Summer 2022 Week 6

WEDNESDAY, 7/6/2022

AcrB Amp MIC

- Plated E.Coli, ISx and AcrB deletion
- Follow Up-
 - Innoculate 3 ISx colonies
 - Innoculate 3 AcrB Colonies
 - o Innoculate 1 E. Coli Colony

LB dilutions

Tem-1 Gene Detector

- Run a PCR of the AcrB Homologies
- Upstream
 - 93 uL water
 - 30 uL Phusion buffer
 - 7.5 uL 027
 - 7.5 uL AcrB Up Forward
 - 4.5 uL DMSO
 - 3 uL mM dNTP's
 - 1.5 uL Phusion polymerase
 - NEB = 65°C
 - Run at 62°C
- Downstream
 - 93 uL water
 - 30 uL Phusion buffer
 - 7.5 uL 029
 - 7.5 uL AcrB Down Revers
 - 4.5 uL DMSO
 - ☑ 3 uL mM dNTP's
 - 1.5 uL Phusion polymerase
 - NEB = 66°C
 - Run at 62°C

NPTII Gene Detector

- Run a PCR of the 2049 Homologies
- Upstream
 - 93 uL water
 - 30 uL Phusion buffer
 - 7.5 uL 001
 - 7.5 uL 025
 - 4.5 uL DMSO
 - 3 uL mM dNTP's
 - 1.5 uL Phusion polymerase
 - NEB= 59°C
 - Run at 59
- Downstream
 - 93 uL water
 - 30 uL Phusion buffer
 - 7.5 uL 026

- 7.5 uL 004
- 4.5 uL DMSO
- ☑ 3 uL mM dNTP's
- 1.5 uL Phusion polymerase
- NEB = 60°C
- Run at 60°C
- Run a gel to determine success of all PCR's
- Gel Order: Ladder-CAU-CAD-C2U-C2D-AcrU-AcrU-AcrD-AcrD-2049U-2049U-2049D-2049D

Screen Shot 2022-07-07 at 3.03.29 PM.png

- AcrB
 - Annealing temps were too high
 - Perhaps try a temperature gradient
- 2049
 - Annealing Temps were too high
 - Try running them at 56°C ?

THURSDAY, 7/7/2022

AcrB deletion-

- Innoculated 3 colonies of ADP1-Isx
- Innoculated 3 colonies of ADP1 AcrB deletion
- Innoculated 1 colony of E. Coli

https://benchling.com/zen_blade/f/lib_VddSY4bs-summer-lab-sessions/etr_SKfJ8Mzm-summer-2022-week-6/view?versionId=579957930¬e=true&protocols=#p... 2/5

- Follow Up-
 - Add 2 uL of each stock to the following dilutions
 - 128 ug/mL
 - 64 ug/mL
 - 32 ug/mL
 - 16 ug/mL
 - 8 ug/mL
 - 4 ug/mL
 - 2 ug/mL
 - 0 ug/mL
- Set Up

Tem-1 Gene Detector

- Follow Up- Run a PCR of the AcrB homologies against a temperature gradient?
 - Run at 60° and 64° ?

NPTII Gene Detector

- Run a PCR of the 2049 Homologies
- Upstream
 - 🧹 93 uL water
 - 30 uL Phusion buffer
 - 7.5 uL 001
 - 7.5 uL 025
 - 4.5 uL DMSO
 - 🗸 3 uL mM dNTP's
 - 1.5 uL Phusion polymerase
 - NEB= 59°C
 - Run at 56°C
- Downstream
 - 93 uL water
 - 30 uL Phusion buffer
 - 7.5 uL 026
 - 7.5 uL 004
 - 4.5 uL DMSO
 - 3 uL mM dNTP's
 - 1.5 uL Phusion polymerase
 - NEB = 60°C
 - Run at 56°C
- Follow Up- Run a gel to determine success of 2049 PCR

Expected-

FRIDAY, 7/8/2022

AcrB deletion-

- Follow Up-
 - Put 200 uL of each dilution into the 96 well plate
 - Run it in the plate reader
 - OD Measures light scattering
 - High OD measurement = High scattering = Large amount of particles = turbidity
 - Low OD = Low Scattering = Low amt of particles = no turbidity

Tem-1 Gene Detector

NPTII Gene Detector

- Ran a gel to determine the success of the PCR
 - Expected Results
 - Upstream = 1266
 - Downstream 1213
 - Gel Order: Ladder-CU-CD-U-U-D-D

NPTII Detector Homologies.png					
				12 38	
Ladder	Up Control	Down Control	Upstream homolgy	Downstream Homolgy	
2kb 1kb		-			

- The Gel looks really good!
- Clean and Concentrate the Product and start a GGA reaction
 - GGA Reaction Conditions
 - 🔽 10.8 uL water
 - 2 uL T4 ligase Buffer
 - 2 uL Bsmbl mix
 - 2 uL Upstream
 - 1.3 uL ng Downstream
 - 1.1 uL PCR 1
 - .8 uL PCR 2
- Grew an overnight culture of 2049 deletion strain
- Follow Up-

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- Run gel to determine if the expected product is seen
- Put 15 uL into culture tube w/ 500 uL LB and 35 uL overnight Culture

Follow Up- Cameron made a plate of the marionette strains, tomorrow I need to innoculate cultures of those

• Follow Up-

- Plate +DNA and -DNA cultures onto AZT plates
 - 10⁻² dilution of +DNA
 - Serial dilution of DNA

Design More Mutations

<u>NPTII (pKD13)</u>

- $\circ~$ 10 bp deletion @ site 1440-1449
- Proposed Deletions-
 - Mutate the catalytic site
 - 10 bp deletion at the catalytic site
 - premature stop codon at the catalytic site

Tem-1 (pBTK404)

- 10 bp deletion @ site 1312-1321
- Mutate Catalytic Serine from AGC -> CTC @ site 1112-1114
- Proposed Deletions-
 - Mutate one of the Leucines to a polar AA at site 1312-1321
 - Premature stop codon at site 1312-1321
 - 10 bp deletion of te catalytic site @1112-1114