



DOE Bioenergy Technologies Office (BETO) 2021 Project Peer Review

SOFAST: Streamlined Optimization of Filamentous *ArthrospiralSpirulina* Traits

March 22, 2021 Advanced Algal Systems

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Lumen Bioscience



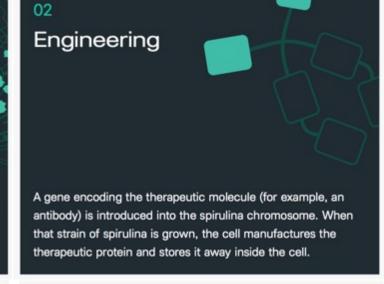
Lumen Bioscience, Inc.

Lumen's interests center around using spirulina to tackle problems in the development of oral therapeutics:

- Commercially viable
- Reduction in costs and risks
- Accelerated time-to-market
- Shortened development process

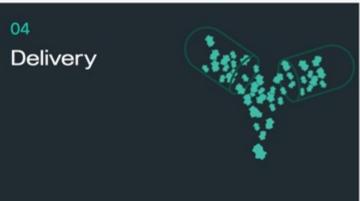
The Organism

Being extraordinarily high in soluble protein, spirulina cells are able to express far higher amounts of therapeutic proteins than any other food crop (>60%).





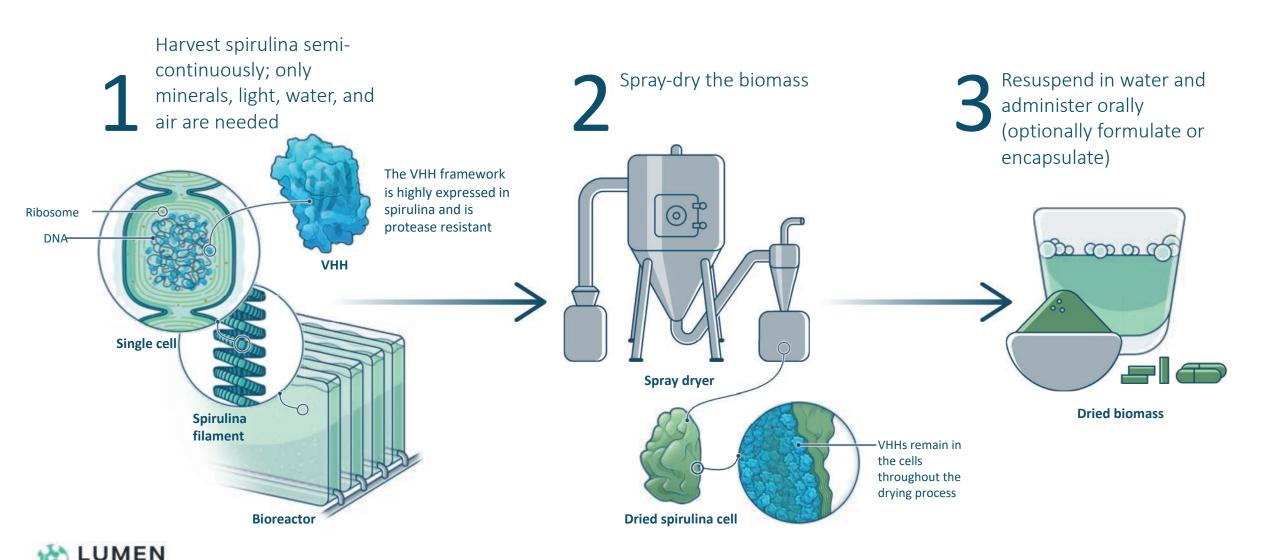
The production system requires only water, salt, CO₂ and light, so it is cheap and rapidly scalable. Harvesting is done by spraydrying the biomass into a powder comprised of spirulina cells, each one filled with a therapeutic protein "payload." The entire process is done at our integrated Seattle lab and cGMP plant, which accelerates pre-clinical and clinical development.



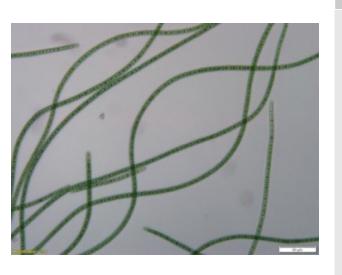
This powder can be packed into dose-specific capsules, which don't require refrigeration and are shelf-stable at room temperature. While the cells do not survive the drying process, the cell membrane protects the therapeutic proteins during transit through the stomach when orally delivered, and releases them in the small intestine where they can bind to and neutralize their disease targets.



Ultra simple manufacturing enables patient safety at de minimis cost



Spirulina (*Arthrospira platensis*): a new biotechnology platform



Lumen developed genetic engineering for Spirulina

Advantages Robust grower, resistant to predation Easier to harvest compared to unicellular

algaeCan be grown in saline or wastewaterbased media

Challenges

- Its protein rich biomass is naturally low in fatty acids and it therefore has a lower energy density than oleaginous algae.
- It is a new chassis. More molecular tools must be developed and demonstrated.
- Its photosynthetic performance is still limited, especially at high culture densities and under stress.



Project Overview – Goal Statement

- Arthrospira species ("Spirulina")
 - Cyanobacteria with proven industrial cultivation capabilities.
 - Lumen has unique, patented technology to genetically enhance *Arthrospira* for improved growth and biomass composition for biofuels and bioproducts.
- <u>Goal:</u> To produce engineered Arthrospira platensis strains with improved photosynthetic traits and increased lipid content.
- <u>Outcome:</u> Demonstrate successful and stable outdoor productivity of 4 engineered *Arthrospira* strains with at least 1 achieving growth rates of at least 19 g/m²/day (AFDW) and at least 14% lipids and 16% carbohydrates.
 - Develop strain improvement toolkits and methods (SOFAST)
 - Increased areal productivity <u>AND</u> projected biofuel yield using these toolkits and methods

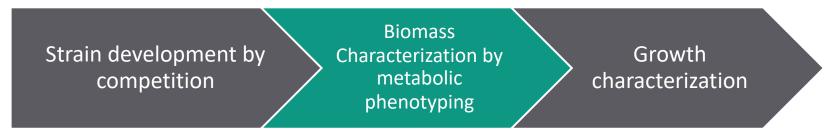


Project Overview

 Goal: To produce engineered Arthrospira platensis strains with improved photosynthetic traits and increased lipid content.

National Renewable Energy Laboratory	Lumen Bioscience
 High throughput methods for biomass characterization 	 Increases lipid production in cyanobacteria (wax esters) Can genetically engineer Arthrospira and select for optimal expression Demonstrated traits that alleviate light and cold stresses in cyanobacteria

This project aims to optimize expression of traits in <u>Arthrospira</u> that increase outdoor biomass productivity and enhance biomass composition for production of biofuel intermediates





1 - Management: Collaboration overview

Project split into three tasks to fulfill project objectives

TASK 2: STRAIN DEVELOPMENT Lumen Bioscience	TASK 3: METABOLIC PHENOTYPING National Renewable Energy Laboratory (NREL)	TASK 4: OUTDOOR GROWTH Arizona Center for Algae Technology and Innovation (ASU)
 Molecular biology and traits Strain and library construction Streamlined selection of traits Strain engineering, sequencing, and characterization Artificial light (indoor) culturing and validation 	 Analytical Biomass characterization by traditional and higher throughput methods Lipidomics characterization in WT and engineered strains Development of single-filament phenotyping and NIR high-throughput methods for Arthrospira 	 Outdoor cultivation Outdoor (sunlight) culturing Media optimization for outdoor culturing
Jim Roberts – Lumen PI Team: Damian Carrieri, Rachelle Lim, Troy Paddock, Lauren Goetsch	Lieve Laurens – NREL co-PI, Algae Platform Team Lead Team: Peter Shanta, Hannah Alt, Kaitlin Lesco	John McGowen - Director of Operations Team: Jessica Forrester

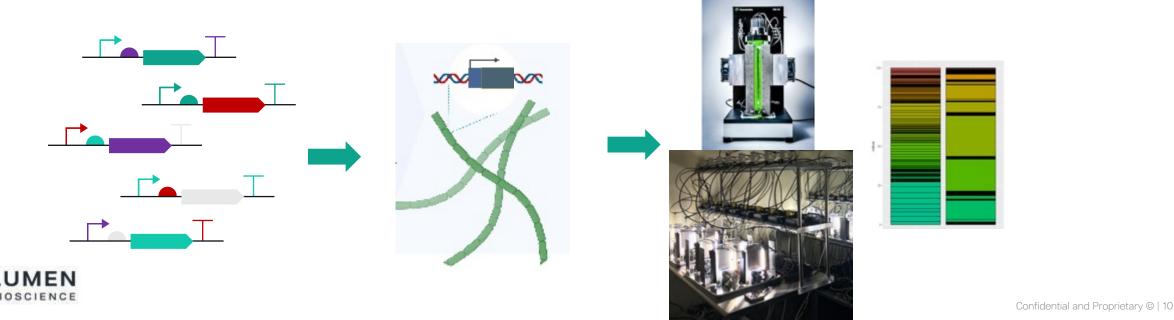
Communication plan:

- Biweekly meetings between NREL and Lumen
- Biweekly calls with ASU during project work
- SharePoint site
- Benchling LIMS
 platform for sample
 tracking and data
 management
- Monthly meetings with BETO project team

2 – Approach Overview

Construct libraries Transform Compete Build Characterize Grow

- 1. Construct libraries of traits with various coding sequences and promoter/terminator strengths (100+ variants in each library).
- 2. Transform libraries of traits into A. platensis using Lumen's patented genetic engineering technology
- 3. Compete transformed libraries against control cells (expressing YFP) or each other and sequence to identify constructs that imparted the best fitness or most wax esters without severe growth defects.



2 – Approach Overview

Construct libraries

Transform

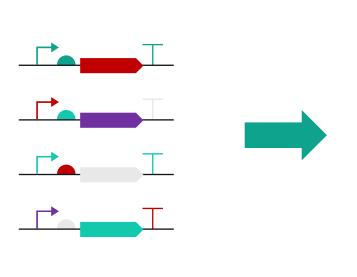
Compete

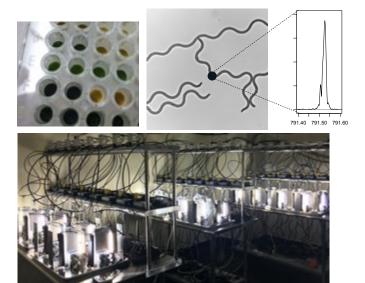
Build

Characterize

Grow

- **4. Build** top constructs into *Arthrospira* to produce strains expressing traits at optimal levels.
- **5. Characterize** newly made strains for improved growth rates or metabolic profiles relative to *wild type* control runs.
- 6. Grow improved strains in indoor and outdoor test beds alongside wild type controls.











2 – Approach: Challenges

Critical Success Factor	Challenge	Strategy	Go/No Go evaluation
Positive effect of stress resistance over-expression resulting in improved growth performance	Stress resistance and cold tolerance traits can adversely affect growth if expressed too strongly or be ineffective if expressed too weakly.	Use traits already demonstrated to improve growth in cyanobacteria. Compete diverse libraries with a wide range of expression levels. Frequently monitor competitions over time and with replicates.	 Successful library construction Rebuild 4 strains Increase in productivity >20% over baseline
Production of wax esters without severe growth defects.	Expression levels of enzymes in the wax ester synthesis pathway must be well-proportioned to avoid accumulation of damaging intermediates.	Same as above and: Use two-gene cassettes, minimizing the number of enzymatic steps.	 Successful library construction Rebuild 4 strains 12% extractable lipids and detectable levels of wax esters
Improvements to strains in the laboratory must translate to improve outdoor performance.	They may not.	Compete and test strains under conditions that closely resemble outdoor cultivations.	PEAK challenge: strain with >50% growth improvement over WT, 14% extractable lipids

3 – Impact

Goals of project: To enhance outdoor growth and lipid content in *Arthrospira platensis* strains through streamlined optimization of genetic traits resulting in at least one achieving growth rates of at least 19 g/m²/day and at least 14% lipids and 16% carbohydrates.

Project Feature	Relevance
Working with Arthrospira platensis	A robust strain, novel to the BETO AAS program, resistant to predation, with commercially demonstrated outdoor production. More easily harvested than unicellular algae.
Optimization of stress-resistance traits	Increased productivity and robustness to factors such as temperature, seasonality, and competition.
	Addressed BETO Barrier Aft-C. Biomass Genetics and Development.
N	Lumen impact: exploration of strain for use as parental background



3 – Impact

Project Feature	Relevance
Optimization of wax ester expression traits	Increased fraction of compounds important for biofuel / bioenergy production thereby enabling algae biomass compositions in environmental simulation cultivation conditions with increased energy content and convertibility.
Development of high- throughput metabolic profiling methods	Increased understanding and characterization of variability in novel feedstock characteristics. Addresses BETO Barrier Aft-E. Algal Biomass Characterization, Quality, and Monitoring Lumen impact: improved TEA model inputs, improved understanding of growth parameter impacts on metabolic profiles
Cultivation targets	Demonstrated translation of laboratory improvements to industrially relevant (outdoor/solar) cultivation approaches, <u>driving down the cost of biomass production</u> . Produced <u>strains that can be licensed</u> with demonstrated enhanced ability to be grown outside for biofuels. Demonstrated <u>traits and engineering tools that can be licensed</u> for enhanced algal biofuel productivity in <i>Arthrospira</i> or other cyanobacteria.

3 – Impacts: Lumen future plans

Lumen Bioscience impacts and future plans:

- 1. exploration of strain for use as parental background
- 2. improved TEA model inputs, improved understanding of growth parameter impacts on metabolic profiles
- 3. Toolkit applications to ACCESS CARBON award

Data sharing methods:

- Publication in scientific journals
- Invention disclosures
- Patent applications



4 – Progress and Outcomes

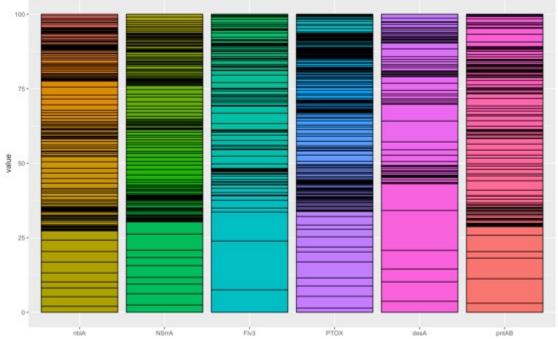
Construct libraries Transform Compete Build Characterize Grow

Deliverable 2.2.1 Successfully create transformation of 4 libraries (theoretical diversity >100) of stress resistance traits (Lumen) [M9]

Promoter Coding Sequence Termin ator

umen) [M9] plasmid library distribution used for transformation

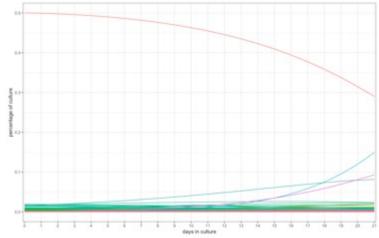
Gene library	Unique Promote rs	Unique Coding Sequence Versions	Terminators present	Theoretical library diversity	Actual library diversity (NGS)
nblA	22	3	3	198	173
flv3	22	2	3	132	121
NSrrA	22	3	3	198	169
PTOX	22	4	3	264	232
desA	22	2	3	132	93
pntAB	22	3	3	198	138



4 – Progress and Outcomes

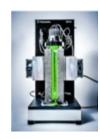
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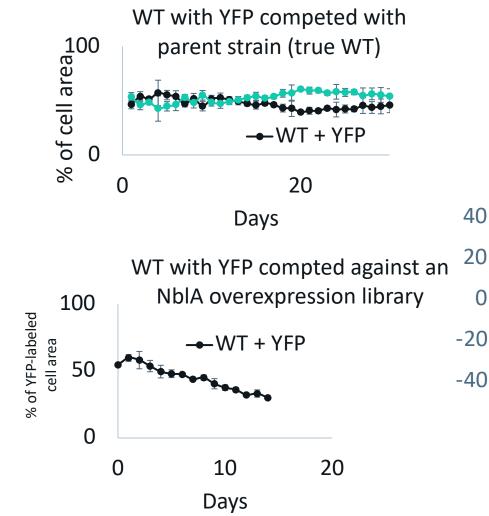
Competition model for growth at Wdt = 13.86 hours: 21 days of growth sufficient to identify winners at 0.1% of the starting population



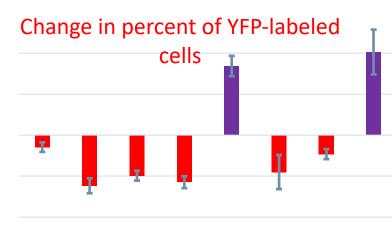
Turbidostat mode: OD = 0.1







5 of 7 competitions led to decrease in share of WT-YFP population relative to stress tolerant libraries



4 - Progress and Outcomes

Construct libraries

Transform

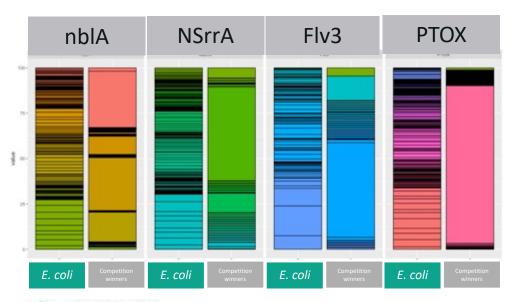
Compete

Build

Characterize

Grow

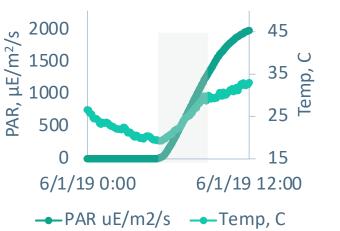
Milestone 2.2.2 Demonstrate productivity improvement over baseline growth for improved strains, by evaluating top strains by indoor, artificial light testing of productivity improvements with continuous monitoring of compositional improvements (Lumen) [M24]



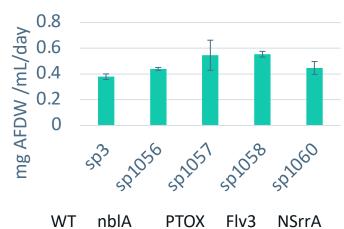
- Rebuilt 20
 strains from
 stress-tolerance
 libraries
- Characterized growth in flask and flat panel reactors



Outdoor growth in Arizona (6/1/2019)



4-day growth rates of rebuilt strains in flask : continuous light, 23C





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4 – Progress and Outcomes

Construct libraries

Transform

Compete

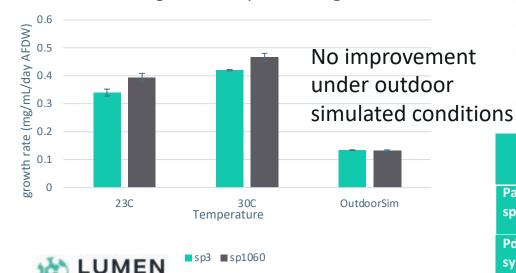
Build

Characterize

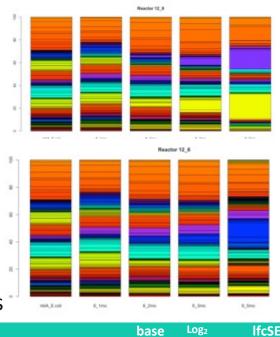
Grow

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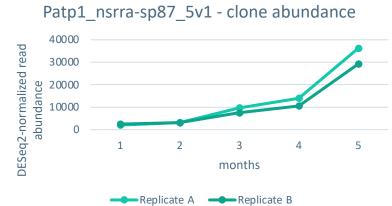
Growth performance at constant and outdoor simulated light and temperature regimes



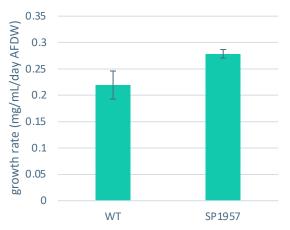
Repeated transformation of SOFAST libraries in flat-panel reactors under outdoor simulated conditions



	base Mean	Log ₂ FoldChan ge	IfcSE	padj
Patp1_nsrra sp87_5v1	1180 2.6	3.86	0.24	9.56E -69
PdnaK_nsrra synPCC coop_4v1	1627 2.2	3.7	0.17	2.08E -120







4 - Progress and Outcomes

Construct libraries

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Deliverable 2.3.1 Successfully create transformation libraries (theoretical diversity >100) of a two-gene wax-ester cassette.

Deliverable 2.3.2 Demonstrate continuous strain selection in ePBR of wax ester libraries for at least 14 days

Milestone 2.3.1 Evaluate top 5 strains by indoor, artificial light testing of productivity. Demonstrate growth of wax ester strains at rates not more than 10% slower than the wild type baseline.

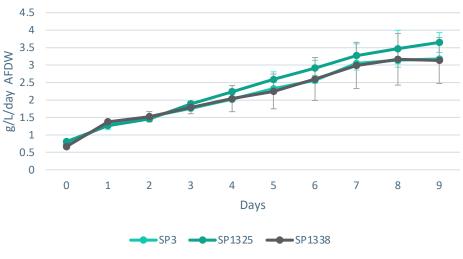


Library	E. coli Sanger- sequencing estimated diversity (Chao1)	MinION- determined Library diversity of A. platensis library at end of competition
FAR12-AtfA45	128	35
FAR14-AtfA45	135	19
FAR16-AtfA45	135	37
FAR18-AtfA45	135	19
FAR12-AtfA67	128	51
FAR14-AtfA67	135	27
FAR16-AtfA67	135	37
FAR18-AtfA67	135	55

LUMEN BIOSCIENCE NREL GC and ESI data informed clones to rebuild potential wax ester-producing strains

No impact to growth for wax-ester producing strain SP1325 under constant light or outdoor simulated conditions

Characterization of wax ester-producing strains in
9L reactors – constant light



Novel Wax Ester strains	ID	Wax ester conten t as % lipid	Growth rate: Constant light/temp in miniarray (mg/mL/day) [%WT]	Growth rate: Outdoor simulation in miniarray (mg/mL/day) [%WT]	Growth rate: Constant light/temp in 9L reactor (mg/mL/day) [%WT]
PpsaF-FAR14coop_Pgap205- AtfA45	SP1325	1.14	0.35 [103.2]	0.159 ± 0.021 [83]	0.324 ± 0.042 [118]
PpetB-FAR16coop_PccmK-AtfA67	SP1338	0.74	0.148 [43.6]	0.145 ± 0.023 [76]	0.271 ± 0.09 [98]

4 - Progress and Outcomes

Construct libraries

Transform

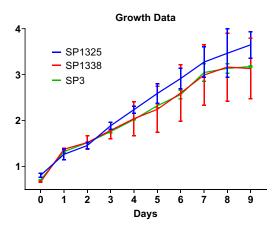
Compete

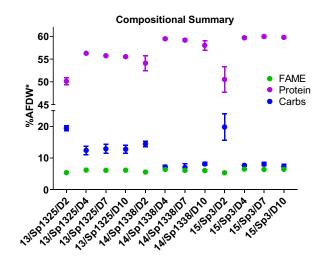
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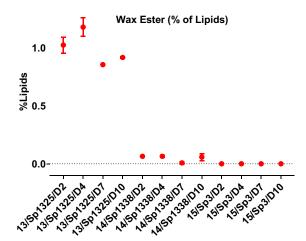
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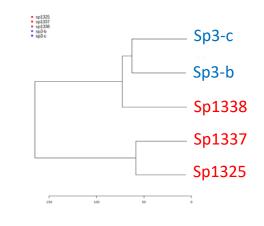


Sample	Fame	Protein	Carbs
17/Sp1325/D4	6.09	56.16	11.46
17/Sp1325/D7	6.01	55.83	13.88
17/Sp1325/D10	6.09	55.46	13.71
14/Sp1338/D4	6.4	59.35	7.6
14/Sp1338/D7	6.13	58.98	7.88
14/Sp1338/D10	6.16	57.27	8.46
18/Sp3/D4	6.54	59.82	7.84
18/Sp3/D7	6.36	60.18	7.68
18/Sp3/D10	6.41	59.63	7.78

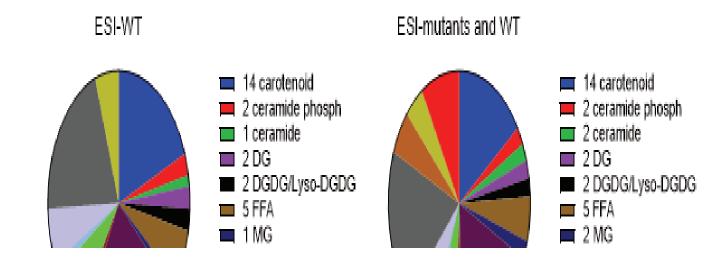


4. Progress and Outcome

- Wax ester producing strains shift biomass composition over time
- SP1325 exhibits significantly increased carbohydrate content at no reduction (minor increase) in growth rate
- Lipidomics profile shows distinct clustering of SP1325 mutant dominated by polar lipid and WE physiological changes
- Detailed metabolomics analysis is ongoing to narrow down carbon sequestration pathways in SP1325



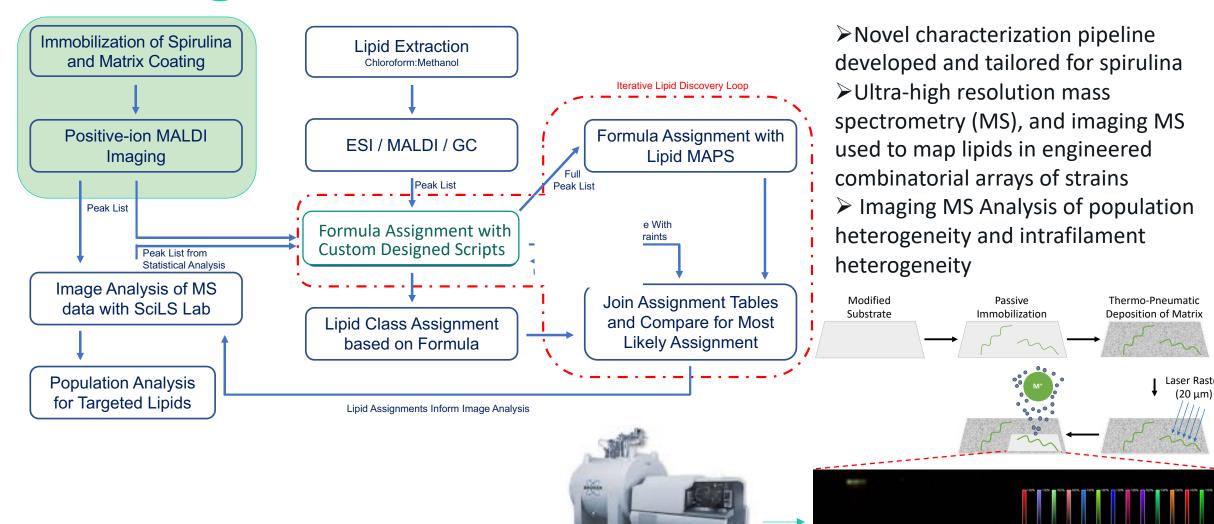
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18/Sp3/D10	6.41	59.63	7.78







4. Progress and Outcome

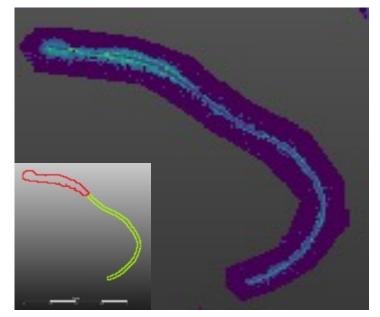




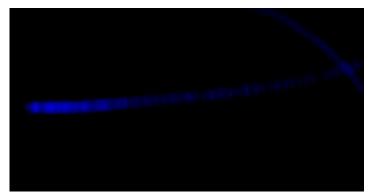


4. Progress and Outcome: Toolkit development

- First demonstration of MALDIimaging MS shows population heterogeneity
- Intrafilament imaging lipidomics shows differential concentration of distinct physiology within spirulina filament
- MS multivariate analysis narrows down lipidome changes associated with filament ends, a total of 26 lipids can be differentially expressed



Imaging MS profile of MGDG (775.5331 m/z) across a filament



Fluorescent D-amino acid dye (7-hydroxycoumarin 3-carboxylic acid (HCC-OH) attached to a D-amino acid backbone, 3-amino-D-alanine, HADA) image, showing peptidoglycan intercellular boundaries on one filament end



4 - Progress and Outcomes

Construct libraries

Transform

Compete

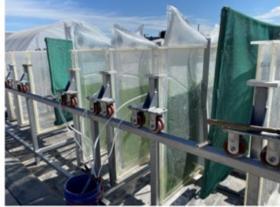
Build

Characterize

Grow



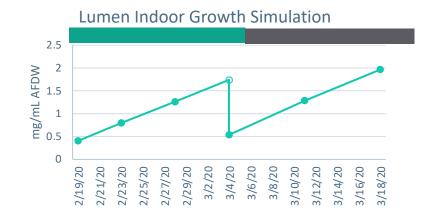




MOBILE MINI - OUTDOOR GROWTH OCT TO NOV 2019



growth g/L/day error 0.102 0.004 .UMEN First 0.0904 0.005 Second



Lumen Growth perforr

n Indoor	growth	g/L/day	error
h	First	0.0953	0.0015
mance	Second	0.102	0.003

Milestone 4.3.1 Test two stress resistant strains in outdoor cultivation (season 1, winter/spring), demonstrating at least 50% productivity improvement in biomass growth relative to WT baseline. (ASU) [M33]

> Best performing stress tolerance and wax ester strains for outdoor testing at AZCATI

SP1060 and SP1325

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Summary

The project aimed to enhance growth and lipid content in an industrially relevant cyanobacterium, Arthrospira platensis.

<u>Approach</u>: combine genetic engineering and selection approaches with high-throughput biomass characterization to select for winning genetic constructs that we transform into *Arthrospira* stains.

Construct libraries Transform Compete Build Characterize Grow

Progress:

- ✓ In situ competitions
- ✓ Metabolic profiling
- ✓ Indoor growth characterization for wax esters
- Outdoor growth seasons

Target: 20% enhancement in growth / >12% lipids
19 g/m²/day, > 14% lipid, >16% Carbohydrate.

Relevance:

- ➤ A robust strain, resistant to predation, with commercially demonstrated outdoor cultivation, that shows improved for outdoor growth AND biomass composition, <u>driving down the cost of</u> biomass production.
- Potential for <u>licensing strains</u>, <u>traits</u>, <u>and genetic</u> <u>tools</u> for algal biofuel production.



Quad Chart Overview (Competitive Project)

Timeline

- October 2017
- June 2021

	FY20 Costed	Total Award
DOE Funding	(10/01/2019 – 9/30/2020) \$377,730	(negotiated total federal share) \$1,193,682
Project Cost Share	\$182,040	\$556,331

Project Goal

To produce engineered *Arthrospira* platensis strains with improved photosynthetic and cold-tolerance traits and increased lipid content.

End of Project Milestone

End of Project Goal: Demonstrate at least 1 improved platform spirulina strain achieves at least a 15 percent improvement in biomass productivity relative to wild type baseline, displaces 25 percent of delivered CO2 with DAC, and has a biomass composition containing at least 2% therapeutic protein and achieves a 100% relative increase in components of biomass that can be used for bioenergy production relative to baseline strain and conditions

Project Partners*

• NREL (34%)

Funding Mechanism

DOE-FOA-0001628: Productivity Enhanced Algae and Tool-Kits (PEAK) 1 Topic Area 1: Strain Improvement

Additional Slides

Responses to Previous Reviewers' Comments

Demonstrated progress on most milestones, unfortunately
the project pivoted away from outdoor raceway trials to
indoor simulated trials instead.

The SOFAST project addresses both the development of tools for strain improvement and characterization as well as the use of those tools to improve an industrially relevant strain. Focusing on Spirulina as a target is logical and relevant given the long history of outdoor cultivation at scale. Key steps have been made towards the project goals around tool development. The use of competition experiments to identify top performing strains under selective pressure is clever and likely to result in a highly efficient methodology. It is unfortunate that the validation of the strain performance will not be conducted in outdoor trials.

Response to Comments

The project pivoted from outdoor raceway trials to outdoor flat-panel trials. Indoor simulations were always planned and remain in the project goals.

Outdoor validation trials will indeed be completed.



Reviewer Comments

Publications, Patents, Presentations, Awards, and Commercialization



LIST ANY PUBLICATIONS, PATENTS, AWARDS, AND PRESENTATIONS THAT HAVE RESULTED FROM WORK ON THIS PROJECT

Patents:

None

Publications / Presentations:

Presentation: Damian Carrieri, Rachelle Lim, Troy Paddock, Jim Roberts, John McGowen, Peter Shanta, and Lieve Laurens. "SOFAST: Streamlined Optimization of Filamentous *Arthrospiral Spirulina* Traits". The Rules of Life: Complexity in Algal Systems; Summer Science Symposium. July 2020.

Manuscript in preparation: "Engineering of Bioenergy Pathways in Cyanobacteria has a Global Metabolic Impact on Photophysiology and Lipid Metabolism" Shanta P., et al.

Manuscript submitted: "Overexpression of NbIA decreases phycobilisome content and enhances photosynthetic growth of the cyanobacterium Synechococcus elongatus PCC 7942" Algal Research. (submission 2020_833). Damian Carrieri, Tracey Jurista, Nina Yazvenko, Adan Schafer Medina, Devin Strickland, and James M. Roberts.

