

COLOR CODING KEY

Spectrophotometry Measurements

Construct Group

Plasmid Group

Interlab

Cell Culture/Plating

Biobrick Group

Cyanobacteria Transformation Group

Experimental Verification

Plasmid & Construct Design Group

Week 4

June 24, 2018

Spectrophotometry Measurements at 12:00 (Stephanie)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 0.679
- UTEX 1% room temp. 6/13 2m A= 0.885

Spectrophotometry Measurements at 23:10 (Lin)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 0.890
- UTEX 1% room temp. 6/13 2m A= 0.872

Plasmid Group (Lin)

- New streaks of 1414, 1579, 2991 bacteria on spectinomycin plates

June 25, 2018

Spectrophotometry Measurements at 8:09 (Priya)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 0.870
- UTEX 1% room temp. 6/13 2m A= 0.847

Spectrophotometry Measurements at 16:00 (Natalie)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 0.480
- UTEX 1% room temp. 6/13 2m A= 0.493

Cell Culturing/Plating (Natalie)

- Supplementing measurement flasks with culture
- Added 20 mL of culture from UTEX 1% room temp. 6/13 1e into UTEX 1% room temp. 6/13 1m

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- Added 20 mL of culture from UTEX 1% room temp. 6/13 2e into UTEX 1% room temp. 6/13 2m

Construct Group (Karthik/Matthew/Natalie/Woody/Dominika)

- G-block Resuspension of PpsbA2_sps and PpsbA2_cscB (followed IDT protocol)
 - Pelleted down the dry DNA for the two tubes
 - Added 25 μL of autoclaved Milli-Q water to PpsbA2_sps and PpsbA2_cscB
 - Left the two tubes in the water bath at 53 $^{\circ}\text{C}$ for 15 minutes
 - Vortexed the two tubes and then centrifuged them
 - Nanodrop:
 - PpsbA2_sps= 3.7 ng/ μL , 5.1 ng/ μL
 - PpsbA2_cscB = 4.9 ng/ μL , 8.0 ng/ μL
- Primers resuspension of 1579 PcscB start, 1579 PcscB end, 1579 Psps start, 1579 Psps end (followed IDT protocol)
 - Pelleted down the dry DNA for all four tubes
 - Resuspended 1579 PcscB start with 796 μL nuclease free water
 - Resuspended 1579 PcscB end with 860 μL nuclease free water
 - Resuspended 1579 Psps start with 872 μL nuclease free water
 - Resuspended 1579 Psps end with 804 μL nuclease free water
 - Left the four tubes in the water bath at 50 $^{\circ}\text{C}$ for 15 minutes
 - Vortexed the four tubes and centrifuged them
- PCR of PpsbA2_sps and PpsbA2_cscB (Short Cycle)
 - PCR Mixture 50 μL
 -

Phire Mastermix	25 μL
Forward Primer	1 μL
Reverse Primer	1 μL
Template DNA	2.5 μL

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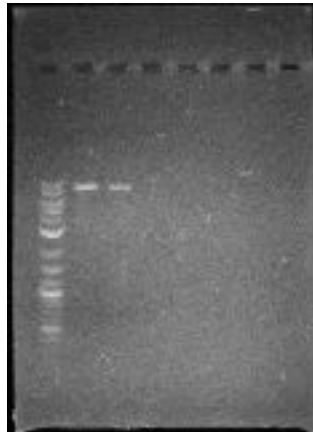
Plasmid & Construct Design Group

H ₂ O	20.5 μ L
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- Nanodrop:
 - PpsbA2_sps = 60.4 ng/ μ L
 - PpsbA2_cscB = 40.5 ng/ μ L

Plasmid Group (Priya/Stephanie/Manvi/Lin)

- MiniPrep x 3
 - 1 miniprep using NEB kit
 - 2 minipreps using Qiagen kit
- Concentration of DNA = 5.4 ng/ μ L, 11.6 ng/ μ L, 58 ng/ μ L (not DNA curve)
- Attempted with Mary Lou and DNA curve \rightarrow 26.7 ng/ μ L
- Split 1579
- made liquid inoculation for 1414 and 2991
- Gel extraction/gel electrophoresis for 1579



- Agarose gel (1%) used
- EcoRV and Sal I with cutsmart buffer
- total = 45 mL rex, incubated 1 hour at 37 $^{\circ}$ C
- Added 9 μ L loading dye
- 1. 1 kb MW maker

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- 2. 1579 DNA
- 3. 1579 DNA

June 26, 2018

Construct Group (Karthik/Matthew/Natalie/Woody)

- PCR purification of PpsbA2_cscB (following Qiaquick protocol)
 - Didn't add sodium acetate (skipped step 1)
 - Step 7, followed the increased DNA concentration step
 - 80 μ L PB buffer
 - 16 μ L used to purify
 - 30 μ L of Milli-Q Water
 - 1 μ L used on nanodrop
 - Nanodrop of PCR purified
 - PpsbA2_csB = 22.3 ng/ μ L
- Resuspension of PidiA_cscB and PidiA_sps (following IDT protocol)
 - Pelleted down the dry DNA for the two tubes
 - PidiA_cscB \rightarrow 25 μ L of autoclaved milli-Q water
 - PidiA_sps \rightarrow 25 μ L of autoclaved milli-Q water
 - Left the two tubes in the water bath at 50 $^{\circ}$ C for 15 minutes
 - Nanodrop:
 - PidiA_sps = 30.3 ng/ μ L, 6.2 ng/ μ L
 - PidiA_cscB = 13.8 ng/ μ L
- PCR of PidiA_cscB and PidiA_sps (Short Cycle)
 - Nanodrop
 - PidiA_cscB = 606.2 ng/ μ L(used 3 μ L)
 - PidiA_sps= error (used 2.5 μ L)
- PCR purification of PpsbA2_sps (following Qiaquick protocol)
 - 80 μ L of PB buffer
 - 16 μ L of PpsbA2_sps
 - 30 μ L of autoclaved milli-Q water
 - Nanodrop:

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- PpsbA2_sps = 8.2 ng/ μ L, 7.5 ng/ μ L
- PCR purification of PidiA_cscB and PidiA_sps (following Qiaquick protocol)
 - 80 μ L of PB buffer
 - 16 μ L of PidiA_cscB and PidiA_sps
 - 30 μ L of autoclaved milli-Q water
 - Nanodrop:
 - PidiA_cscB = 8.4 ng/ μ L
 - PidiA_sps = 5.6 ng/ μ L

Spectrophotometry Measurements at 16:08 (Lin/Natalie)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 0.633
- UTEX 1% room temp. 6/13 2m A= 0.665

Plasmid Group (Priya/Stephanie/Lin)

- 1579 Miniprep success
 - Nanodrop: 102.8 ng/ μ L
- Split 1414/2991 cultures
- Testing restriction enzyme with digest
- Gel electrophoresis \rightarrow failure as EtBr was not added to solution, but found that PstI worked.

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-
- 1. MW Marker (1 kb)
- 2. PstI+EcoRV
- 3. PstI
- 4. PstI + SalI
- 5. PstI + PvuII

Cell Culturing/Plating (Natalie)

- Split the Syn. UTEX 2434 in half
- 10 mL of culture into new test tube with 10 mL of BG-11
- Supplemented the original culture with 10 mL of BG-11

Cell Culturing/Plating (Natalie)

- Split the UTEX Collier 5/19 culture sup. With 20 mL BG-11
- 25 mL of culture into new flask labeled UTEX room temp. 6/26 (split) and supplemented it with 25 mL of BG-11
- Supplemented original culture with 25 mL of BG-11

June 27, 2018

Construct Group (Karthik/Matthew/Natalie)

- Nanodrop of PCR DNA

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- PidiA_sps = no good results
- Used 2 μL
- PCR purify of PpsbA2_sps
 - 80 μL PB buffer
 - 16 μL of PpsbA2_sps
 - 30 μL of EB buffer
 - Nanodrop:
 - PpsbA2_sps = 9.9 ng/ μL
 - Used 1 μL
- Made .7% agarose gel
 - .4 g of agarose and 500 μL of 100X TAE
 - Left in fridge because when pulling out comb, wells broke

Plasmid Group (Priya/Stephanie/Sara/Lin/Manvi)

- Mini-prepped 1579
 - 102.0 ng/ μL DNA (Had to nanodrop 2X) \rightarrow 27 μL remaining
- Gel electrophoresis for 1579 with ethidium bromide staining

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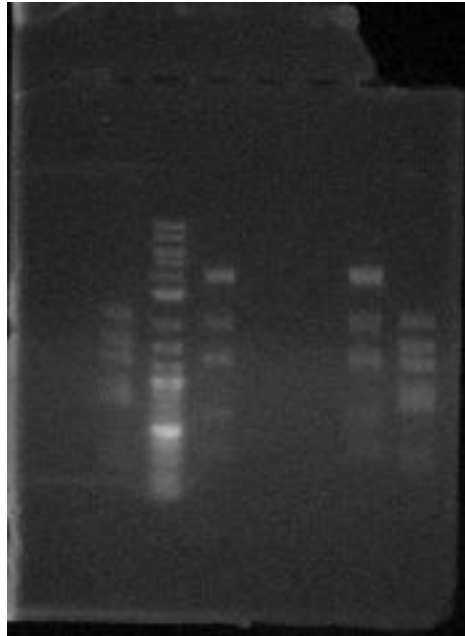
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-
- 1. PstI + PvuII
 - PstI + PvuII, confirmation that both worked
- 2. MW Marker
- 3. PstI+EcoRV
 - Both worked
- 4. PstI + Sall
 - Both worked
- 5. PstI + PvuII
 - Did this twice because we thought this lane could have been punctured
- Gel electrophoresis w/ 1% Agarose for 1579
 - .6 g of agarose and 500 μ L of 100X TAE

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-
- 1. PstI + PvuII
- 2. MW Marker
- 3. PstI+EcoRV
- 4. PstI + Sall
- 5. PstI + PvuII
- 6 g of agarose
- Stained the gel with Diamond nucleic acid Dye that is 10,000 x concentrated
- Diluted 100X TAE into 1X TAE
- Diluted 20 μ L of 10,000X Diamond Dye with 200 mL of 1X TAE
- When transferring gel into box for shaking/staining it broke into pieces and also when transferring gel into UV box, it also broke into pieces
- Split 1579 *E. coli* in half

Spectroscopy Measurement at 17:50 (Natalie)

- Done at 750 nm with 1500 μ L of culture
- UTEX Collier 5/19 Culture sup. with 20 mL BG-11 A=.697
- UTEX room temp. 6/13 (split) A= 1.009

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- UTEX room temp. 6/26 (split) A= .760

Spectroscopy Measurement at 17:56 (Natalie/Sara)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 0.756
- UTEX 1% room temp. 6/13 2m A= 0.751

June 28, 2018

Spectroscopy Measurement at 9:56 (Sara)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A=0.851
- UTEX 1% room temp. 6/13 2m A= 0.715

Spectroscopy Measurement at 18:20 (Natalie)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A=0.933
- UTEX 1% room temp. 6/13 2m A= 0.827

Construct Group (Karthik/Natalie/Manvi/Priya/Woody)

- Made a .7% agarose gel
 - Added .5 μ L 10,000X Diamond Nucleic Acid Dye
 - .4 g of agarose
 - 500 μ L of 100X TAE
 - Ran gel with diamond dye
 - Buffer had 18 μ L of 1X diamond dye, 3.5 mL of 100X TAE, 346.32 mL of Milli-Q water
 - Added MW marker to two wells
- Resuspension of Q3cscB and Q3sps (following IDT protocol)
 - Q3cscB \rightarrow 100 μ L of autoclaved milli-Q water
 - Q3sps \rightarrow 100 μ L of autoclaved milli-Q water
 - Left the two tubes in the water bath at 50 $^{\circ}$ C for 15 minutes

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- Nanodrop (used 3 μ L):
 - Q3cscB= 161.0 ng/ μ L, 92 ng/ μ L (no photos, but really smooth curves), 31.5 ng/ μ L
 - Q3sps = 292.9 ng/ μ L
- Resuspension of 1414 promoters (following IDT protocol)
 - PcpC → 50 μ L of autoclaved milli-Q water
 - PcpC560 → 50 μ L of autoclaved milli-Q water
 - PidiA → 25 μ L of autoclaved milli-Q water
 - PpsbA2 → 25 μ L of autoclaved milli-Q water
 - Prbc → 50 μ L of autoclaved milli-Q water
 - Left the 5 tubes in the water bath at 50 °C for 15 minutes
 - Nanodrop:
 - PcpC560 = 4.4 ng/ μ L, 5.4 ng/ μ L (2 μ L used)
 - PidiA = 9.5 ng/ μ L
 - PpsbA2 = 10.7 ng/ μ L
 - Prbc = 9.0 ng/ μ L, 7.7 ng/ μ L (2 μ L used)
 - PcpC = 3.4 ng/ μ L, 38.9 ng/ μ L (2 μ L used)
- Resuspension of primers (following IDT protocol)
 - 1414 promo start → 704 μ L of autoclaved milli-Q water
 - 1414 promo end → 800 μ L of autoclaved milli-Q water
 - 1579 cscB start → 912 μ L of autoclaved milli-Q water
 - 1579 cscB end → 824 μ L of autoclaved milli-Q water
 - 1579 sps start → 904 μ L of autoclaved milli-Q water
 - 1579 sps end → 908 μ L of autoclaved milli-Q water
- PCR of Q3 cscB (after mixing) (CSCB cycle)
- PCR of Q3 sps (after mixing) (SPS Cycle)
- PCR of PcpC560 (EYPF Cycle)
- PCR of PpsbA2 (SHORT Cycle)

Plasmid Group (Manvi/Priya/Stephanie)

- Miniprep with 1414 plasmid

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- Nanodrop → 203.4 ng/μL
- Miniprep with 2991 plasmid
 - Nanodrop → 92.6 ng/μL
- Made 1% agarose gel for staining with
 - Gel ended up breaking and having to be thrown out, might be a problem with the TAE solution

Interlab (Lin/Matthew/Natalie)

- Finished pipetting and setting up 96 well plate for calibration 1 and calibration 2, but were unable to measure the plates because we didn't know the instrument information
- For Calibration 2, we changed tips when mixing for row E and used the same tips for the entire row for F, G and H
- Left the plate in room temperature overnight

June 29, 2018

Spectrophotometer Measurement at 10:45 (Sara)

- Done at 750 nm with 1500 μL of culture
- UTEX 1% room temp. 6/13 1m A=0.507
- UTEX 1% room temp. 6/13 2m A= 0.353

Spectrophotometer Measurement at 18:28 (Lin/Lukas)

- Done at 750 nm with 1500 μL of culture
- UTEX 1% room temp. 6/13 1m A=1.124
- UTEX 1% room temp. 6/13 2m A=0.892

Construct Group (Woody/Matt/Natalie/Sara)

- PCR of 1414 PidiA (SHORT Cycle)
- PCR of 1414 Prbc (EYFP Cycle)
- PCR of 1414 Pcpc (EYFP Cycle)
- Resuspension of Geneblocks (following IDT protocol)
 - Q3 PisiAB_cscB → 50 μL of autoclaved milli-Q water

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- Q3 PisiAB_sps → 50 μL of autoclaved milli-Q water
- Left the two tubes at 49 °C for 15 minutes
- Nanodrop:
 - Q3 PisiAB_cscB = 13.0 ng/μL
 - Q3 PisiAB_sps = 11.2 ng/μL
- Resuspension of Geneblocks (following IDT protocol)
 - Opto EYFP → 100 μL of autoclaved milli-Q water
 - Orig EYFP → 100 μL of autoclaved milli-Q water
 - Left the two tubes at 50 °C for 15 minutes
 - Nanodrop:
 - Opto EYFP = 15.5 ng/μL
 - Orig EYFP = 8.3 ng/μL
- Resuspension of Q1 Primers (following IDT protocol)
 - 2991 start lone → 860 μL of autoclaved milli-Q water
 - 2991 end lone → 912 μL of autoclaved milli-Q water
 - Left the two tubes at 51 °C for 15 minutes
- PCR of Q3 PisiAB_cscB (EYFP Cycle)
- PCR of Q3 PisiAB_sps (EYFP Cycle)
- PCR of Opto EYFP (EYFP Cycle)
- PCR of Orig EYFP (EYFP Cycle)

Plasmid Group (Priya/Stephanie/Manvi/Sara)

- Tried making a gel with the stationary agitation method with diamond nucleic dye
 - Broke twice
- Made a gel with diamond nucleic acid dye → SUCCESS!!
 - Incubated for 20 minutes
 - Loaded gel with RE digest of plasmid DNA
 - Banding was smeary but indicated that ECORI and BamHi were both working

Interlab (Lin)

- Made fluorescein stock for Calibration 3

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June 30, 2018

Spectrophotometer Measurement at 13:28 (Lin)

- Done at 750 nm with 1500 μ L of culture
- UTEX 1% room temp. 6/13 1m A= 1.275
- UTEX 1% room temp. 6/13 2m A= 0.984