA, 0.8NL primer, 6.2NL millipore H<sub>2</sub>O)

ANV224 → P<sub>λRLacI</sub> 3TS1 with VF<sub>2</sub> (2.5NL DNA, 0.8NL VF<sub>2</sub> primer, 8.7NL H<sub>2</sub>O)

ANV226 → P<sub>λRLacI</sub> 3TS2 with VR (2.5NL DNA, 0.8NL VR primer, 8.7NL H<sub>2</sub>O)

Digest from IDT:

2.5 uL DNA

2 uL E or X

1 uL S or P

12.5 uL H<sub>2</sub>O

Ligation from IDT:

1.4 uL DNA

2 uL ligase buffer

1 uL ligase

14.4 uL H<sub>2</sub>O

1 uL IC3

Hydrating constructs from IDT:

\* 10uL H<sub>2</sub>O for 100ng/uL concentration

\* Digest

\* Ligate

Send for sequencing:

800 ng DNA for sequencing

0.8 uL primer VF<sub>2</sub> & VR

bring to 12uL  
w/ H<sub>2</sub>O

# TSPurple (T<sub>3</sub>Purple DAS, & TSPurple LAA)

8/30/17

20 NL ~~Digest~~ starting off  
- 1.4 NL for ligation  
- 6 NL for gel  
12.6 NL left

Gel 8/30/2017

1.	-
2.	-
3.	-
4.	-
5.	+ T <sub>3</sub> Purple LAA
6.	2 Log ladder
7.	+ T <sub>3</sub> Purple
8.	+ T <sub>3</sub> Purple DAS
9.	-
10.	-

expected lengths: 788 bp <sup>DAS</sup>  
LAA  
750 bp Ø

results: lengths look  
good! successful  
digest

100 V, 60 min

## 8/31/2017 Ligation of T<sub>3</sub>Purple, T<sub>3</sub>Purple DAS, T<sub>3</sub>Purple LAA

Program: 002 on thermocycler

4.5 uL DNA      <sup>plate on</sup>  
1 uL ~~T<sub>3</sub>C~~ AT3 or AK3      <sup>both</sup>  
+ 5.5 uL sticky-end ligase      MM  
11.0 uL      ↳ 2X, refer to protocol

the same  
(<sup>+</sup>T<sub>3</sub>Purple or  
T<sub>3</sub>Purple DAS, LAA)

other tubes  
14.6 uL H<sub>2</sub>O  
1.4 uL DNA

stored in freezer w/ DNA

\* Ran out of IC3, used digested backbones from one year ago  
↳ Digest more backbone

## 9/1/2017 - Transformation of tsPurple, tsPurple DAS, tsPurple LAA

- 2μL DNA
- 50μL DH10 beta cells
- Heat shock for 30 sec @ 42°C, ice for 2 min
- plated on Tet & Kan plates

9/5/2017

Results from Transformation:

- ↳ Tet plates: no growth on control nor tsPurple (+ DT) plates
- ↳ Kan plates: growth on all four plates (control, tsPurple, tsPurple DAS, tsPurple LAA)

9/6/17 ::

Transformations:

P<sub>KR</sub> LacZ (18.5ng/L), R0011C<sub>1</sub> XPC1, GFP (from RNA mini-prime Box or Control), RFP

(Following NEB<sup>10</sup>, Beta Transformation Protocol)

9/6/17 - lunch period

- colony PCR

- control (C) labels C
- ts purple Ø Ø
- tspurple DAS (DAS)
- tspurple LAA (LAA)

PCR setup

C Ø DAS

LAA

- julia

DNA dilutions in ~~Slow microtote rack~~ (freezer)  
julia/emily/laurie's box

sequencing - 9/4/2017

RUOII (1PXP CI (63.2 ng/μL): VF<sub>2</sub> - AHA782  
VR - AHA774

9/8/17 gel of colony PCR's  
- trypsinase

- | 1. control
- | 2. tr purple S
- | 3. trpurple Sac
- | 4. tr purple Sac
- | 5. ladder (2-log)
- | 6. PCR Sac I
- | 7. RUOII CpxPCU

12 μL reactions

10 μL DNA

2 μL loading dye

4 μL 2-log ladder

expected lengths

S ~ 760

Kas ~ 790

Sac ~ 790

PCR Sac I ~ 1115

RUOII CpxPCU ~ 2000 2012

\* tubes in weigh boat in top left

RFP Transformation 9/18/17 NS

- RFP → 5uL, 10uL  $\exists$  P<sub>X</sub>R LacI R011RFP → DH10 Beta  
P<sub>c</sub>19 → 5uL  $\exists$  (+) control → DH10 Beta  
→ DH10 Beta (-) control

\* put in incubator for 1 hr at 2:28 p.m.

- × spread on Kan and plain plates
- P<sub>X</sub>R LacI R011RFP 5uL → ~~Kan~~ plain plate
  - P<sub>X</sub>R LacI R011RFP 10uL → Kan plate
  - P<sub>c</sub>19 5uL → plain plate
  - plain DH10 Beta → plain plate Carb plate  
↳ expect to see nothing b/c on a  
Carb plate

9/13/17 RFP Streak to Singles

- \* results from Friday → lawns on both RFP plates  
→ controls: colonies on Carb plate or DH10  
lawn / colonies of P<sub>c</sub>19  
- colonies (look visually different from  
lawn - colonies) most likely contamination
- P<sub>X</sub>R LacI R011RFP 5uL → plated on Kan / streak to singles
  - P<sub>X</sub>R LacI R011RFP 10uL → plated on Kan / streak to singles

9/13

- inoculated liquid cultures of p<sub>X</sub>R lacI and R0011Cp<sub>X</sub>P<sub>c</sub>  
need to retransform tryptone constructs

9/13 lunch  
transformation of tryptone G, DAS, CAA  $\exists$  puc19

↳ did not work

↓ control

9/14/17 - going back to 8/18 miniprep to  
transform, cPCR, and turbidite

ligation  
-2  
-2  
1/2 left

9/14/17 transformation <sup>JL</sup> ~~of trypanosomes from ST227~~

Transformations: tsPurple AK3, tsPurple DAS AK3, tsPurple LTA AK3

↳ plated 100uL on amp + kan plates (+ puc19 DNA control)

\* Note: heat shock occurred a little later than 30 min.

LH  
Transformations: R11C1pXP 3T5, B73 CJ 1C3

↳ plated 100uL on <sup>↑</sup>tet and <sup>↑</sup>cam plates (+ puc19 DNA control)

\* Note: heat shock occurred a little later than 30 min

\* Next Step: Colony PCR

9/15/17

transformation results. did not work  
↳ re-digesting from hydrated (trypanos)   
digested trypanser  
(transfer into purple microcentrifuge tubes)

9/15/17 - Gaurav B.

Miniprep of R001C1pXP C1 and PKRblad  
Results -

R001C1pXP C1 - A<sub>260</sub> (10mm): 3.859  
A<sub>260</sub>/A<sub>280</sub>: 1.86

Concentration: 193.0 ng/μL

PKRblad - A<sub>260</sub> (10mm): 0.429  
A<sub>260</sub>/A<sub>280</sub>: 1.91

Concentration: 21.5 ng/μL

9/15/17 Julia

3rd - ran gel of trypsin digests w/ ladder.

standy's ladder

ts purple o

tr purple eas

ladder

tr purple eas

SMI eas, not loading tape

expanded length

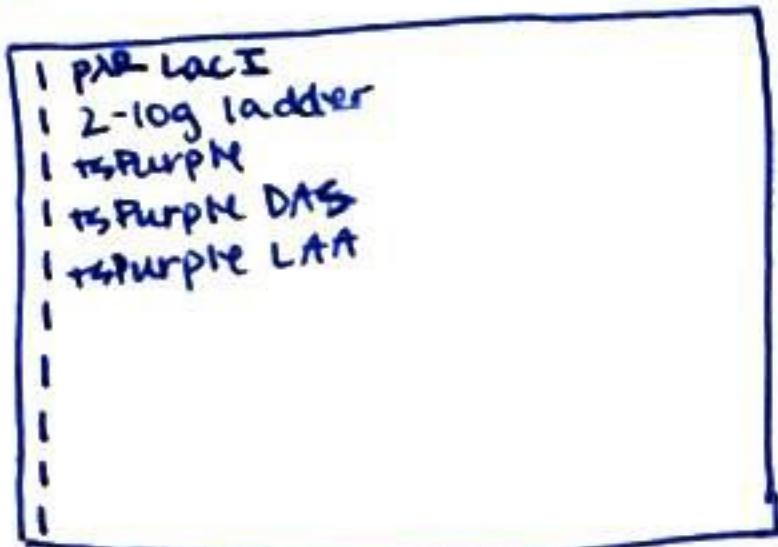
o - 750

dark - 780

9/16/2017

Digests → stored in pink PCR-tube box in junior drawer!

Sample: pAR lacI IC3 E2S } 25 μL rxns  
 + tsPurple / tsPurple DAS / tsPurple LAA X & P  
 Ctrl



5 μL DNA, 1 μL loading dye

4 μL 2-log ladder

RESULT: pAR lacI did not show up  
 tsPurples look good ✓  
 [ ] too low

Ligations → stored in pink PCR-tube box in junior drawer!

Parts	length of Insert	length of Vector	Total	CONC	μL	
IC3	1000	2200	3200	0.31	1	2 μL ligation buffer
pAR lacI	1115	2204	3324	0.30	3.1	+ 1 μL ligase
tsPurple	690	0	690	1.45	0.64	
tsPurple DAS	920	0	920	1.087	0.86	
tsPurple LAA	920	0	920	1.087	0.86	
→ + 12.26 μL mg H <sub>2</sub> O → 12.04 μL mg H <sub>2</sub> O						

\* Part length of tsPurple should be 759 bp, 788 bp w/ DAS & LAA \*

9/19

calculations were off → wrong tspurple part lengths  
 pCR conc. too low → re-inoculating from  
 9/17/17 GB pCR lacI transformation  
 lac: no colonies      0 colonies, purple (RFP: : )  
 das: colonies, white      puc19 - no growth ;)

inoculating D & C lysis to run digest gel diagnostics

10/2 Hydrated amilcp, amajLime, & CjBlue  
- stored in freezer (bag) in box that says Standy's

10/3/2017

<u>parts</u>	<u>1 of Insert</u>	<u>1 of vector</u>	<u>Total</u>	<u>-[DNA]</u>	<u>μL</u>	
IC3	1982	2070	4052	0.248	2	1 μL ligase
pBR Laci	1227	3189	4416	0.226	2.10	+ 2 μL lig buffer
~purple	788	0	788	1.325	0.38	12.42 μL dH <sub>2</sub> O
IC3	2015	2070	4085	0.245	2	1 μL ligase
pBR Laci	1227	3189	4416	0.226	2.16	+ 2 μL lig buffer
~purple DAS/LAA	788	0	788	1.269	0.38	2.46 μL dH <sub>2</sub> O

10/4/17

Transformation results (pmc19 over表达了 LacZ表达量)

\* control =

everything else = - cry -

10/4/17 (GFP)

Digoxin

pBR - 16.5 μL DNA in 25 μL reaction

RBS - 12.2 μL DNA in 50 μL reaction

16.5 DNA

2.5 water

5 EcoRI-HF

5 SphI-HF

5 mg/L

12.2 DNA

5.0 water

1.0 G-Taq

1.0 PstI

30.9 μL H<sub>2</sub>O

10/14/17 GB

ligation of p $\lambda$ R, ~~lacZ~~, and RBS  
 \* almost out of T4 ligase

Parts	L of Part	Lot Vector	Total	"kb	uL
IA3	60	2155	2355	0.443	1
p $\lambda$ R	49	2079	2129	0.152	2.94
RBS	11	2070	2081	0.481	2.76

1 uL IA3

1 uL Ligan

2.94 uL p $\lambda$ R Dig11.3 uL mgH<sub>2</sub>O

2.76 uL RBS Dig

2 uL Ligan Buffer

10/9/17 - GB

Restriction Digest

p $\lambda$ R + RBS	LacI
DNA 11.99 uL	4.40 uL
wtSmart 2.5 uL	2.5 uL
EcoRI-HF 0.5 uL	0.5 uL
SphI-HF 0.5 uL	0.5 uL
mgH <sub>2</sub> O 9.51 uL	7.1 uL

Program OD3

10/17/2017

Miniprep: 2 (60.9 ng/uL, A260/A280: 1.93)	#2
1 (61.8 ng/uL, A260/A280: 1.92)	#1
3 (106.8 ng/uL, A260/A280: 1.89)	#3

10/18/17

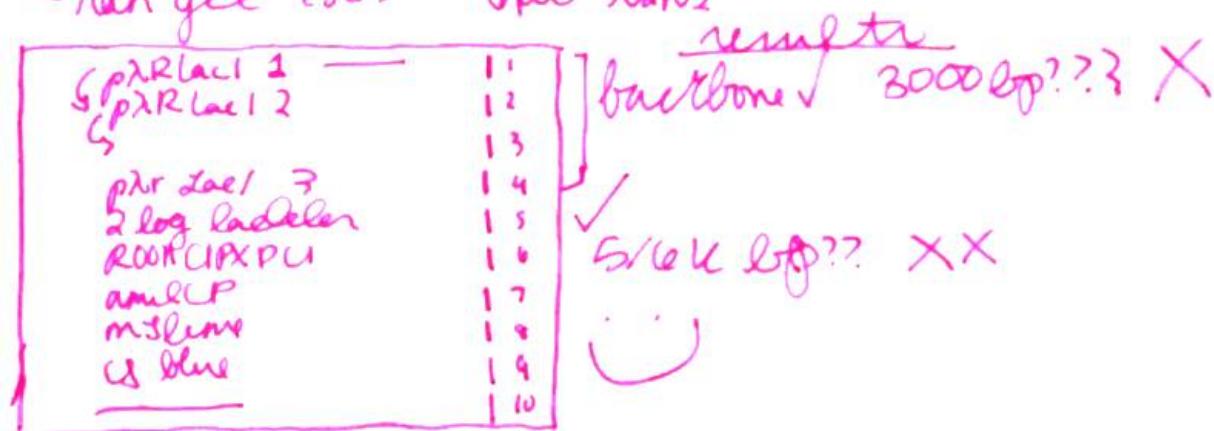
Digest of p<sub>X</sub>R

- : p<sub>X</sub>R LaeI 1
- : p<sub>X</sub>R LaeI 2
- : p<sub>X</sub>R LaeI 3
- : R<sub>600</sub> II ClpXPCU 2
- : amil CP
- : mS11uv
- : Cg Blue

GIB

~nan gel (JL)

6μl runs



## 10/17 - Ligation Calculations

Part	Lot 1	L of Vector	Total	'/kb	uL
IT3	<u>1942</u> 1975	2461	<u>4403</u> <u>4436</u>	<u>0.227</u> <u>0.225</u>	1
PfRLacl	1187	2204	3391	0.295	$\frac{0.769}{0.763} \times 3 = \frac{2.307}{2.289}$
TsPurple	755	0	755	1.325	$0.171 \times 3 = 0.513$
TsPurple DAS	788	0	788	1.269	$0.177 \times 3 = 0.531$
TsPurple LAA	788	0	788	1.269	$0.177 \times 3 = 0.531$

PfRLacl-TsPurple

1 uL IT3

2.307 uL PfRLacl

0.513 uL TsPurple

2 uL Ligate Buffer

1 uL Ligase

13.19 uL nucH<sub>2</sub>O

PfRLacl-TsPurple DAS/LAA

1 uL IT3

2.289 uL PfRLacl

0.531 uL TsPurple DAS/LAA

2 uL Ligate Buffer

1 uL Ligase

13.18 uL nucH<sub>2</sub>O

10/18

- forward/reverse RODC1PAPC1 and all primers annealed  
hydrated

→ 5 uL in sequencing box

(→ 5 uL in Stand's DNA box closet)

rent for sequencing

- RODN1PXP (1 (middle area) (two tubes, F & R)

forward reverse  
ANV213 ANV214

- 4 uL dNTP (193 ng/uL)

- 7.2 uL water (milli-Q)

- 0.8 uL primers

10/18 Segmentation (continued):  
 19 CJB VF2 22 ACP VF2  
 20 CJB VR 23 ACP VR

Alyssa  
 ↘

10/25/17 JL

liquid cultures of:

- J23101 (2) cam
- GFP (2) cam
- R0011::lexPXP1::reg (2) ampr
- R0011::lexPXP1 (2) ampr
- pARlacI::tetracycline (2) tet

10/25/17 - (Gramm) continuation of ligation of TsPurple into IC3

Part	Lot 1	Lot V	Total	1/kb	µL	(Digestion by Lauren)
IC3	755	2070	2825	0.354	1	
TsPurple A	755	0	755	1.325	0.802	15.198 µL mg H <sub>2</sub> O
---	---	---	---	---	---	---
IC3	788	2070	2858	0.350	1	
TsPurple	788	0	788	1.269	0.827	15.173 µL mg H <sub>2</sub> O
DNA Ladder						
Gel-						
					2 µL Buffer	
					1 µL Ligase	



10/26/2017 - Virginia Violet TS/Amp/pUC/HC

- 1) Centrifuge tube briefly before opening to ensure DNA is not in the lid.
- 2) Add 20 μL sterile water or 1X TE, and incubate 10 min at room temp.
- 3) Dissolve DNA by pipetting up and down a few times.
- 4) Long-term storage at -20°C.

\* 1 ng/μL stock  $\Rightarrow$  1 μL solution, 99 μL H<sub>2</sub>O

20 μL    2 μg dna

$$2000 \text{ ng} / 20 \mu\text{L} = 100 \text{ ng} / \mu\text{L}$$

$$1 \mu\text{L} + 99 \mu\text{L H}_2\text{O} = 1 \text{ ng}/\mu\text{L}$$

10/26 LH

## transformation

- Virginia violet
- meadow trillium
- trillium DAS

>10/26/17 -

Grauman/Lauren Minifrops

Part	Cone	A260/A280	A260
1) J23101 #1	74.9 ng/uL	1.85	1.538
2) J23101 #2	95.9 ng/uL	1.82	1.919
3) R11C4 XPC1 #1	312.3 ng/uL	1.84	6.244
4) PHRlac1 TsPurple	30.1 ng/uL	1.47	0.402
5) R11C1 XPC1 seq #1	243.1 ng/uL	1.85	4.874
6) <del>R11C4 XPC1 seq #2</del>			
7) GFP #1	106.8 ng/uL	1.89	2.137
8) R11C1 XPC1 #2	159.0 ng/uL	1.82	3.180
9) GFP Unknown #1	248.0 ng/uL	1.83	4.973
10) GFP Unknown #2	204.8 ng/uL	1.74	4.097
11) GFP #2	173.5 ng/uL	1.41	3.470

# Parts

\*TSp = TS Purple

			<u>Backbones</u>
49	P <sub>λ</sub> R	Lambda Promoter	
11	B33	Weak RBS	2204 iK3
34	B34	Strong RBS	2070 IC3
1064	LacI	Protein generator	
717	eGFP	Green Fluorescent Protein	
129	B15	double terminator	
816	CI	Lambda Repressor	
753	GFP-DAS	GFP w/deg tag	ts purp 755
753	GFP-LAA	GFP w/deg tag	ts purp-das 788
624	CIP	Part 1 of Protease	ts purp-laa 788
1317	CIPX	Part 2 of Protease	
55	R0011	PLac Promoter	
→ 155	TSpurp	Purple Chromoprotein	
→ 188	TSP DAS*	↓ w/deg tag	
→ 188	TSP LAA*	↓ w/deg tag	
33	LAA+DAS	deg tags (each separate)	
11iK3	P <sub>2</sub> R - LacI		
1875	P <sub>2</sub> R - B33 - LacI - B34 - GFP		
2147	B15 - R0011 - B33 - CIPX - B33 - CIP		
1963	B33 - CIPX - B33 - CIP		
1253	B15 - P <sub>2</sub> R - B33 - LacI		
827	B33 - CI		
→ 787	B34 - GFP - DAS		
→ 787	B34 - GFP - LAA		
→ 1848	P <sub>2</sub> R - B33 - LacI - B34 - TSpurp		
→ 2079	P <sub>2</sub> R - B33 - LacI - B34 - TSpurp DAS		
→ 2078	P <sub>2</sub> R - B33 - LacI - B34 - TSpurp LAA		
2974	B15 - R0011 - B33 - CIPX - B33 - CIP - B33 - CI		
→ 724	B34 - TSpurp		
1911	P <sub>2</sub> R - B33 - LacI - B34 - GFP LAA		

nmnstop

$$\text{ACP} = \frac{\text{A}260(10\text{mm})}{3} - 0.625 \\ \text{A}260/\text{A}280 - 2.00 \\ 31.3 \text{ ng/mL}$$

$$\text{ACP} = \frac{\text{A}260(10\text{mm})}{1} - 1.697 \\ \text{A}260/\text{A}280 - 2.01 \\ 89.9 \text{ ng/mL}$$

~~$$\text{ACP} = \frac{\text{A}260(10\text{mm})}{2} - 25.370 \\ \text{A}260/\text{A}280 - 1.05 \\ 1268.9 \text{ ng/mL}$$~~

$$\text{mL} = \frac{\text{A}260(10\text{mm})}{1} - 1.513 \\ \text{A}260/\text{A}280 - 1.84 \\ 75.7 \text{ ng/mL}$$

$$\text{mL} = \frac{\text{A}260(10\text{mm})}{3} - 0.773 \\ \text{A}260/\text{A}280 - 2.12 \\ 38.6 \text{ ng/mL}$$

$$\text{CJB} : \frac{\text{A}260(10\text{mm})}{2} - 3.574 \\ \text{A}260/\text{A}280 - 1.49 \\ 178.7 \text{ ng/mL}$$

$$\text{CJB} : \frac{\text{A}260(10\text{mm})}{1} - 1.236 \\ \text{A}260/\text{A}280 - 1.69 \\ 61.8 \text{ ng/mL}$$

$$\text{CJB} = \frac{\text{A}260(10\text{mm})}{2} - 1.836 \\ \text{A}260/\text{A}280 - 1.93 \\ 91.8 \text{ ng/mL}$$

$$\text{mL} = \frac{\text{A}260(10\text{mm})}{2} - 0.943 \\ \text{A}260/\text{A}280 - 1.67 \\ 47.2 \text{ ng/mL}$$

### Digest 9/20/2017

CJB:  $\frac{178.7 \text{ ng}}{1 \mu\text{L}} = \frac{1 \mu\text{g}}{5 \mu\text{L}}$  37.4  $\mu\text{L}$  1.  $\mu\text{L}$  5  $\mu\text{L}$  cutSmart

mL 1:  $\frac{75.7 \text{ ng}}{1 \mu\text{L}} = \frac{1 \mu\text{g}}{13.21 \mu\text{L}}$  29.79  $\mu\text{L}$  1  $\mu\text{L}$  X, 1  $\mu\text{L}$  P 5  $\mu\text{L}$  cutSmart

ACP 1:  $\frac{34.9 \text{ ng}}{1 \mu\text{L}} = \frac{1 \mu\text{g}}{11.78 \mu\text{L}}$  31.22  $\mu\text{L}$  1  $\mu\text{L}$  X, 1  $\mu\text{L}$  P 5  $\mu\text{L}$  cutSmart

# Gel 9/21 (of digest)

C3B	?
2 Log Ladder	
MJL	
2 Log Ladder	
ACP	

~~gauge~~

C3B3	
2-LL	
MJL	
2-LL	
ACP	

# Gel 9/15

C3B	11
C3B2	12
2 Log 3	13
MJL1	14
MJL2	15
MJL3	16
2 Log	17
ACP1	18
ACP2	19
ACP3	10

failed w/ smears

← looked decent, rest trash  
... maybe PCR materials

↳ liquid cultures → draw from new  
colonies on plates but use  
MJL 1 + 2 dilutions instead  
of plates

Ordered parts from IDT & C time

## Hydration:

w/ ordered  
parts

$$1000 \text{ ng} \quad | \quad 40 \mu\text{L} = 25 \text{ ng}/\mu\text{L}$$

mJL  
CJB  
ACP

10/2/2017: Hydrated amilcp, amay lime, cjbline  
- stored in big freezer in standy's bix

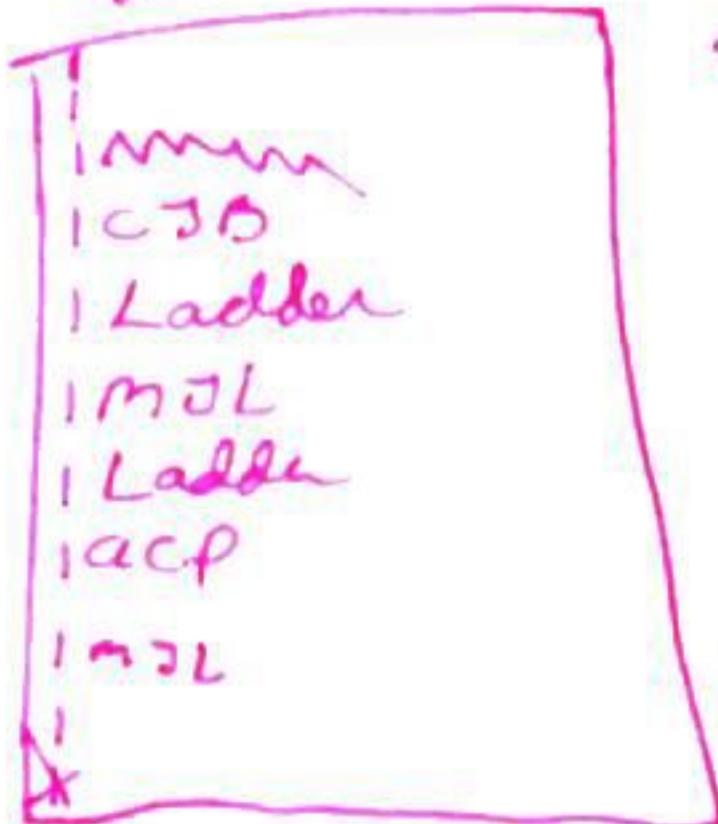
Digestion: 10/3/17

10 mL DNA of 25 ng/ $\mu$ L  
~~2  $\mu$ L E~~  
 1  $\mu$ L P  
 7 mL H<sub>2</sub>O

gel 10/4

25 mL 10/10/17  
 10 DNA  
 2.5 2.1 buffer  
 .5 EcoRI  
 .5 PstI  
 11.5 H<sub>2</sub>O

success & correct (the acp didn't show well)



Ligation: AF 10/5 → failed → due to wrong digestion

Parts	L part	L vector	total	kb	uL	
IC3	822	2070	2.892	.346	1	1 uL Ligase
CJB	822	0	.822	1.21	.8575	2 uL Li. Buff 15.143 uL H <sub>2</sub> O
IC3	822	2070	2.859	.3498	1	1 uL Ligase
ACP	789	0	.787	1.26	.833	2 uL Li. Buff 15.167 uL H <sub>2</sub> O
IC3	822	2070	2.883	.347	1	1 uL Ligase
MJL	813	0	.813	1.23	.846	2 uL Li. Buff 15.154 uL H <sub>2</sub> O

Transformation 10/6/17

CJB IC3 - growth, no color

ACP IC3 - growth, no color

MJL IC3 - growth, no color

control - no growth

ligation failed, IC3 closed on self.

digest → phosphatase IC3  
retransform & religate

Digest 10/9/17

linearized IC3

↳ iGEM protocol

Digest 10/10/17

colors: MJL

25 uL sda

CJB

10 uL DNA

+ 11.5 uL H<sub>2</sub>O

ACP

2.5 uL 2.1 Buffer

.5 uL EcoR1

.5 uL Pst1

Ligation 10/10/17

using 10/5 calculations

failed

positive control worked  
check digestion on gel

Gel 10/11/17



all are correct, ligation is what failed  
(+ transfection)

Digatur 10/10/17

S nt overnight LQ → miniprep → sequencing

10/18/2017 - 10/19/2017

- Digest of Amil CP, Amaj Lime, CJ Blue

10 uL DNA

5 uL NEBuffer 2.1

1 uL EcoRI-HF

1 uL PstI

33 uL dH<sub>2</sub>O

- Gel  $\Rightarrow$  Part lengths look correct

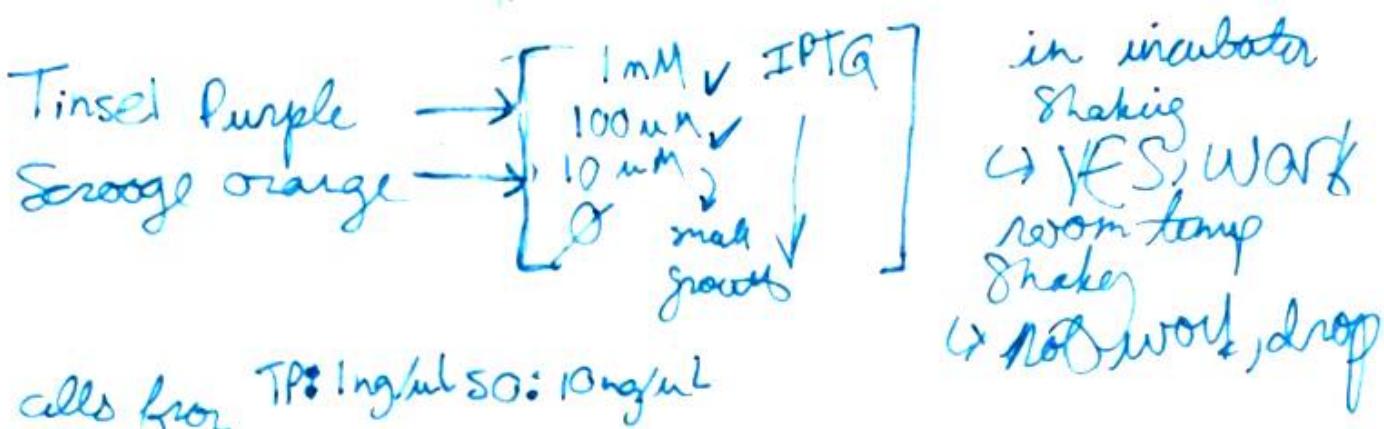
- Ligate to IA3

Part	Part length	Vector length	Total length	kbp	uL	
IA3	822	2155	2977	0.336	1	+ 2uL lig buffer 1uL ligase 15.172uL H <sub>2</sub> O
CJ Blue	822	0	822	1.217	0.828	
IA3	789	2155	2944	0.340	1	+ 2uL lig buffer 1uL ligase 15.195uL H <sub>2</sub> O
Amil CP	789	0	789	1.267	0.805	
IA3	813	2155	2968	0.337	1	+ 2uL lig buffer 1uL ligase 15.178uL H <sub>2</sub> O
Amaj Lime	813	0	813	1.230	0.822	

23°C for 10 min, 45°C for 10 min

10/25/17

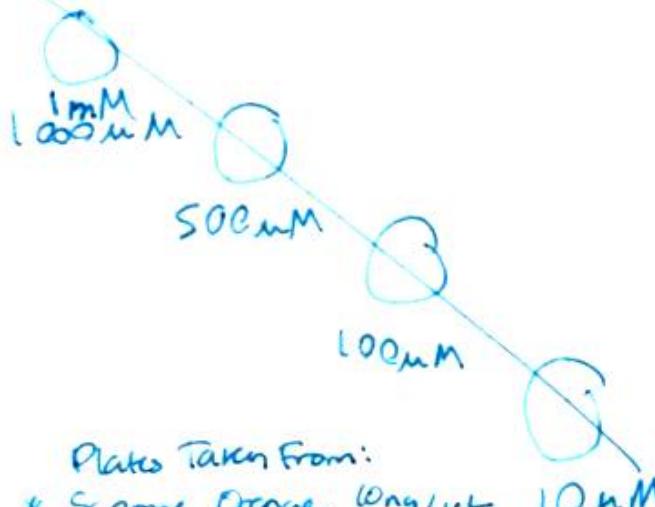
Liquid Culture (amp resistance)



10/26/18

Doubling change both 1mM + 100µM so adding 500nM  
(amp resistance)

triplicates



- Plates Taken From:  
\* Scrooge Orange 1ng/ml  
(10/24 Trans TAS)  
\* Tinsel Purple 1ng/ml  
(10/24 Trans TAS)  
\* added (-) control  
to test sterility LB

TP	from:	1mM	IPTG
500nM	ng/ml	10nM IPTG	
100nM		IPTG	
SO	resistance	1mM	IPTG
10nM	ng/ml	10nM IPTG	
IPTG		IPTG	