

DOE Bioenergy Technologies Office (BETO) 2019 Project Peer Review

Engineered reversal of the β -oxidation cycle in clostridia for the synthesis of fuels and chemicals

March 5, 2019
Biochemical Conversion

Michael C. Jewett, Northwestern University (PI)
Michael Köpke, LanzaTech (Co-PI, Presenter)



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Quad Chart Overview

Timeline

- Project start date - 10/1/2018
- Project end date - 9/30/2021
- Project complete 7% (1/14 milestones)

	Total Costs Pre FY17*	FY 17 Costs	FY 18 Costs	Total Planned Funding (FY 19-Project End Date)
DOE Funded	\$0	\$0	\$0	\$1,600,000
Project Cost Share*	\$0	\$0	\$0	\$400,000

•**Partners:** Northwestern University (34%), LanzaTech (33%), University of South Florida (25%), Georgia Institute of Technology (8%)

Barriers addressed

- Ct-D. Advanced Bioprocess Development
- Ct-H. Gas Fermentation Development
- Ct-L. Decreasing Development Time for Industrially Relevant Microorganisms
- Ct-N. Multiscale Computational Framework toward Accelerating Technology Development
- Ot-A. Availability of Quality Feedstock

Objective

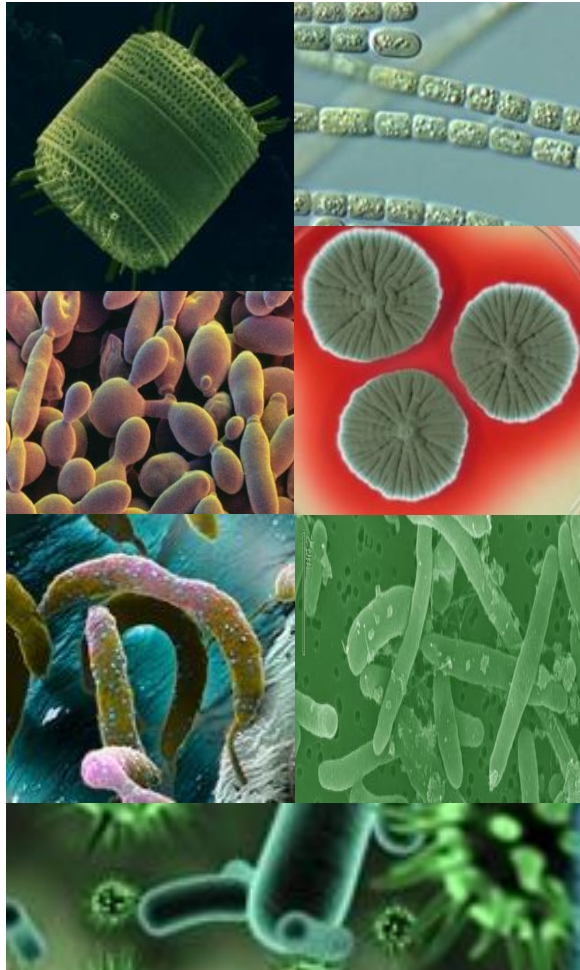
- Our project objective is to develop clostridia to ferment synthesis gas into a range of advanced bioproducts.

End of Project Goal

- We will manufacture one product from engineering a reversal of the β -oxidation cycle in clostridia at a metric of $>0.1\text{g/l/h}$ in $>80\text{L}$ scalable pilot reactor.
- We will assess environmental, community and rural economic development impacts




1 - Project Overview

The need for low-cost biofuels and bioproducts from sustainable resources is intensifying



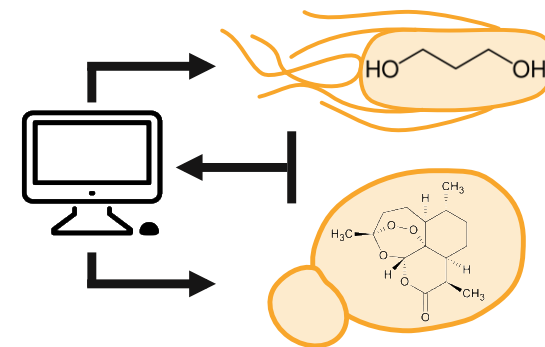
Unfortunately, designing, building, and optimizing biosynthetic pathways in cells remains a complex and formidable challenge

- Development cycles to optimize biosynthetic pathways can be slow, especially for non-model organisms

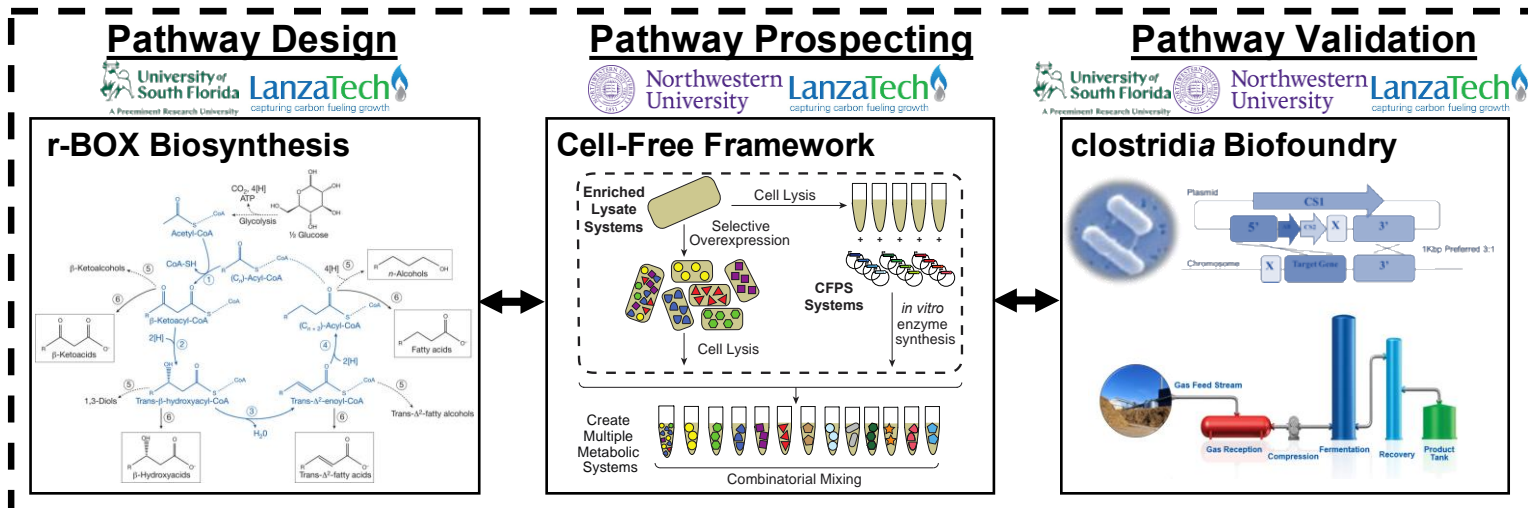
Molecule	Institutions	Time	Person Years	Cost
1,3-Propanediol 	DuPont, Genencor, Tate & Lyle	15 years (1992-2007)	>550	>\$130M
Artemisinin 	UC Berkeley, Amyris, Sanofi	13 years (2000-2013)	>130	>\$50M
Farnesene 	Amyris, TOTAL	4 years (2008-2012)	>250	>\$30M

Nielsen J & Keasling JD, *Cell* (2016) DOI: 10.1016/j.cell.2016.02.004; Karim AS, Dudley QM & Jewett MC *Industrial Biotechnology* (2017) DOI: 10.1002/9783527807796.ch4

- Platform organisms, accessible feedstocks, target molecules, and stable environments in which to work are limited
- Integrated computational frameworks for biodesign need improvement



Our project objective is to develop clostridia to ferment synthesis gas produced from cellulosic biomass by established gasification technologies, into a range of advanced bioproducts



Rural Economic Development & Sustainability Analysis



We will target products used as **drop-in fuels, fuel additives, and chemical building blocks** with a **\$14Bn US market**.

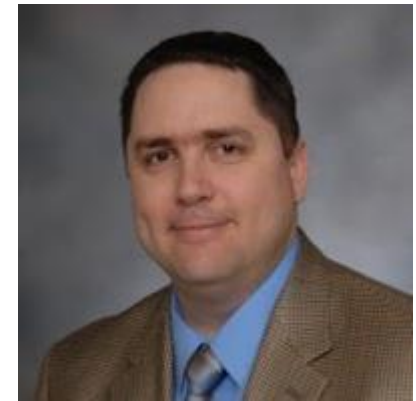
2 – Approach (Management)



Northwestern:
Michael Jewett
Bioengineering



LanzaTech:
Michael Koepke
Industrial Biotech



Univ. South Florida:
Ramon Gonzalez
Chemical Engineering



LanzaTech:
Robert Conrado
Technoeconomic Analysis



GaTech:
Valerie Thomas
Technology Assessment

Several activities ensure efficient coordination within the team and reporting to DOE/USDA

- Northwestern and LanzaTech have **collaborated on numerous projects since 2015** and have developed successful mechanisms for technical coordination, data sharing, and integration
- Bi-weekly meetings to review the team's progress and discuss any matters requiring action
- Project structure enables multiple, parallel paths to achieve 4 tasks and the milestones
 - **Task 1.** Develop and apply informatics and computer-aided design tools to choose molecules, enzymes, and pathways in clostridia (**USF, LT**)
 - **Task 2.** Establish a cell-free framework for rapid pathway prototyping and analysis (**Northwestern, LT**)
 - **Task 3.** Develop optimized production strains of clostridia (**LanzaTech, NU, USF, GT**)
 - **Task 4.** Rural economic development and sustainability analysis (**GaTech, LT**)
- Interface with other DOE projects to accelerate goals
 - **Complement cBioFab project, utilize DOE user facility JGI for gene synthesis**

2 – Approach (Technical)

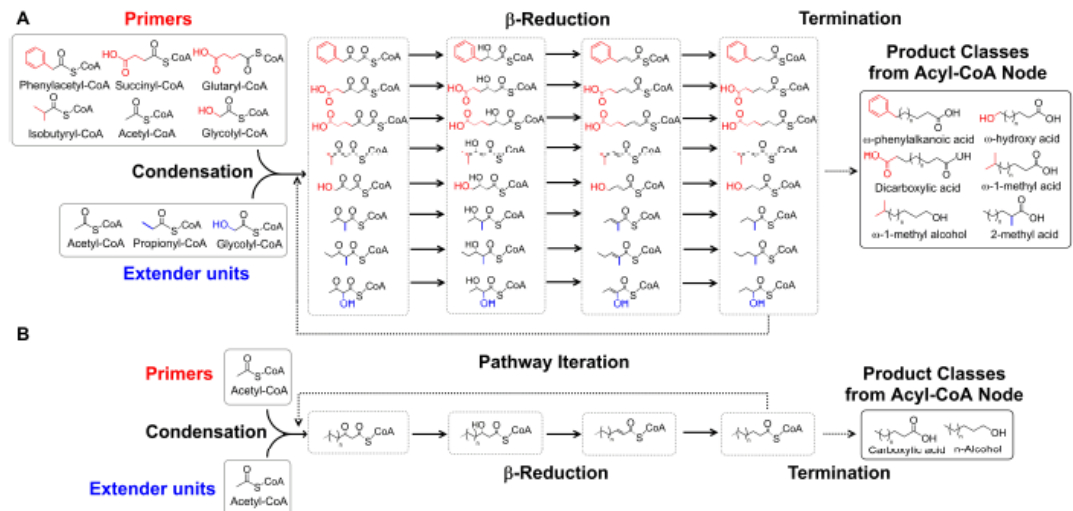
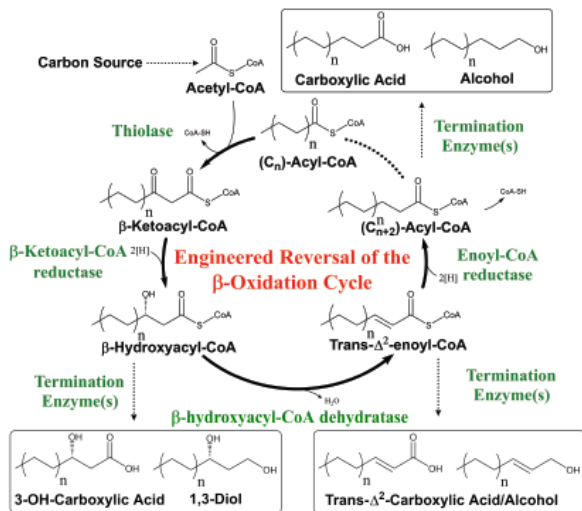
To achieve our vision, we will:

- **Aim 1.** *Develop and apply informatics and **computer-aided design tools** to choose molecules, enzymes, and pathways for reverse β -oxidation cycle (r-BOX) in clostridia.*
- **Aim 2.** *Establish a **cell-free framework** for rapid pathway prototyping and analysis*
- **Aim 3.** *Develop optimized production strains of **gas-fermenting clostridia**.*
- **Aim 4.** *Technoeconomic and rural economic development and sustainability analysis.*

Embedded in these aims, are several key innovations that will allow us to combine *in vitro* (cell-free) and *in vivo* work to interweave and advance state-of-the-art pathway design, prospecting, validation, and production in an integrated framework

Reversal of the β -oxidation cycle (rBOX) offers unique access to thousands of molecules

- Most bioengineering approaches to date rely on linear pathways specifically designed for a single molecule.
- The cyclic and iterative r-BOX pathway with **highly modular architecture and combinatorial nature** offer the unique advantage of providing **access to thousands of molecules** with different chemistries and chain lengths.



Dellomonaco C, Clomburg JM, Gonzalez R, *Nature*(2011) DOI: 10.1038/nature10333

Cheong S, Clomburg JM, Gonzalez R, *Nature Biotech* (2016) DOI: 10.1038/nbt.3505

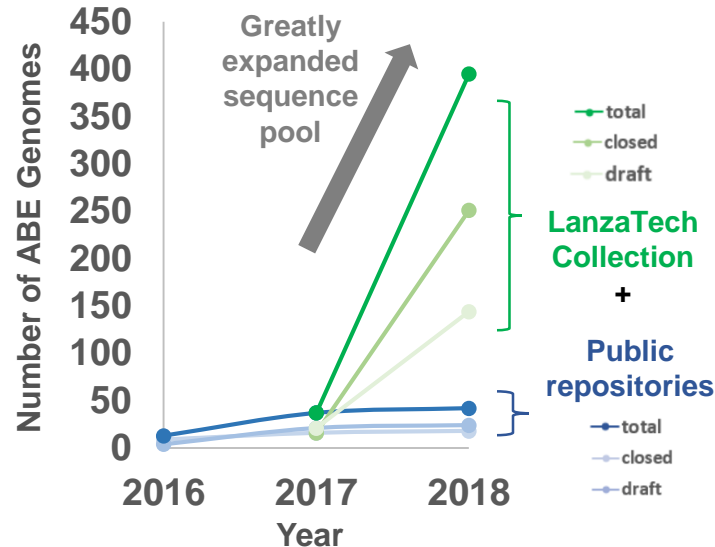
Largest collection of industrial *Clostridium* strains provides unique access to highly-evolved gene variants

- LanzaTech owns the largest collection of industrially-deployed acetone-butanol-ethanol (ABE) clostridia strains
- The collection dates back to 1944 and contains hundreds of highly evolved strains, **spanning several decades of development at several large-scale commercial plants**
- In collaboration with JGI, the entire collection have been sequenced



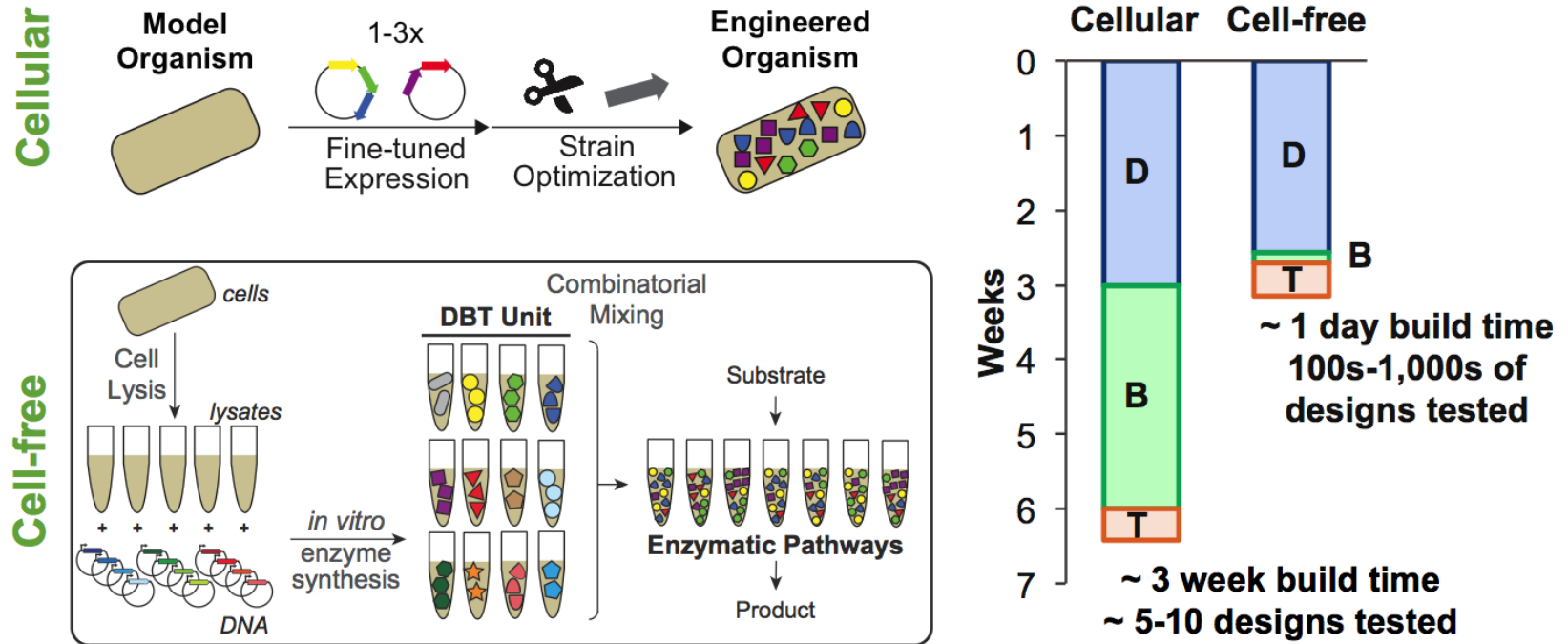
NCP commercial ABE plant
1935-1982

Jones DT and Woods DR, Microbiol Rev (1986).



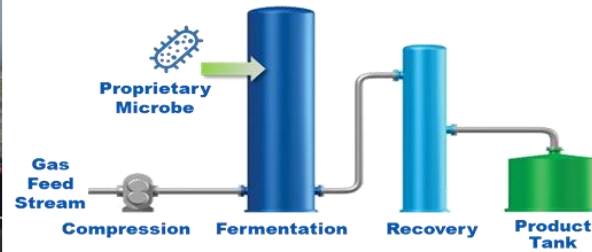
Cell-free Framework allows for rapid pathway prototyping

- Cell-free systems enable **acceleration of new enzyme and pathway assessments**, bypassing transformation idiosyncrasies and low-throughput workflows impeding progress on many non-model microorganisms



Clostridia gas fermentation allows high yield conversion of lignocellulosic feedstocks

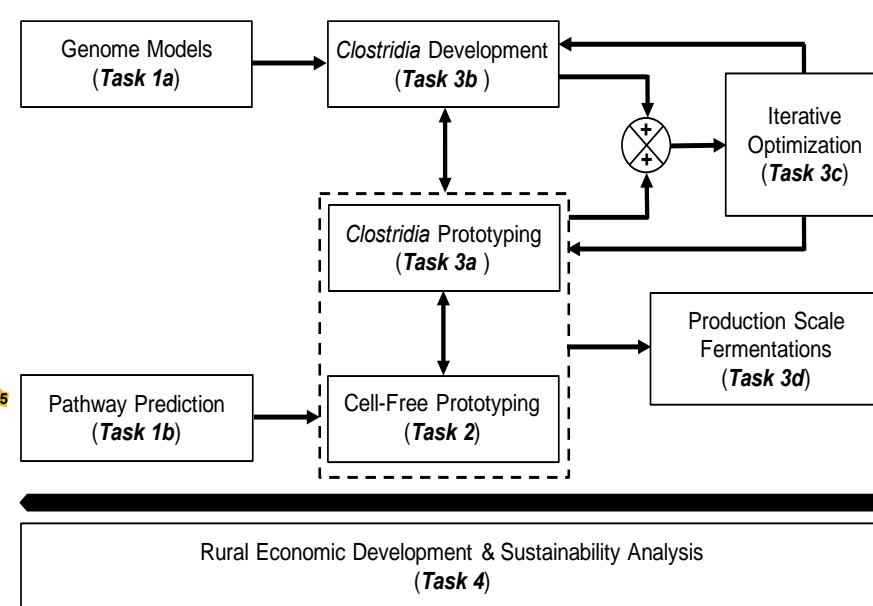
- A hybrid thermochemical (gasification) - biological (gas fermentation) pathway utilizes all biomass components (lignin and cellulose), maximizing yields and overcoming barriers such as biomass recalcitrance.
- Integrated gasification-fermentation has been demonstrated in extended continuous operations using multiple types of lignocellulosic material.
- Syngas fermentation uses the same fermentation process implemented in LanzaTech's first commercial scale gas fermentation facility (started May 3 2018).



How are we going to achieve our goals?

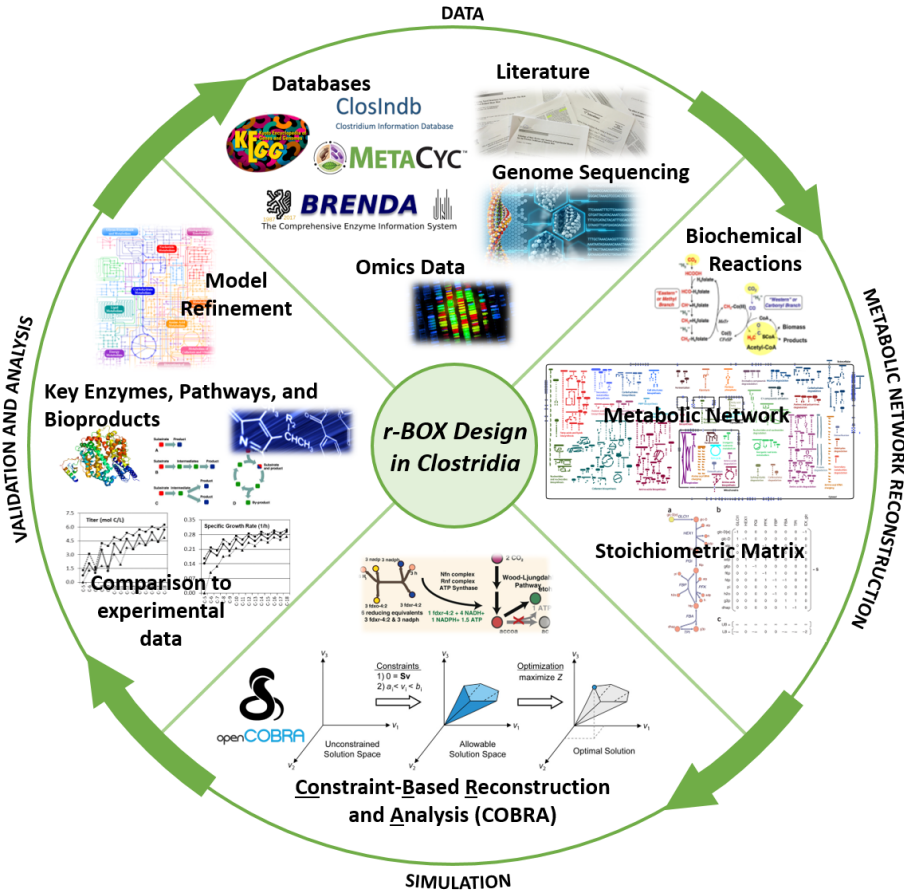
		Phase 1		Ph 2
		Y1	Y2	Y3
Task 1	Develop and apply informatics and computer-aided design tools to choose molecules, enzymes, and pathways for r-BOX in clostridia			
	1a: Determine optimal r-BOX designs in clostridia supporting maximum bioproduct synthesis	1.1	1.2	
	1b: Identify candidate r-BOX enzymes facilitating optimal r-BOX designs		1.3	
Task 2	Establish a cell-free framework for rapid pathway prototyping and analysis			
	2: Demonstrate methods for systematic pathway optimization using cell-free cocktails.	2.1	2.2	2.3
Task 3	Develop optimized production strains using a clostridia biofoundry that leverages cell-free systems			
	3a: Testing r-BOX pathway variants informed by cell-free prototyping in clostridia	3.1	3.2	3.3
	3b: Genome-scale model (GEM) predictions to optimize flux and inform cell-free work.			3.4
	3c: Iterative pathway and strain optimization			3.5
	3d: Process development and scale-up			
Task 4	Rural economic development and sustainability analysis			
	4a: Technoeconomic analysis (TEA) and Life cycle analysis (LCA)	4.1	4.2	
	4b: Rural economic development analysis			4.3

Legend	NU Northwestern University: (Jewett)	GT Georgia Tech (Thomas)
★ -Milestones	LT LanzaTech (Köpke, Conrado)	
..... -Interactions between tasks	RU Rice University (Gonzalez)	★ -Go/No-go decision



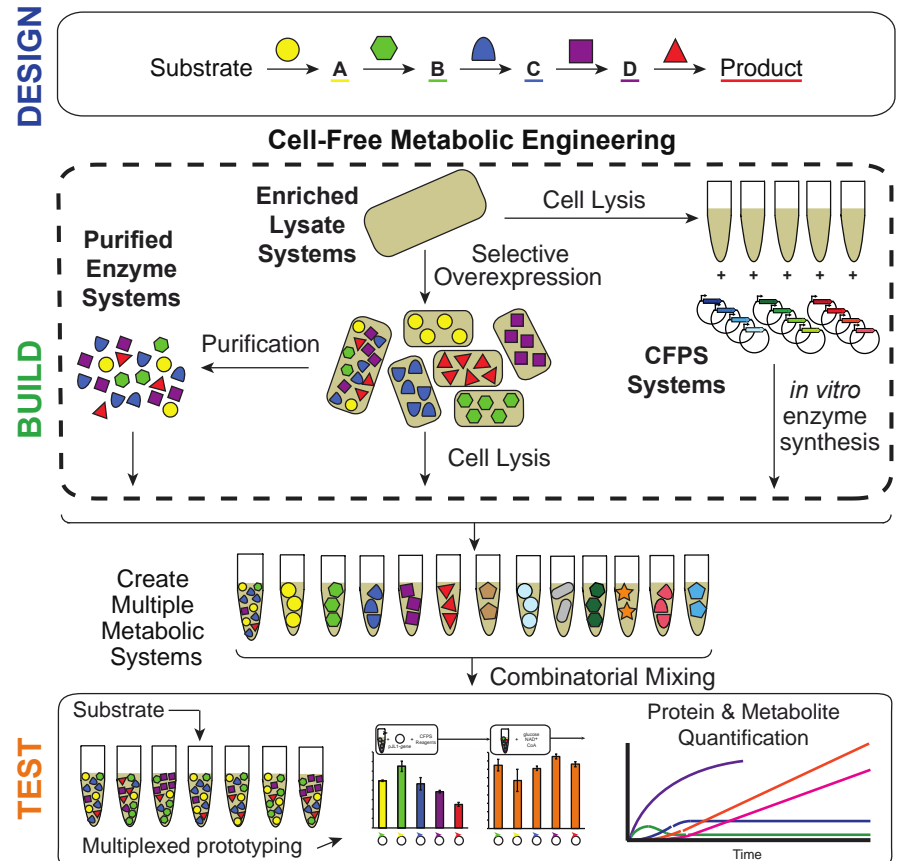
Task 1: Use pathway design to provide a focused set of r-BOX input molecules

- Milestone 1.1.** Identify and compare r-BOX enzyme variants by data-mining complete *Clostridium* collection and public databases: **>100 additional unique r-BOX and Ptb-Buk putative gene variants over the public domain identified (Y1/Q4)**
- Milestone 1.2.** Quantify theoretical product yields and **generate optimal strain designs with > 100,000 simulations** carried out per design (Y2/Q2)
- Milestone 1.3.** **Optimize computational framework** for generating novel pathways based on feedback from other Tasks and refine pathway design (Y3/Q2)



Task 2: Develop methods for accelerating systematic pathway optimization using cell-free cocktails

- **Milestone 2.1.** Develop, implement, and demonstrate methods for the **cell-free mix-and-match approach to optimize biosynthetic pathways** that are **2x faster than the state of the art in vivo** approach (Y2/Q2).
- **Milestone 2.2.** Demonstrate **expression of pathway enzymes** for at least one r-BOX pathway at levels of **greater than 50 $\mu\text{g/mL}$** using the cell-free framework (Y2/Q3).
- **Milestone 2.3.** Study and **optimize pathways using our cell-free framework**, and refine and optimize pathways with **at least 2-fold improvement** (Y3/Q2).



Task 3: Develop optimized strains for production of r-BOX products from biomass syngas

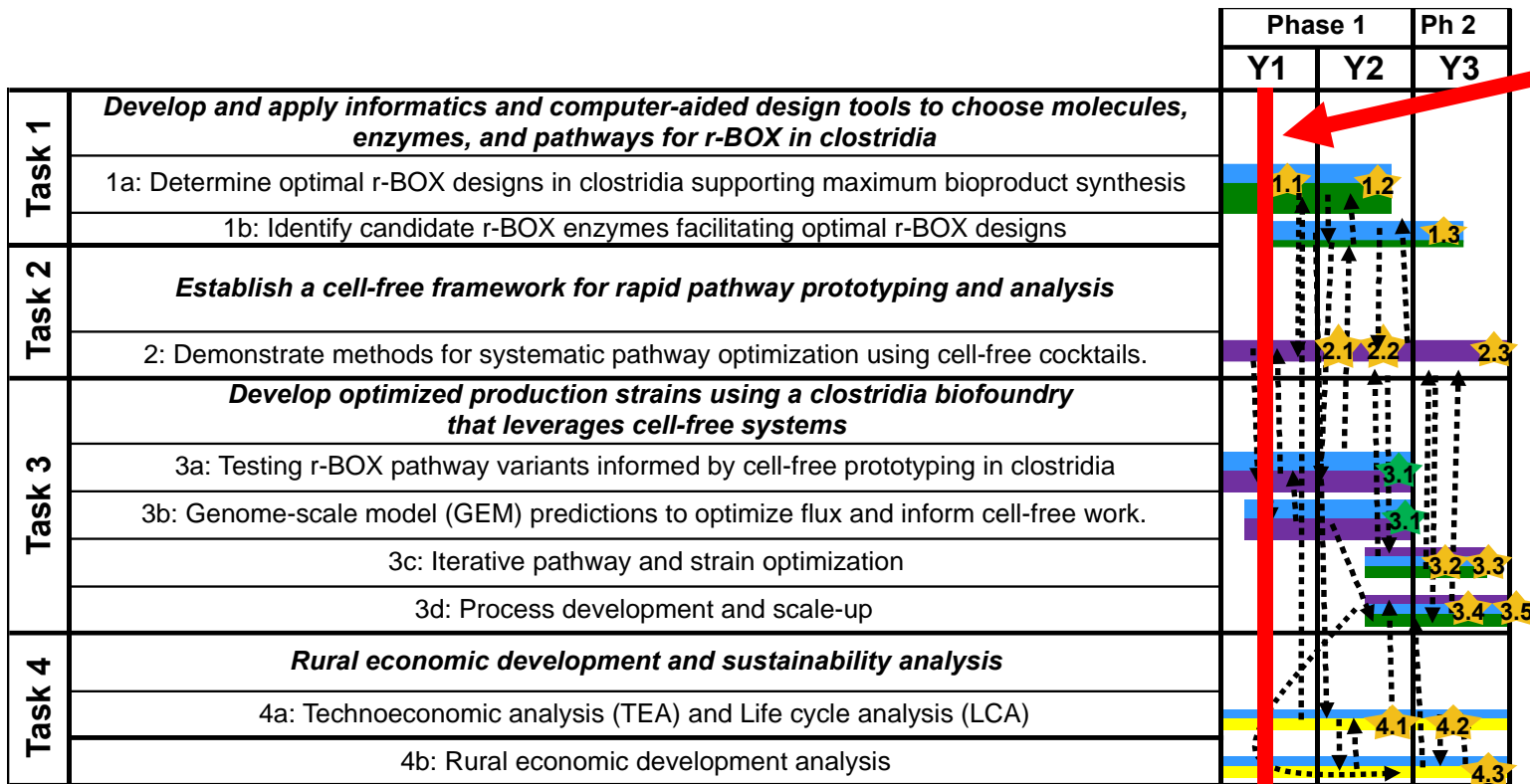
- **Milestone 3.1.** (go/no-go decision): Construct and evaluate **50 unique pathway designs** for target molecules *in vivo* and *in vitro*. Our **metric is 100mg/L/d** of one target product. (Y2/Q4)
- **Milestone 3.2.** Proof of concept for **additional rBOX target products** on syngas. (Y3/Q2)
- **Milestone 3.3.** Construct and evaluate an **additional 150 unique pathway designs** for target molecules *in vivo* and *in vitro*. (Y3/Q3)
- **Milestone 3.4.** Comparison of best performing engineered strain for on **synthetic syngas against real biomass syngas** in 1.5L lab scale reactor and demonstration of a target **metric of >0.1g/l/h**. (Y3/Q3)
- **Milestone 3.5.** One selected rBOX product at a target **metric of >0.1g/l/h** in **>80L scalable pilot reactor**. (Y3/Q4)



Task 4: Assess environmental, community, and rural economic development impacts of r-BOX synthesis to facilitate design

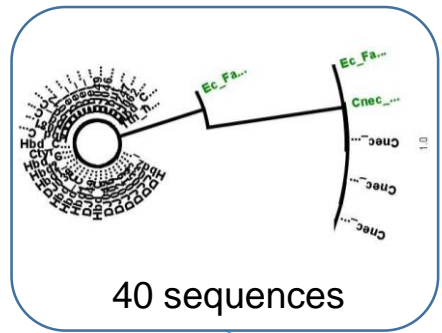
- **Milestone 4.1.** Complete **2 workshops to inform lifecycle impact analysis** using TRACI model. All stakeholders will be invited to both workshops; aims are to gather input from multiple parties on potential economic, community, and environmental impacts. (Y2/Q4)
- **Milestone 4.1.** Completed LCA for two rBOX molecules. (Y3/Q2)
- **Milestone 4.1.** Completed assessment of infrastructure and supply chains for biomass feedstock supply of two rBOX molecules in the US southeast. (Y3/Q4)

3 – Technical Accomplishments/ Progress/Results

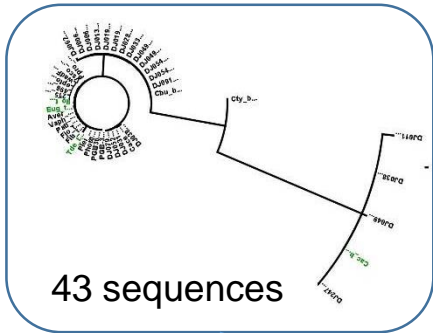


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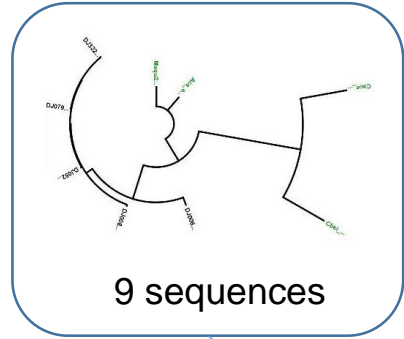
Task 1: Based on initial enzyme set, mined enzyme candidates for r-BOX in *Clostridium* collection



40 sequences

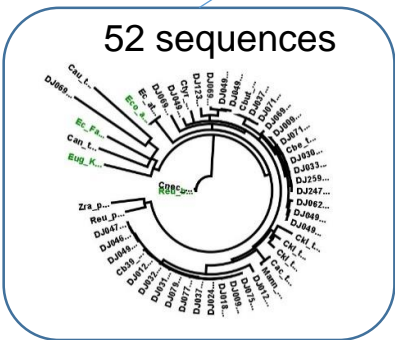
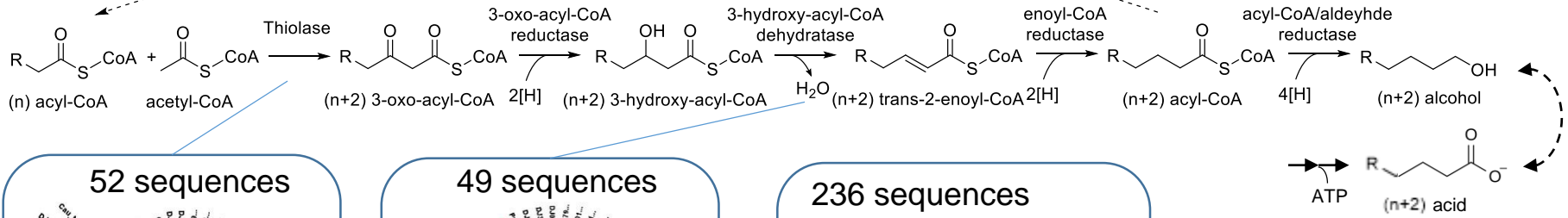


43 sequences

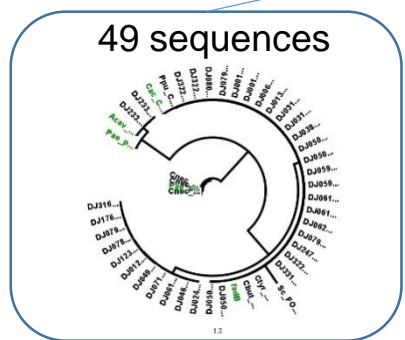


9 sequences

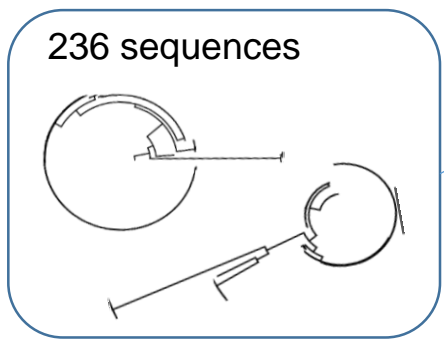
Pathway iteration



52 sequences



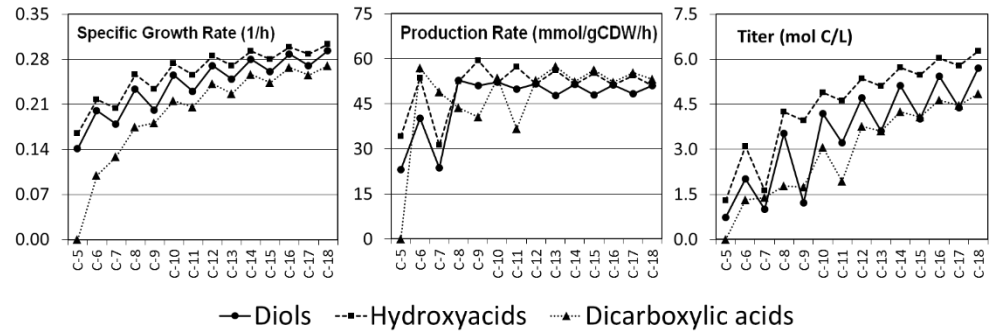
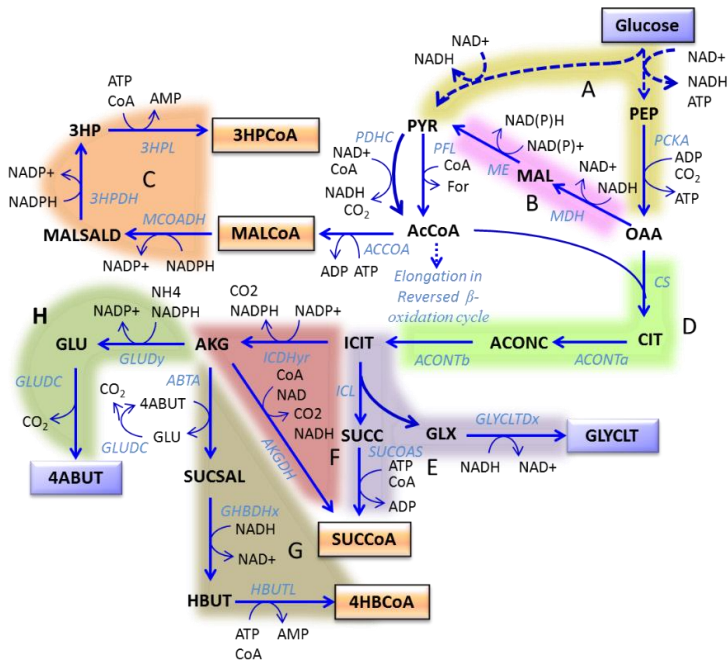
49 sequences



236 sequences

More than 200 unique rBOX gene variants identified that have been codon adapted and are currently being synthesized through the JGI DNA synthesis program

Task 1: Genome-scale modeling (GEM) and metabolic flux balance analysis to define optimal r-BOX inputs

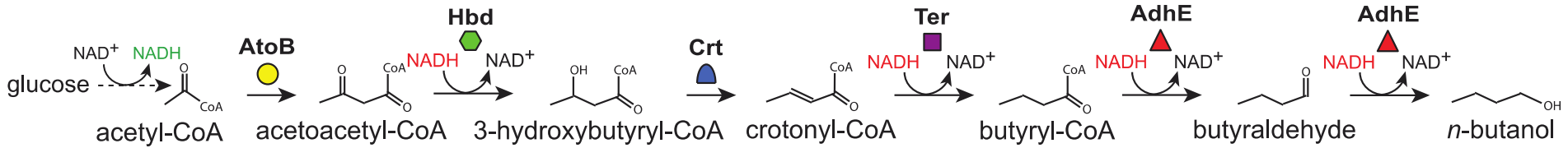
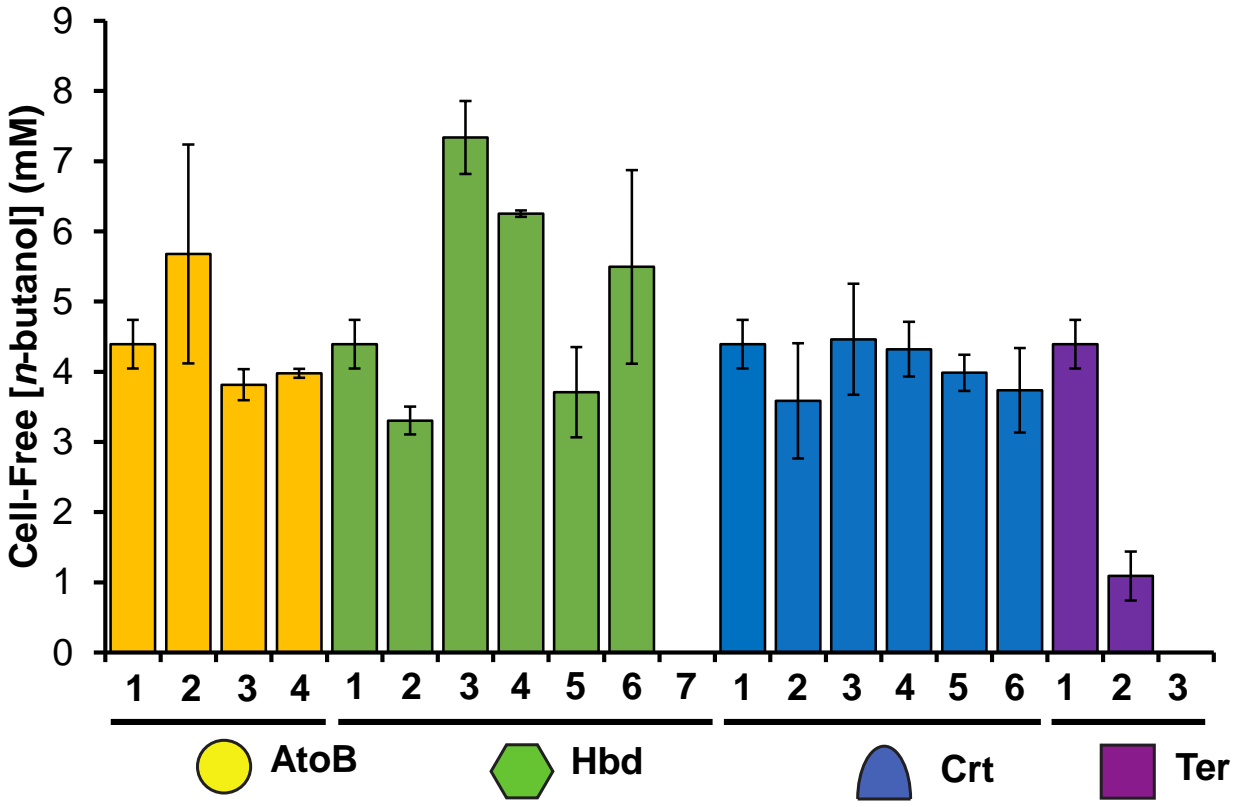
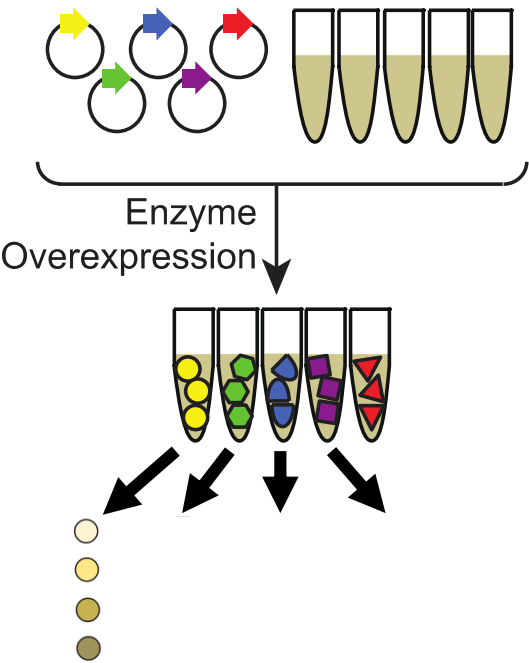


- *E. coli* GEM has previously been exploited to determine optimal routes for omega-functionalized product synthesis and input molecules (left) to maximize target product synthesis from glucose (right)

Reactions from *E. coli* model have been integrated to *Clostridium* GEM to enable modeling of target product synthesis in desired genetic background

Task 2: Developed cell-free platform to test homology activity and beginning to apply it to r-BOX

(example for butanol pathway below)



Task 4: We plan two workshops to guide the project's strategic direction and product emphasis

- Workshop 1. Atlanta, October 2019. Invitees will include Georgia Rural Economic Development Authority representatives, industry representatives, environmental NGO representatives, experts on community impacts, ethanol manufacturers
- Workshop 2. Soperton area, April 2019. Invitees will include: Treutlen County Development Authority, Heart of Georgia Altamaha Regional Commission, Site Tour

4 – Relevance

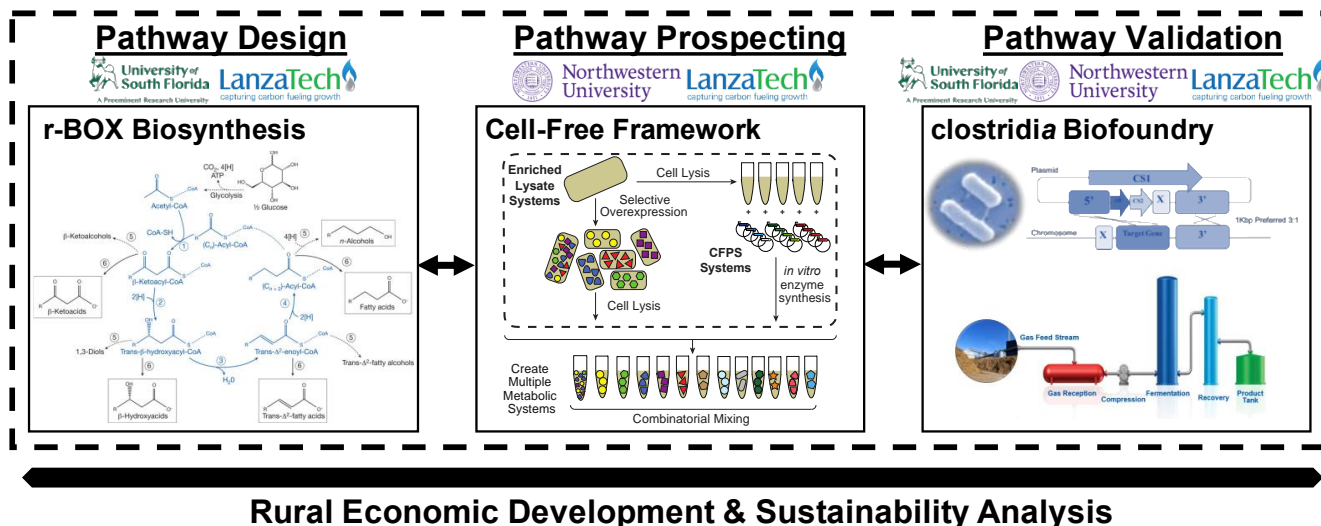
- Enable high-level, sustainable synthesis of next-generation biofuels and bioproducts by **developing clostridia to ferment synthesis gas**
 - Advance syngas fermentation, a cost-effective technology for the use of cellulosic biomass that is broadly applicable in the production of biofuels and biobased products (Ct-H)
 - Expand the diversity of products that can be produced and co-produced via syngas fermentation (Ct-H)
 - Gain new knowledge of metabolism in obligate anaerobes and traits related to “industrial fitness” of clostridia (Ct-H, Ct-D)
- Create a cell-free framework to **decrease development time for industrially relevant microorganisms**
 - Advance bioprocess development by reducing the time to new biosynthetic pathways (Ct-D, Ct-L, Ct-N)
 - Provide a key case study for the bioenergy industry by establishing r-BOX for production of advanced biofuels and value-added chemicals
- Expand the scope of biomanufacturing practice, **enabling regional and global economic growth**
 - Develop rural economic and sustainability analysis frameworks to guide product selection
 - Accelerate commercialization of new gas fermentation products from lignocellulosic biomass, with specific application to forestry residues in the Southeast (At-A)
 - Demonstrate pilot scale synthesis of one r-BOX product (Ct-H, Ct-D)

5 – Future Work

- **Key upcoming milestones (18 months):**
 - Use genome-scale metabolic models for r-BOX pathway design
 - Establish a cell-free framework for rapid pathway prototyping of r-BOX pathways to accelerate design in clostridia
 - Complete workshops to inform life cycle analysis
- **Go/No-Go Point (9/30/2020):**
 - Construct and evaluate 50 unique pathway designs for target molecules *in vivo* and *in vitro* to enable 100mg/L/d of one r-BOX product in clostridia
 - *Already created compatible vector system to facilitate evaluation*
- **Project completion steps**
 - Demonstrate synthesis of one r-BOX product in clostridia at a metric of >0.1g/l/h in >80L scalable pilot reactor.
 - Complete rural economic development & sustainability analyses

Summary

- We will develop a new technology platform for engineering r-BOX in clostridia for fuel and chemical synthesis from biomass syngas
- Our approach to integrate pathway design, pathway prospecting, pathway validation, and economic analysis will **enable a new paradigm of biological design.**



- Our current project progress is **ahead of schedule.**
- The impact will be to **reduce the time required to develop new biosynthetic products** and **expand the scope of biomanufacturing practice through gas fermentation**

Thank you!

U.S. DEPARTMENT OF
ENERGY

Energy Efficiency &
Renewable Energy

BIOENERGY TECHNOLOGIES OFFICE

Technology Manager: Jay Fitzgerald

Project Monitor: Clayton Rohman

Grants Management Specialist: Nicholas Oscarsson