#### (RT2-1) Investigating a Plasmid and Recombination-Based System for Genome Editing in *Enterobacter Iudwigii*

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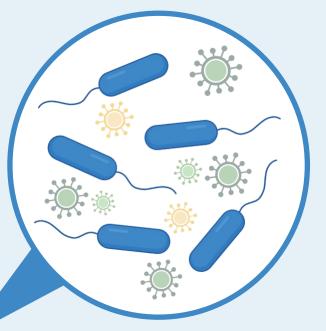






Use of chemicals is prevalent and problematic

IMI





**Context & Overarching Project Goals** 

- 1. Develop better biological understanding of sink-adapted *Enterobacter ludwigii*
- 2. Genetically engineer *E. ludwigii* to be more probiotic and/or antimicrobial



### Computational

- 1. Identify possible locations for insertion
- 2. Return information about potential downstream effects of insertion

## Wet Lab

Conduct foundational genome editing experiments, attempting recombination at chosen locations



Methods

Identify positions for insertion in genome (Prokka-annotated E10 genome and chose ampE, poxB, and hcp1)

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MOM

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Plasmid construction (Design, PCR, and Gibson assembly)

**(x3)** 

Delivery into PIR1 *E. coli* cells (Electroporation)

**(x3)** 

Delivery to E10 *E. ludwigii* cells (*Electroporation*)

**(x3)** 

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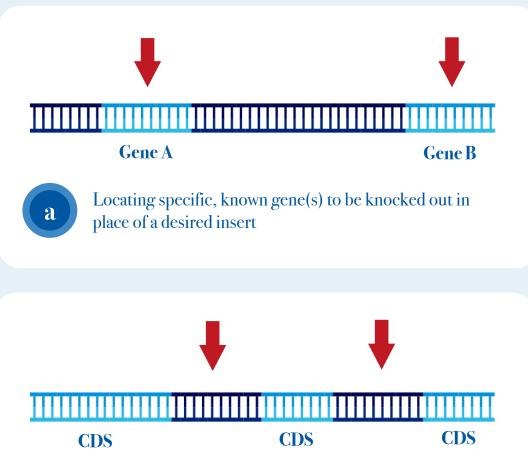
Analysis of recombination success (4 colonies per knockout type)

**(x3)** 





Results





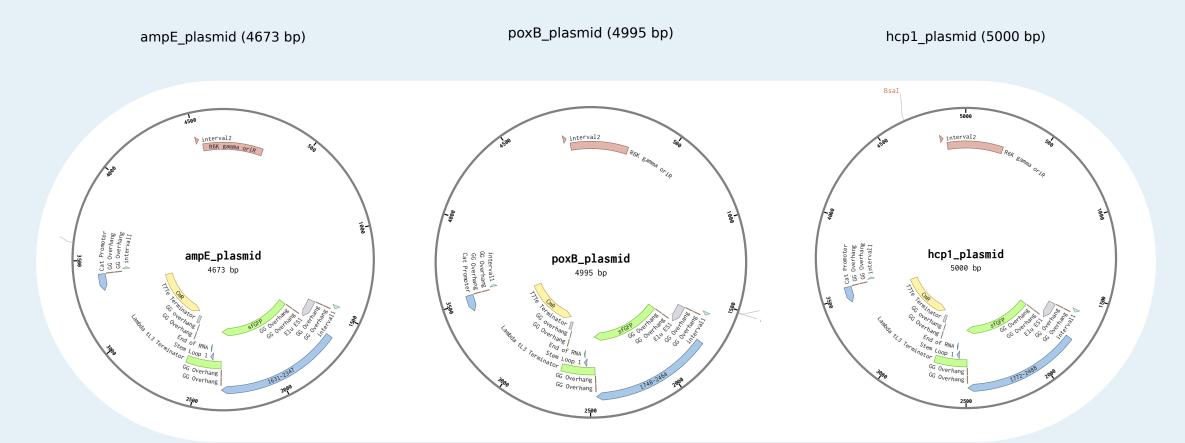
Locating intervals that fall between coding sequences (CDSs), insertion will *not* directly disrupt a gene

#### **Results Pt. 1** *Successes in Computational Outputs*

- Program to locate sites for insertion
- Program to identify genes that may be lost or gained due to an insertion
- Shared and documented on GitHub

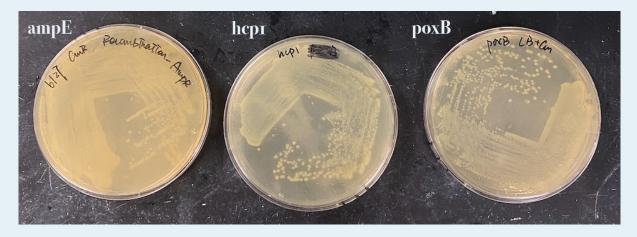
**Designed Plasmids** 

Specific origin, GFP, CmR, and homologous arms

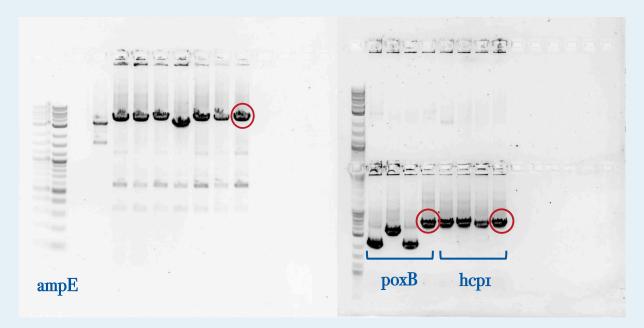


#### **Results Pt. 2** *Successes in Plasmid Delivery*

- Successful plasmid design and creation
  - Illumina sequenced
- Transformation into PIR1 (*E. coli*)
- Likely successful delivery of plasmids into E10
  - Reasonable voltage and time of electroporation
  - Chloramphenicol resistance



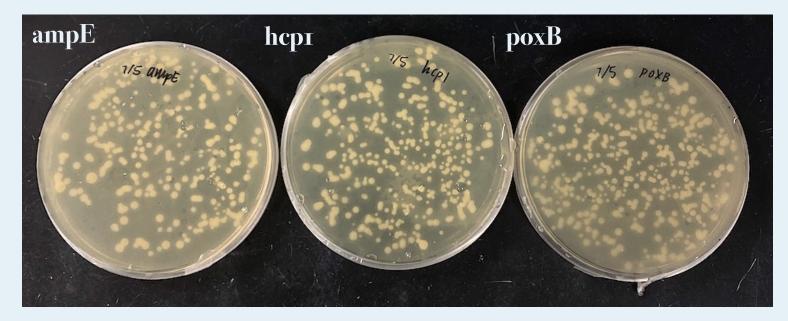
Survival of transformed PIR1 (*E. coli*) on selective media



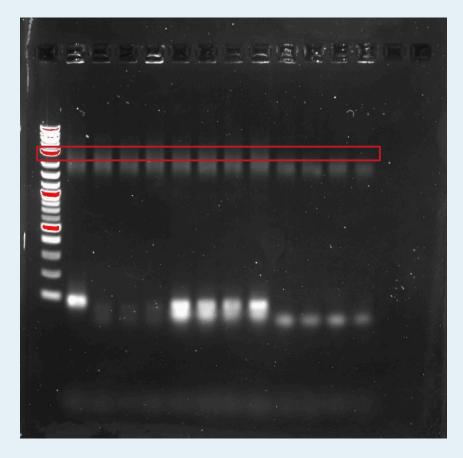
Validation of plasmids' presence in sample PIR1 colonies

#### **Results Pt. 3** *Recombination Outcomes*

- 4 colonies isolated per transformation (12 total)
- Detection of false positives
- Inserts were *not* integrated into our samples' genomes



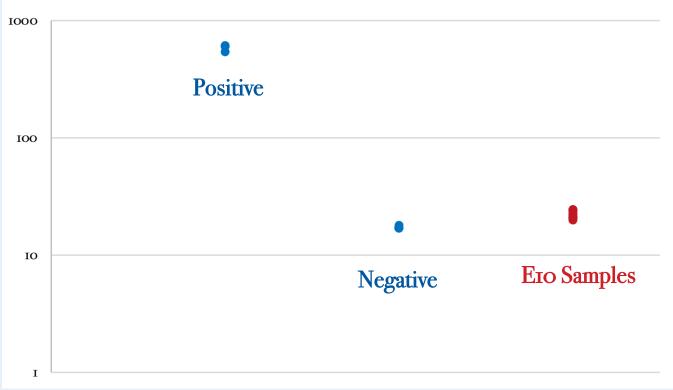
Survival of transformed E10 on selective media



No apparent amplification of target sequence (roughly 3 kb)

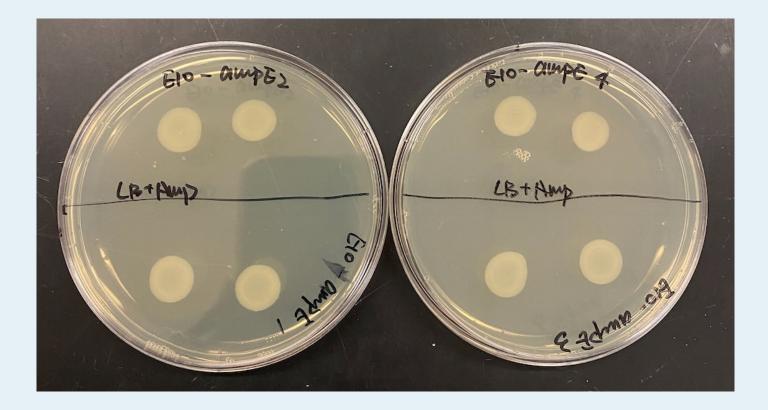
#### Results Pt. 3 (Continued) All Attempted Knockouts





Median fluorescence levels of E10 samples resembled those of negative controls

- Following colony PCR of transformed E10 samples, no bands of appropriate length on gel
- No fluorescence detected via Flow cytometry



Results Pt. 3 (Continued) ampE Attempted Knockout

• Unintended growth on selective media (ampicillin)

# Future Directions

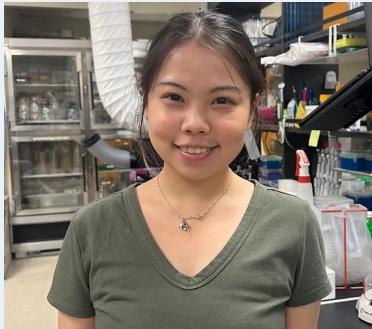
#### Immediate

- Continue utilizing program to find locations for insertion and analyze desirability
- Repeat experiment, tweaking certain aspects of protocol
  - Ex: increase length of homologous arms in designed plasmids, sample more *E. ludwigii* colonies following transformation, etc.
- Explore other possible methods of genetic engineering
  - Ex: CRISPR technology

#### Long Term

- Determine useful mechanisms for *E. ludwigii* to employ within sink microbiome
- Identify and harness specific functional genes to increase *E. ludwigii* robustness and probiotic/antimicrobial characteristics





# Thanks to Dr. Crook and Qiaochu Li!



