# Expanding the Dynamic Range of Methanosarcina acetivorans through **Recombinant Expression of an RNA Polymerase** Laura Kirshenbaum<sup>1</sup>, Connor Hines<sup>1</sup>, Tyler Bartolome<sup>2</sup>, Nicole Buan<sup>1</sup>

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## Abstract

Methanogens are anaerobic archaea that grow and conserve energy through the production of methane from acetate or C1 carbon substrates such as methanol and carbon dioxide. They play an important role in the global carbon cycle, contributing between 350 and 420 Tg of methane to the atmosphere annually. When used in anaerobic digesters, methanogens ferment organic waste to produce biogas containing methane, which can be used as a renewable fuel source. Due to their use of low-cost feedstocks and their unique metabolism, methanogens can also be a cheap and efficient host organism for the synthesis of organic molecules such as bioplastics and isoprene, a precursor to rubber. One limiting factor for the industrial use of methanogens is a limited understanding of archaeal transcription and translation, and limited genetic tools available for their regulation. This study aims to expand the dynamic range of gene expression of Methanosarcina acetivorans through the recombinant in vivo expression of an RNA Polymerase. Using *in vitro* transcription, we show that the RNA Polymerase effectively transcribes genes in the presence of *M. acetivorans* Iysate. We then demonstrate that the RNA Polymerase produces high levels of transcription in M. acetivorans in vivo by expressing the reporter gene YFAST under the polymerase's specific promoter. Additionally, we demonstrate a new tool for understanding transcriptional and translational regulation in methanogens. Going forward, a new inducible expression system will be introduced in *M. acetivorans* as well to push its dynamic transcriptional capabilities further.

## **Research Questions**

- Will a heterologous RNA polymerase be inhibited by *Methanosarcina* acetivorans?
- Can the dynamic range of gene expression in *Methanosarcina acetivorans* be increased using a recombinant RNA polymerase system?

## Background



Figure 1. Scanning electron micrograph (SEM) of Methanosarcina acetivorans

### **Applications for Environmental** Security

- Methanogens can be used as a renewable fuel source.
- Methanogens have been shown to be a renewable source of isoprene, indicating they could produce terpenes, which are predicted to become a billion dollar industry by 2028.
- Methanogens can synthesize products such as terpenes from cheaper substrates than most bacteria and plants, and are a cleaner source than fossil fuels.

### Methanogens

- Anaerobic archaea that grow and conserve energy by producing methane.
- *Methanosarcina acetivorans* is a species of methanogen isolated from marine sediment, often used for genetic engineering.
- Have isoprenoid lipids in their cell membrane rather than the fatty acid lipids found in bacteria and eukaryotes.



Figure 2. Isoprene, an organic compound that can be synthesized by methanogens, can be used as a precursor to rubber.

## Methods



### Results

### **RNAP** is Not Inhibited *in vitro*



Figure 8. qPCR fluorescence shows *M. acetivorans* cell lysate is no less inhibitory than *E. coli* cell lysate. Plasmid DNA was transcribed, reverse transcribed, and amplified with qPCR to compare transcription levels.

E. coli DH10B: Protein expression strain *E. coli* DH5 $\alpha$ : Plasmid expression strain \*\* = p < 0.005 \* = p < 0.05 ns = p > 0.05 n = 10, 15, 7, 6, 15, 7

Strain

NB34

NB647

NB658

NB659

NB645 pLK15

pLK28

Pmcr-YFAST

Figure 5. Plasmids with YFAST expressed under various promoter systems are transformed into M. acetivorans. Cells are grown to mid-log phase, incubated with HMBR, and their fluorescence is

Figure 6. Inducible expression of the RNA polymerase gene and the

A) Genetic elements of pLK28 without the inducer. The permease and operator are constituitively expressed while the RNA polymerase gene

operator, and the RNA polymerase gene are all expressed. The RNA

**Figure 7.** The inducer molecule is used to induce expression of the RNA polymerase and of YFAST. It is imported into cells through a permease and binds to the repressor protein. Once bound, the repressor

### **RNAP Expresses YFAST** *in vivo*



<b>,</b>	
** = p < 0.005	
* = p < 0.05	
n = 8, 7, 7, 6, 8, 8	

## pLK30 5085 br pLK18 4969 bp Tag tmcr yfast (2757) ApaI (2741) BamH

Figure 10. The primary plasmids used for this project. pLK30 expresses YFAST under Pmcr to test native expression levels from a highly expressed promoter. pLK27 expresses the RNA Polymerase within *M. acetivorans* and uses it to transcribe YFAST under Prnap. pLK18 has no promoter to act as a promoterless control. pLK28 adds elements needed to establish a repressed system to pLK27 to allow for inducible control of expression from the recombinant RNA Polymerase

- Compare protein expression levels between strains with YFAST under control of *Pmcr* versus *Prnap* using fluorescence assays.
- Detect YFAST expression using Western Blots.
- regulation in Methanosarcina acetivorans be better understood.
- Further explore tools to help transcription, translation, and their • Establish inducible expression system for tighter control over gene
- expression.
- Develop fully orthologous gene circuits in *M. acetivorans.*

# Acknowledgements

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### Prnap-YFAST, Pmcr-RNA Polymerase PmcrB-repressor, permease





### **Future Directions**

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