



U.S. ARMY RESEARCH, DEVELOPMENT AND ENGINEERING COMMAND

Microbial Reactors - Indigenous Feed Stocks to Functional Materials

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Team Lead, SBME ARAP Technical Lead

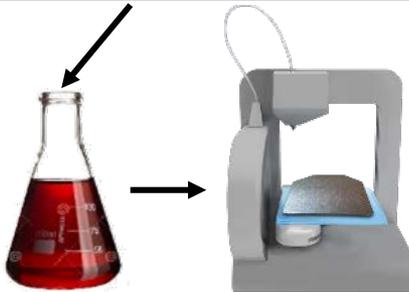
Biotechnology Branch



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ARMY CHALLENGE

To win in the deep future operational environment, the Army will need to be more adaptive and with less logistic demand.



- Cost/benefit specialty/commodity materials is different than commercial sector
- Synthesis from indigenous resources/waste streams enables:
 - cost reduction
 - minimize materials burden
 - Benefits from waste remediation
- Requires development of biological systems and production pathways
 - robust in Army environments.
 - Process complex inputs
 - High value products/precise hierarchal materials

- 1. A component of ARL Essential Research Programs: Tactical unit energy independence, Manufacturing at the point of need, Discovery.**
- 2. Enables CSA Priority #6 Soldier Lethality: Expeditionary Soldier Power Generation**



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UNCLASSIFIED

TECHNOLOGIES FOR FUTURE DISMOUNTED SOLDIER



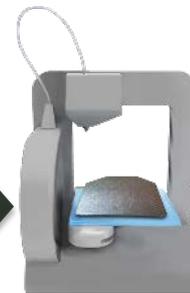
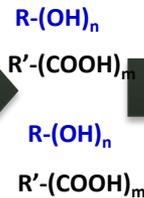
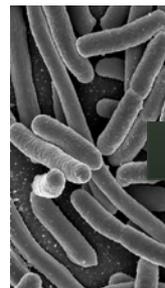
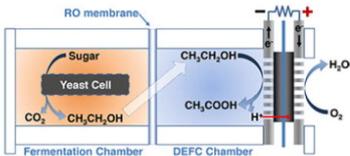
Cumulative, Connective,
Converging w/ Outcomes



Soldier System Interfaces & Integration / Sensored Soldier / Expeditionary Power Generation: Self-Sustaining Autonomous System

Engineered Organism: Flexible feedstocks to controlled reactant output for advanced polymer systems and sensor reagents

Selected Consortia Bioreactor: Conversion of indigenous materials/ wastes for energy generation at expeditionary scale

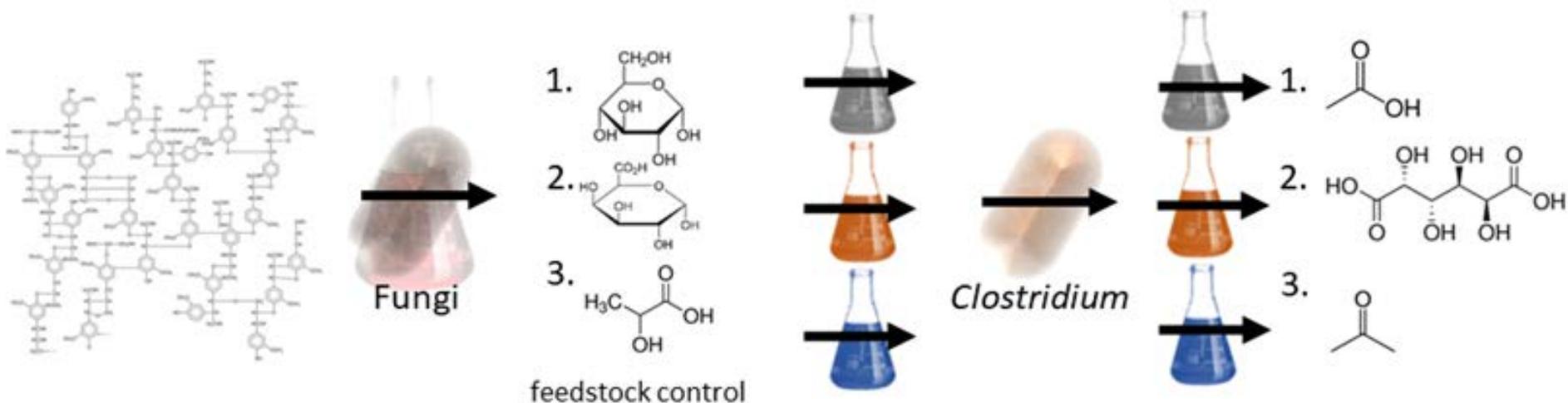




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Technical Challenge

Designed Consortia



Microbial consortia synergistic production of materials and remediation

Organics, Biosensor Reagents,
Probiotics, Enzymes / Catalysts, Adhesives,
Polymer Precursors
Energetics Precursors, Solvents

Nano Particles
Rare earths reclamation

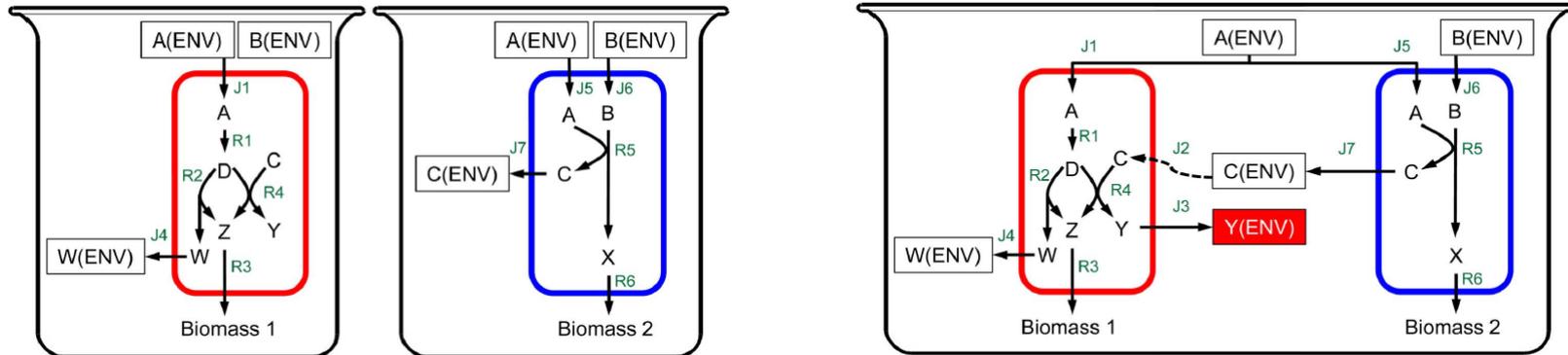
- **Develop production pathways :**
 - **robust in Army environments**
 - **reconfigurable**



From Monocultures to Consortia

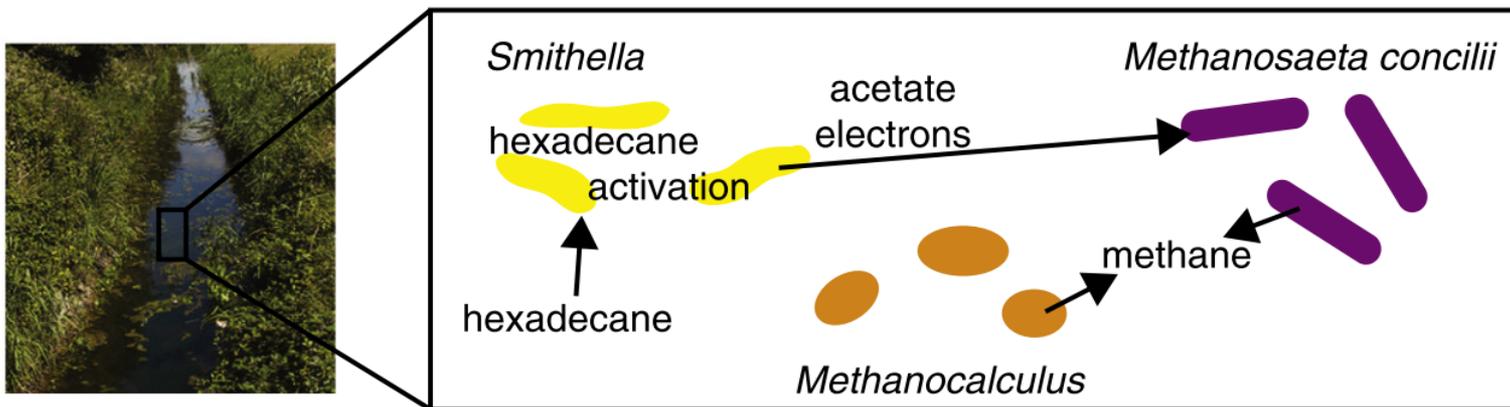
Hypothesis: Microbial consortia will outperform monocultures for waste and indigenous resource conversion to commodity/specialty chemicals

Theory:



Chiu s et al. PLoS Comput Biol. 2014.

Nature:



Hays et al. Current Opin Biotechnol. 2015.



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Bottom-up Assembly of Consortia

Food waste



Indigenous resources
(e.g. grass)



Commodity / Specialty Chemicals

- Leverage advancements in human microbiome research in new context.
- Plethora of genome-scale metabolic models
- Transitions to Natick Soldier Research and Development Center
- Partnership with John La Scala (WMRD)

- Leverage rumen microbiome that degrade lignocellulose
- Leverage past ARL research on *Clostridium acetobutylicum*
- Untapped potential, particularly lignocellulolytic fungi – collaboration with Michelle O'Malley at UCSB



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Non-homogenous
food waste



Tunable outputs

Key questions:

Hundreds of species, so how do we choose members for consortia?

Do combinations of microbes outperform monocultures?

Approach: Flux balance analysis (FBA) of genome-scale metabolic models



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Genome-scale Metabolic Modeling

- Modeling predicts potential of microbes to cooperate/compete
- Plethora of available models:
 - Assembly of gut organism
 - reconstruction and analysis (AGORA)
 - **773 microbes**
 - Includes 205 genera and 605 species
 - Simulate steady-state growth on Western diet

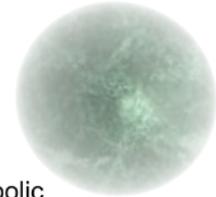
Magnusdottir *et al.* *Nat Biotechnol.* 2016.

- **Simulated monocultures and all co-cultures**
 - 298,378 co-cultures
 - Computation time = 7 days
 - Would be a major experimental challenge
 - 96 well plate format – 3,109 plates
 - With automation ~ 50 plates/day
 - 62 days to setup
 - Metabolite analysis: 10 min/sample
 - ~ 5.5 years total

Genome 1



Genome 2



Metabolic reconstruction based on enzyme annotations

Stoichiometric matrix, S_1

Metabolites	Reactions				Flux vector, v_1
	1	2	3	4	
A	-1	-1	0	1	v_A
B	-1	0	0	0	v_B
C	1	-1	0	0	v_C
D	0	0	0	1	v_D

X

S_2

Metabolites	Reactions				Flux vector, v_2
	1	2	3	4	
A	1	-1	0	1	v_A
B	-1	0	0	0	v_B
C	1	1	0	0	v_C
D	0	0	0	1	v_D

X

Example Rxn 1: $A + B \leftrightarrow C$

FBA
 $S_1 \times v_1 = 0$
Optimize biomass
rxn

Essential nutrient inputs for growth

Metabolic outputs

FBA
 $S_2 \times v_2 = 0$
Optimize biomass
rxn

Essential nutrient inputs for growth

Metabolic outputs

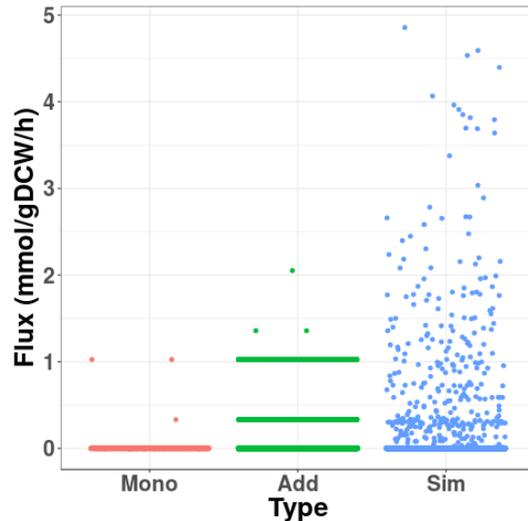
Competition
Cooperation



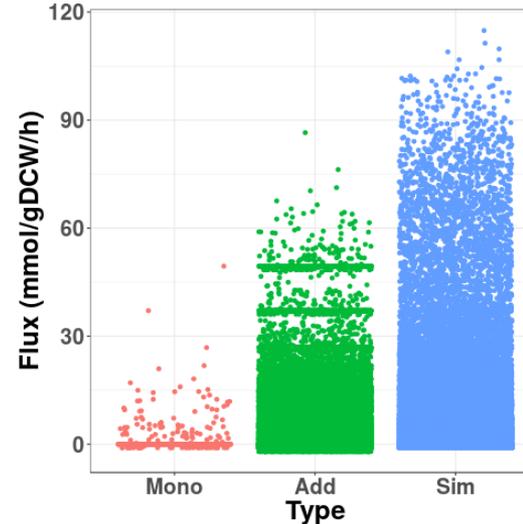
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Co-cultures Outperform Monocultures

Butanol



H₂

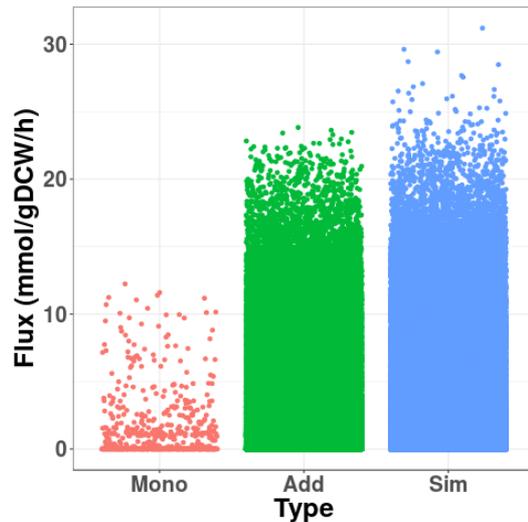


Mono: all monoculture fluxes

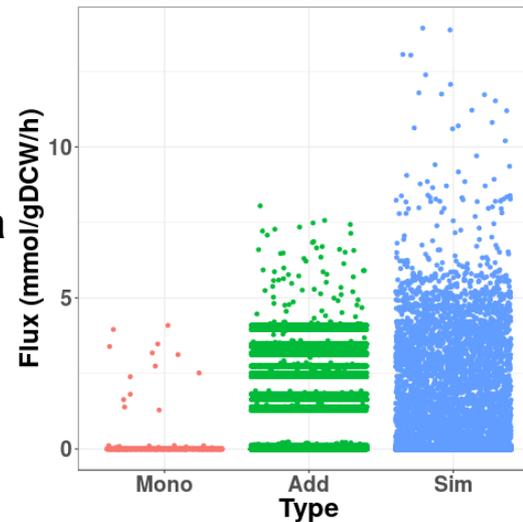
Add: All pairwise monoculture fluxes added together

Sim: All pairwise simulated co-culture fluxes

Propionate



Urea



Models: co-cultures capable of greater than additive production



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Microbe production of Useful Chemicals

Products based on single microbe model simulations:

<u>Organics</u>	<u>Alcohols</u>	<u>Nitrogen</u>	<u>Fatty acids</u>	<u>Gases</u>
Acetate	Ethanol	Ammonium	Tetradecanoate	H ₂
Formate	Butanol	Amino acids	Octadecanoate	N ₂
Succinate	Methanol	Indole	Hexadecanoate	CO ₂
Propionate	Propylene glycol	Urea		CH ₄
Butyrate				
Lactate				
Glycolate				
Pyruvate				

Emergent products from co-culture simulations:

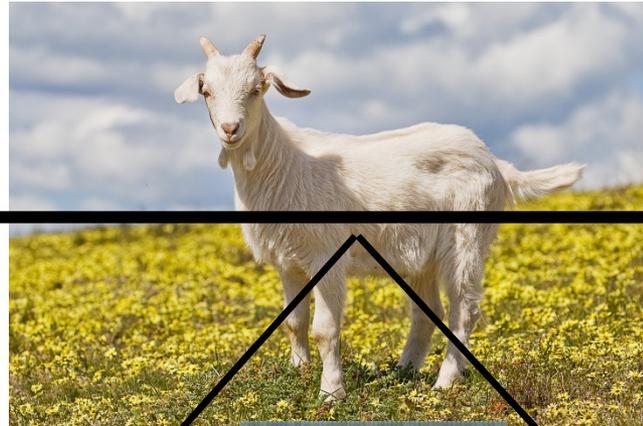
Citrate	2,3-butanediol	Vitamin B ₁	Laurate	N ₂ O
Diacetyl		Vitamin B ₂		
Oxaloacetate		Vitamin B ₉		

**Di-carboxylic acids and alcohols have potential for additive manufacturing
(John La Scala - WMRD)**



RUMEN MICROBIOME AS GUIDE

Non-homogenous
lignocellulose
feedstock

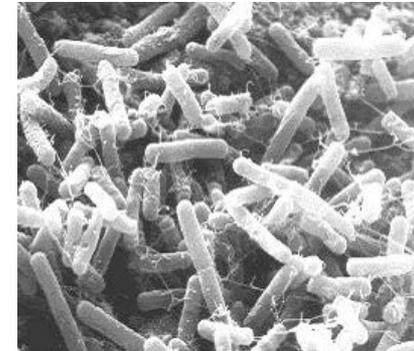
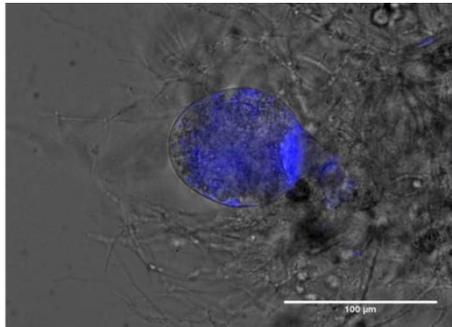


→ Tunable outputs



Anaerobic, lignocellulolytic fungi

Clostridium acetobutylicum



Haitjema et al. *Biotechnol Bioeng.* 2014.



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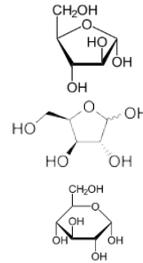
Functional Ecological Approach

Ecological strategies: Division of labor, temporal organization

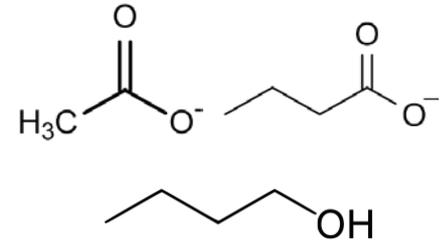
2 Stage

Fungi
(21 days)

Cac
(10 days)



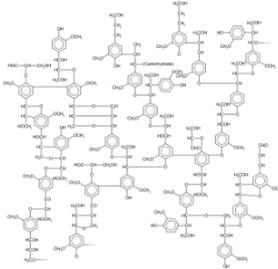
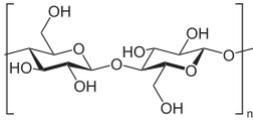
Useful products



Fungi + *Cac*
(31 days)

1 Stage

Reed canary grass



Fungi: *Neocallimastix californiae* *Cac*: *Clostridium acetobutylicum*

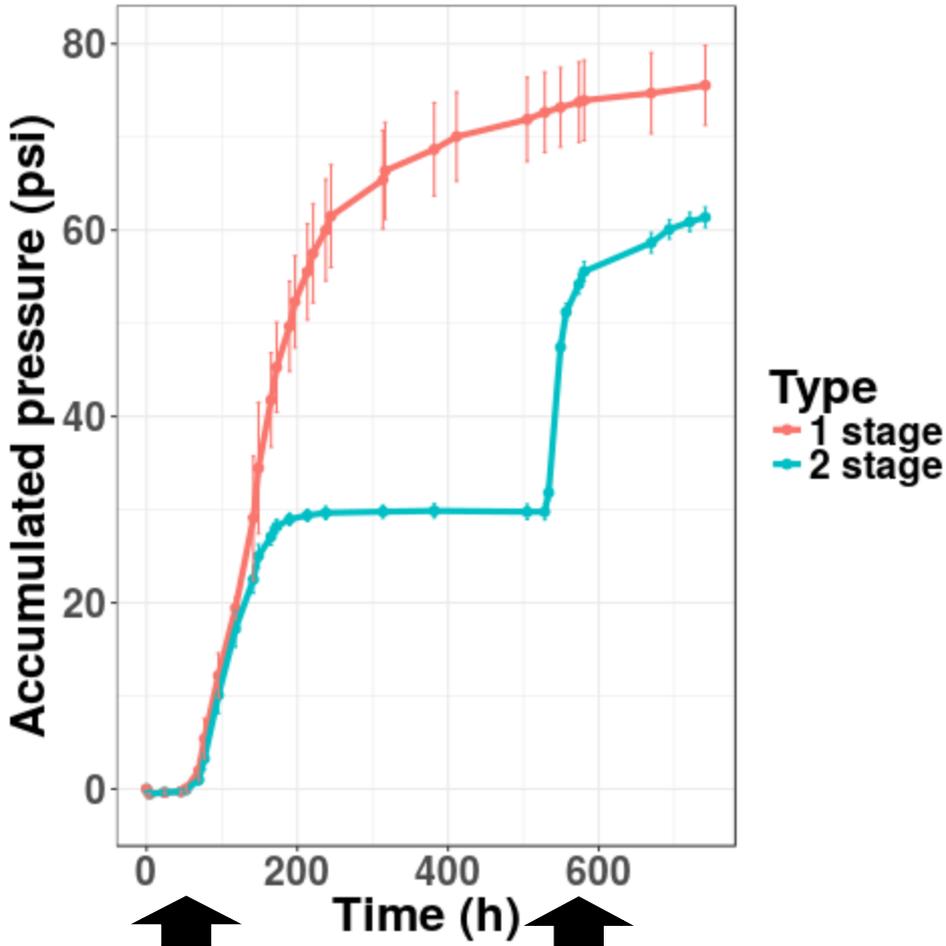


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Fungi/Cac Co-cultures Result In Synergistic Growth

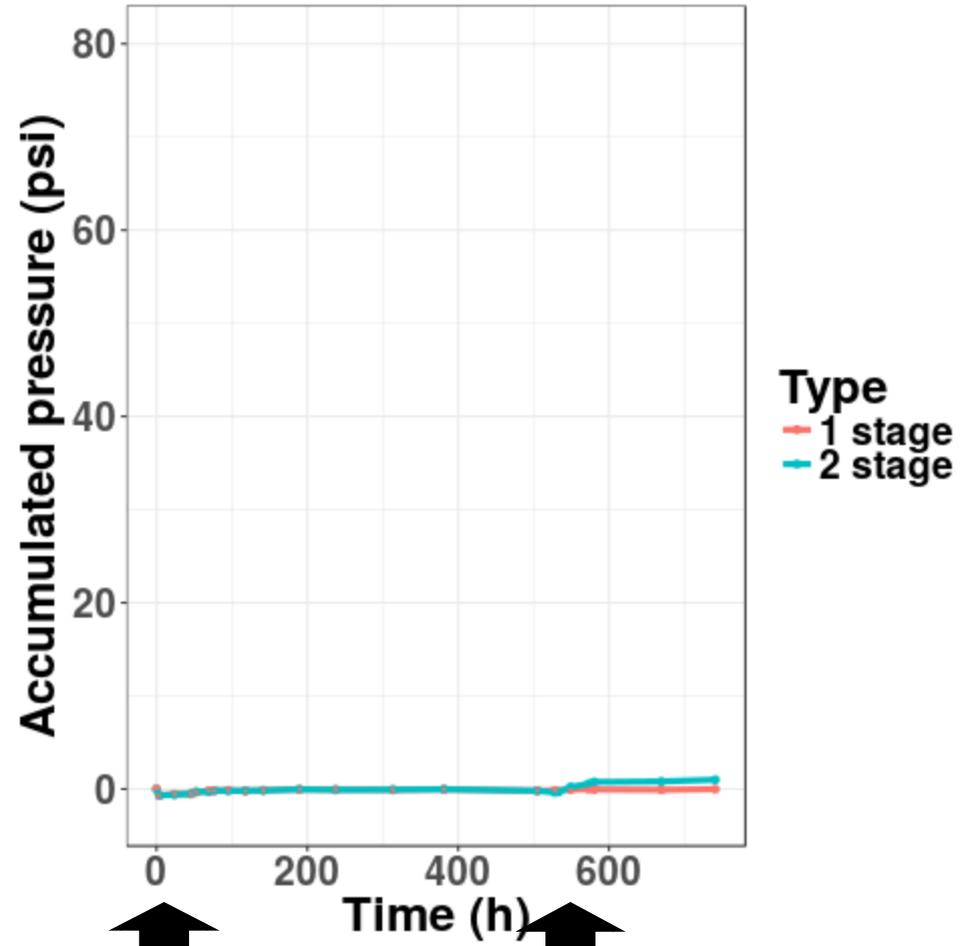
N. californiae + Cac

Cac alone



1 stage: Cac + fungi
2 stage: fungi

2 stage: Cac



1 stage: Cac + media
2 stage: media

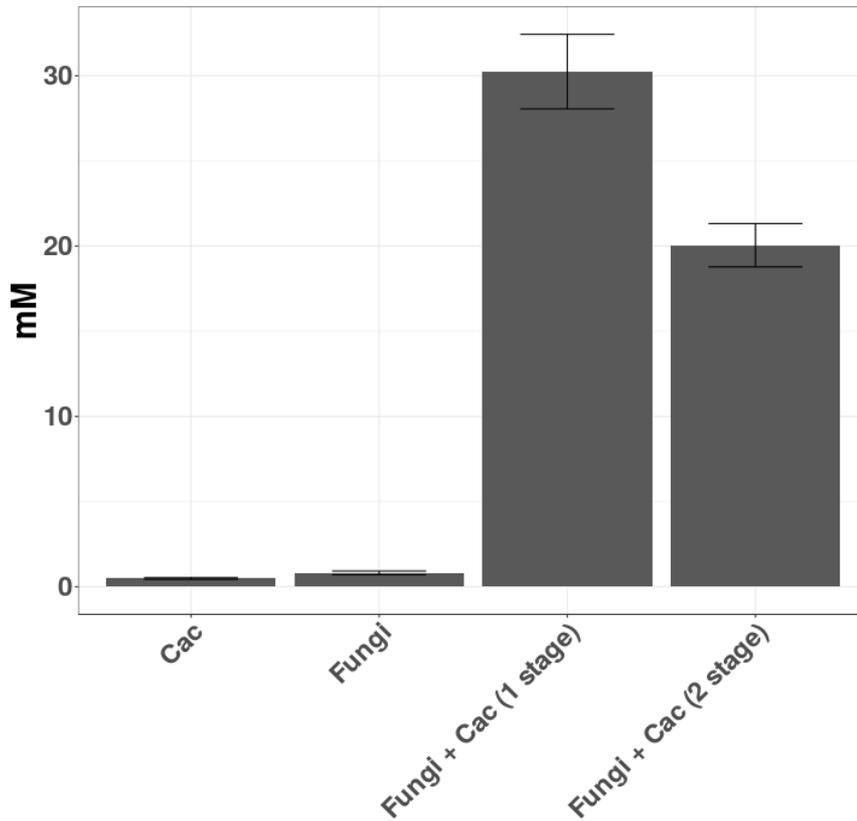
2 stage: Cac



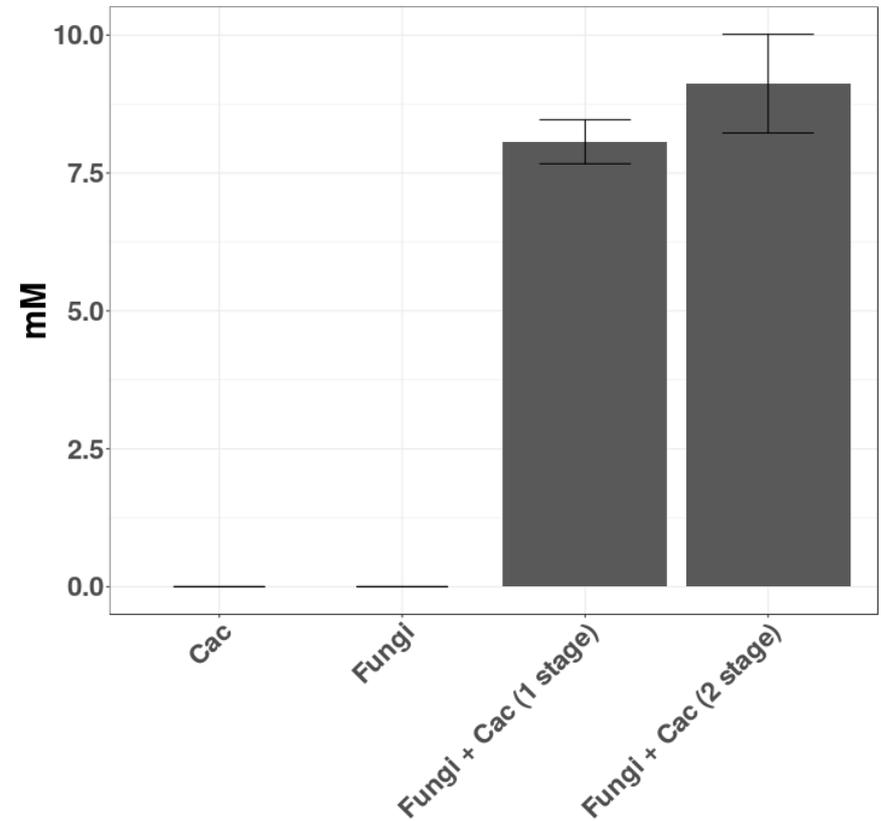
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Cac Metabolic Outputs

Butyrate



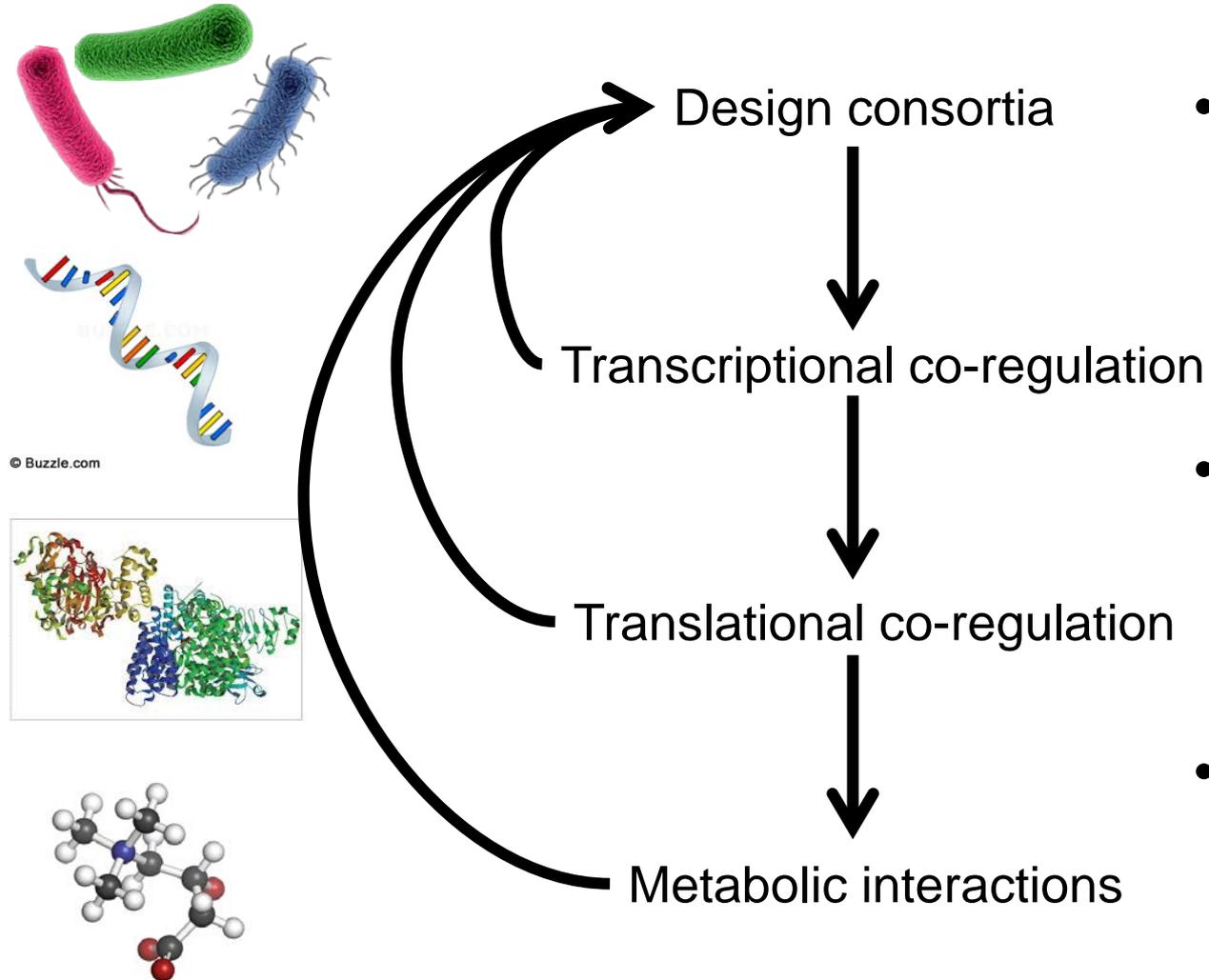
Butanol





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Path Forward



- **Understand synergistic interactions that result in robust, productive consortia**
- **Develop intervention strategies to control consortia assembly and outputs.**
- **Convert food waste and indigenous resources to useful chemicals.**



Discussion & Conclusions

- Co-culturing as alternative to genetic manipulation for robust microbes recalcitrant to genetic changes
- Consortia design allows for:
 - synergistic growth
 - targeting of new output
- Engineer functional and stable interactions between consortia and abiotic components
- Work with NRL, AFRL, MIT to transition functional chemical targets for further analysis and scale up.

Team- Dr. Matthew Perisin, Dr. Theresah Zu, Dr. Sanchao Liu, Dr. Rebecca Renberg, Dr. Nathan Schwalm, Dr. Christian Sund (DARPA detail), Elliot Gerlach, Marcus Benyamin