DOE Bioengery Technologies Office (BETO) 2019 Project Peer Review

Success through Synergy: Increasing Cultivation Yield and Stability with Rationally Designed Consortia

> March 6, 2019 Advanced Algal Systems

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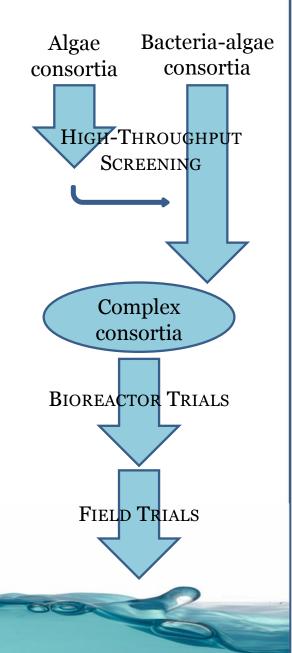
Goal Statement

<u>Project Goal</u>: to increase productivity of open, outdoor *Nannochloropsis* cultures from 7 g/m²/d to >14 g/m²/d (doubling of fall State of Technology value) via **development of (1) rationally-designed intrageneric** *Nannochloropsis* **consortia &** (2) *Nannochloropsis*-bacteria consortia

Outputs: molecular toolkits, pipelines, consortia

Outcomes: increased productivity & stability

<u>Industry Relevance</u>: increased yield, fewer crop losses, & enhanced economic feasibility



Quad Chart

Timeline

- start date -- February 1, 2018
- end date -- January 31, 2021
- ➢ 30% complete (with verification)

	FY 17 Costs	FY 18 Costs	Total Planned Funding (FY 19-21)
DOE Funded	\$0	LANL: \$351 K NMC: \$173 K	LANL: \$1.19 M NMC: \$1.29 M
Project Cost Share	\$0	LANL: \$0 NMC: \$41 K	LANL: \$0 NMC: \$276 K

Barriers Addressed

Biomass Genetics & Development (Aft-C); to improve the productivity and robustness of algae strains against perturbations

Objective

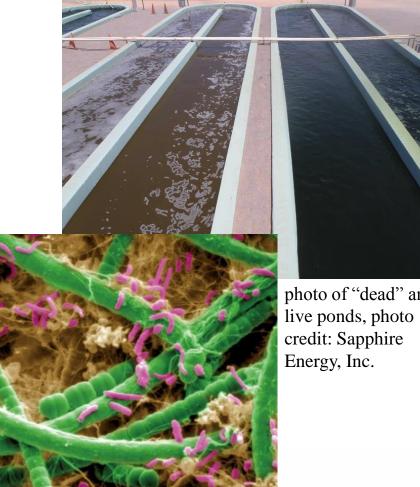
 to rationally design intrageneric Nannochloropsis consortia & Nannochloropsis-bacteria consortia to increase productivity, stability and yield of open, outdoor cultures

End of Project Goal

to reach a productivity target
 >14 g/m²/d with consistent
 biomass composition

1. Project Overview -- *Context and Project History*

- industrial economics of production systems limited by low crop productivity and stability
- consortia of microbial assemblages may address this limitation
 - \succ combining algal strains with complementary traits increases yield and stability (Shurin 2014, Corcoran & Boeing 2012, Stockenreiter et al. 2011)
 - addition of bacteria enhances productivity and reduces crashes (Sapp et al. 2007, Croft et al. 2005, LLNL work)



learning from nature: microbial consortia photo credit: A. Dohnalkova/ PNNL

photo of "dead" and

1. Project Overview -- Context and Project History

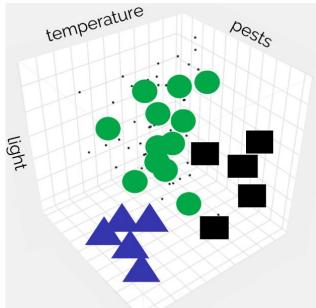
- despite strong theoretical and empirical underpinnings, consortia are not commonly cultivated at scale
- we aim to develop intrageneric algal consortia as well as algal-bacteria consortia (important for consistent biomass composition, downstream processing)
- novel approaches
 - rational design
 - high-throughput screening

Types of consortia

Intrageneric algal – group of algae containing strains from the same genus

Intergeneric algal – group of algae containing strains from different genera

Algal-bacteria – group of algae & bacteria



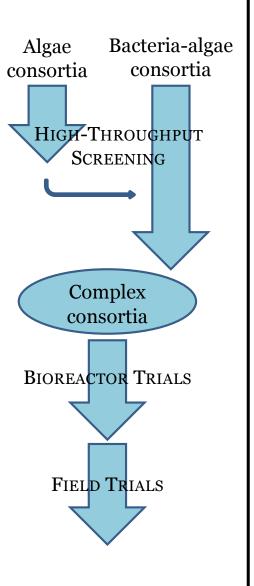
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conceptual model of taxa represented by different colored shapes filling up niche space

2. Approach -- Management

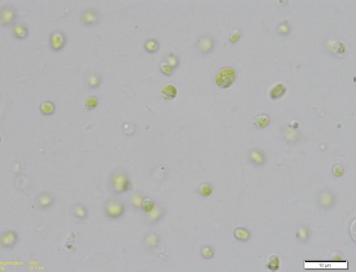
- WBS, SOPO, and Gantt chart with Milestones and Go/No-Go Points serve as <u>guiding documents</u> for the team
- PIs enable communication and collaboration
 - > weekly PI meetings
 - weekly team meetings to plan work, review
 progress towards
 milestones, and
 discuss technical
 barriers
- team members drive progress on specific tasks (see chart)

Task	Key Team Members
1. Verification	All
2. Design of <i>Nannochloropsis</i> Consortia & Tracking Tools	Garcia (Postdoc), Hanschen (Postdoc)
3. Screening & Identification of Growth-Promoting Bacteria	Hovde (Research Scientist), Ohan (Post-MS Student)
4. Verification & Optimization of Consortia Performance	Garcia (Postdoc), Hanschen (Postdoc)
5. PEAK Challenge	Holguin (Assistant Professor), Boeing (Professor) Rodriguz- Uribe (Postdoc)

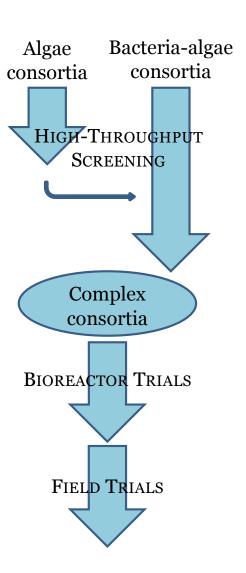


<u>Task #2</u>: Design of Candidate *Nannochloropsis* Consortia & Tracking Tools

- rational design
 - match strains to outdoor regimes by combining taxa with different phenotypes
 - use Sapphire Energy, Inc.* data and strains *original award to Starkenburg at LANL and Corcoran at Sapphire
 - molecular tracking of strains
 - needed to distinguish consortia members
 - cox1 amplicon sequencing

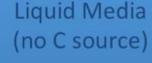


light micrograph of Nannochloropsis strain



<u>Task #3</u>: Screening & Identification of Growth-Promoting Bacteria

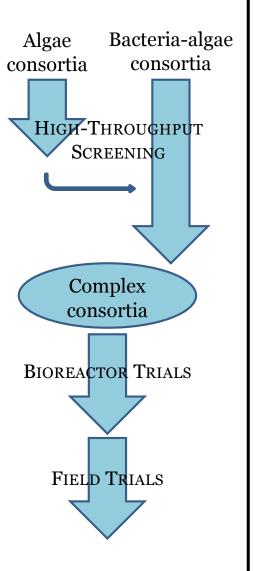
- High-throughput Screening of
 Cell-to-cell Interactions (HiSCI)
 system
- sorting, scale-up, and ID of beneficial bacteria
- sourcing of environmental samples



Solid Agarose

1 algal cell

3-5 bacterial cells

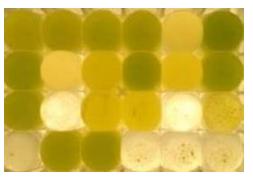


Task #4: Verification & Optimization of Consortia

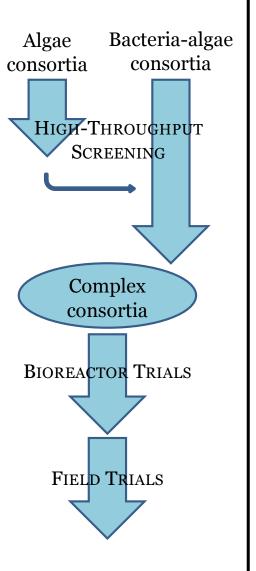
Performance

- individual/complex consortia tested at increasing scales
 - high-throughput screens to confirm co-existence (no stressors)
 - high-throughput screens
 with perturbations
 (temperature fluctuations, invasions)
 - bioreactor trials to better simulate field conditions
 - β-field trials in micro- (100
 L) or mini- (300 L) ponds









Task #5: PEAK Field Challenge

- best performingconsortiavalidated throughoutdoor field trials
- 300-L miniponds
 located at the
 Fabian Garcia
 Center in Las
 Cruces, NM
- metrics of
 productivity (e.g.,
 AFDW) and
 stability (e.g.,
 temporal CV)



Aerial view (top left) of miniponds (top right, bottom) at the Fabian Garcia Center at NMSU

Potential Challenges

- insufficient phenotypic variability across algal strains
- data gaps
- challenges in scaling
- variability in media due to sourcing
- environmental variability difficult to simulate in lab
- control failures

Critical Success Factors

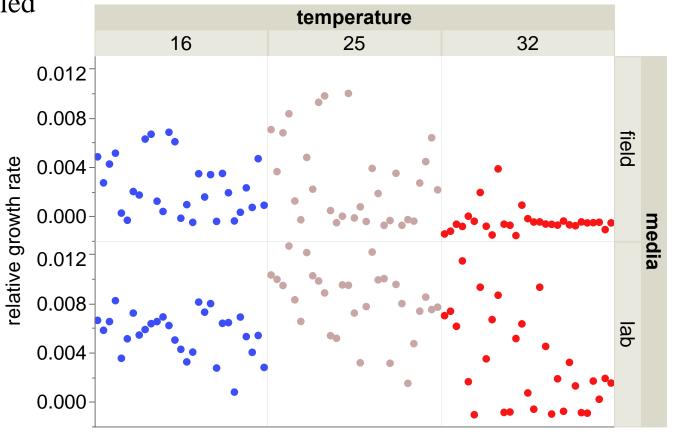
- access to data/analysis methods
 - creative design approaches
- > access to bioreactors, ponds, infrastructure
 - trained pond operators/technicians
 - coordination across team members/sites

The success of this project hinges on consortia performance exceeding the baseline -- *this is the critical success factor for commercial viability.*



1) strain data compiled

Growth rate bioassay data from Sapphire Energy, Inc. for ~30 Nannochloropsis strains (along x-axis) in different media types and temperatures were compiled. Strains grew better in lab media at low or intermediate temperatures.

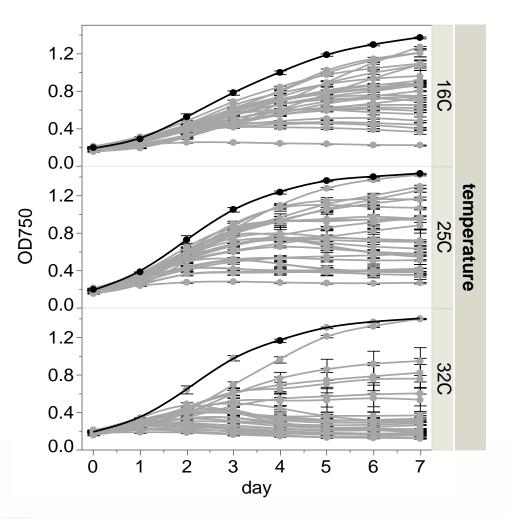


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strain

1) strain data compiled

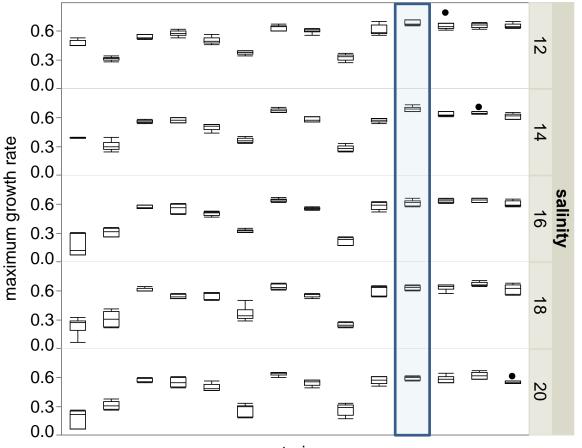
Within the bioassay data, a single top performer (black line) was revealed. This strain was a production strain at Sapphire Energy and is our baseline strain for consortia comparisons.



- 1) strain data compiled
- 2) screenings to enhance database (added assays to overcome data gaps)

salinity

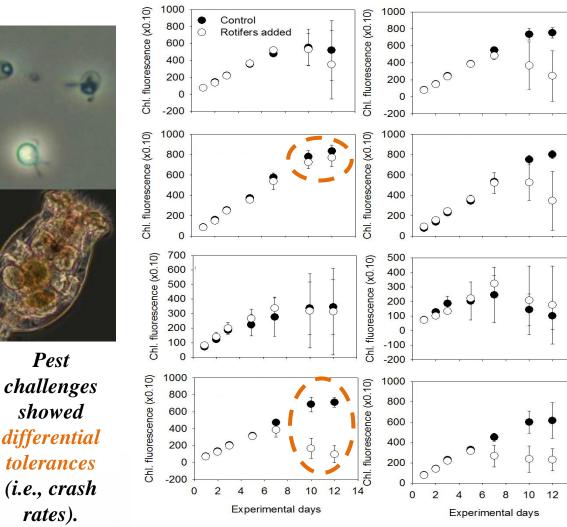
Salinity screenings showed some differential tolerances by some strains. Baseline strain (indicated by box) grew well under all salinities.



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strain

- 1) strain data compiled
- 2) screenings to
 enhance database
 (added assays to
 overcome data gaps)
 - ➤ salinity
 - > pests



- 1) strain data compiled
- 2) screenings to
 enhance database
 (added assays to
 overcome data gaps)
 - ➤ salinity
 - > pests
- 3) 9 consortia designed

#	Rationale
1	top performing strains from each salinity
2	top performing strains from each salinity plus baseline strain
3	top performing strains from temperature screenings
4 & 5	top performing strains from temperature and salinity screenings (two species compositions)
6	consortia with new field isolates
7	kitchen sink (non-rational design for comparison)
8	strains with different intrinsic growth rates
9	strains with known genetic diversity

- 1) strain data compiled
- 2) screenings to
 enhance database
 (added assays to
 overcome data gaps)
 - ➤ salinity
 - > pests
- 3) 9 consortia designed
- 4) molecular tracking tool developed and validated (changed target gene to overcome technical barrier)

Gene/ loci	Amplicon size	Species SNPs	Strain SNPs
cox1	322	5-40	0
rbcL	990	4-76	0
ITS	324	6-50	0-6
ccsA	415	6-50	N. salina N. oceanica

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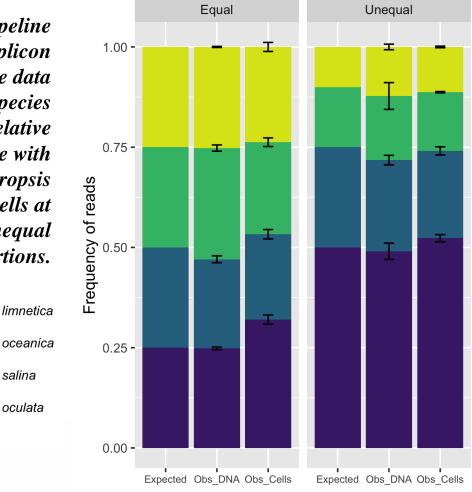
40 Nannochloropsis strains including the following species: australis, gaditana, salina, oculata, granulata, oceanica, limnetica

Evaluation of conserved genes to differentiate species and strains revealed chloroplast ccsA gene a good candidate, compared to cox1 with multiple copies in Nannochloropsis.

salina

- strain data compiled 1)
- screenings to 2) enhance database (added assays to overcome data gaps)
 - \succ salinity
 - \succ pests
- 9 consortia designed 3)
- 4) molecular tracking tool developed and validated (changed target gene to overcome technical *barrier*)

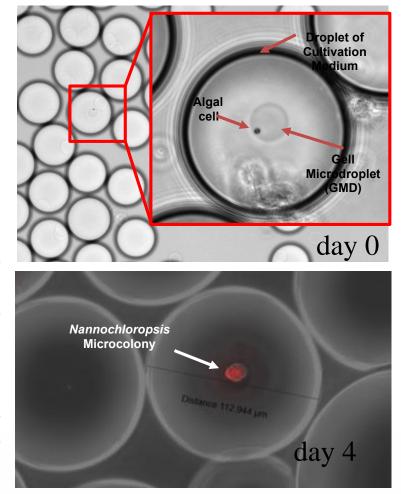
Custom pipeline with amplicon sequence data determined species and relative abundance with Nannochloropsis DNA and cells at equal and unequal proportions.



3. Accomplishments/Progress/Results ^{19 of 30} *Task 3 (Screening & ID of Growth-Promoting Bacteria)*

1) HiSCI system optimized for *Nannochloropsis*

Novel platform HiSCI (<u>H</u>ighthroughput <u>s</u>creening of <u>c</u>ellto-cell <u>i</u>nteractions), previously used for larger cells, optimized for Nannochloropsis cells and used with field medium.



3. Accomplishments/Progress/Results 20 of 30 Task 3 (Screening & ID of Growth-Promoting Bacteria)

- 1) HiSCI system optimized for *Nannochloropsis*
- 2) environmental samples collected

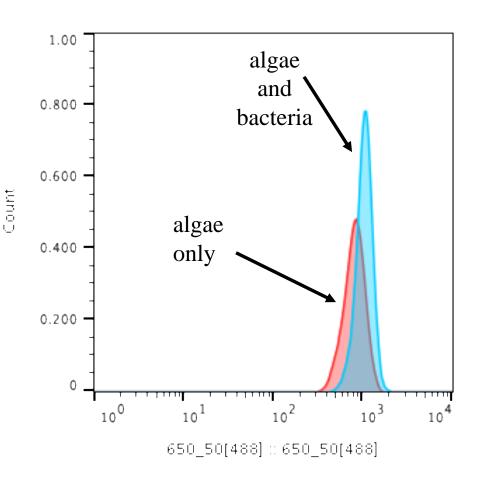


Environmental samples collected from 30+ locations across NM, including Sapphire and Fabian Garcia ponds.



3. Accomplishments/Progress/Results ^{21 of 30} *Task 3 (Screening & ID of Growth-Promoting Bacteria)*

- 1) HiSCI system optimized for *Nannochloropsis*
- 2) environmental samples collected
- 3) screenings conducted
 - 3 iterations with environmentally-sourced bacteria
 - 3 isolates being scaled for further testing



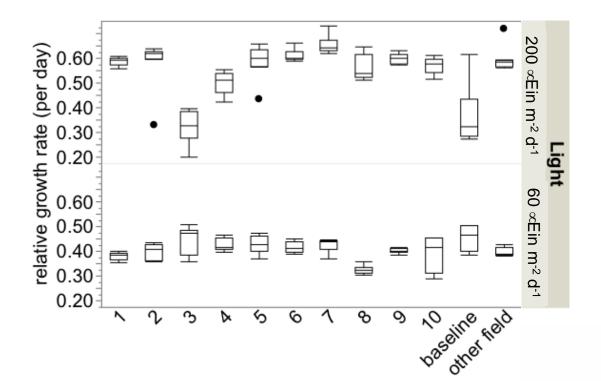
Initial data show increased algal productivity, verified by cell sorting and microscopy during each round of selection when compared to controls.

3. Accomplishments/Progress/Results

Task 4 (Verification & Optimization of Consortia Performance)

- intrageneric Nannochloropsis (algae only) consortia testing in progress
 - 1) high-throughput bioassays

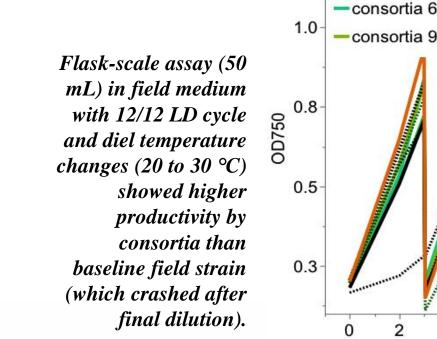
High-throughput assays in field medium with 12/12 LD cycle and diel temperature changes (20 to 30°C) showed higher productivity by some of the consortia (labeled 1-10) than the baseline field strain under higher irradiance.

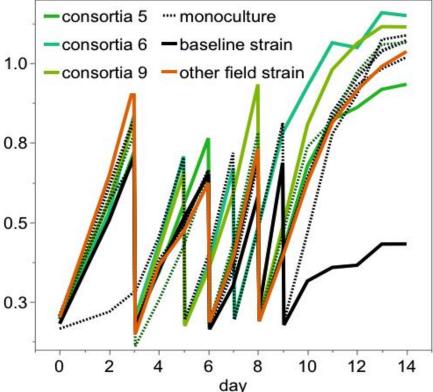


3. Accomplishments/Progress/Results

Task 4 (Verification & Optimization of Consortia Performance)

- intrageneric *Nannochloropsis* (algae only) consortia testing in progress
 - 1) high-throughput bioassays
 - 2) flask-scale bioassays



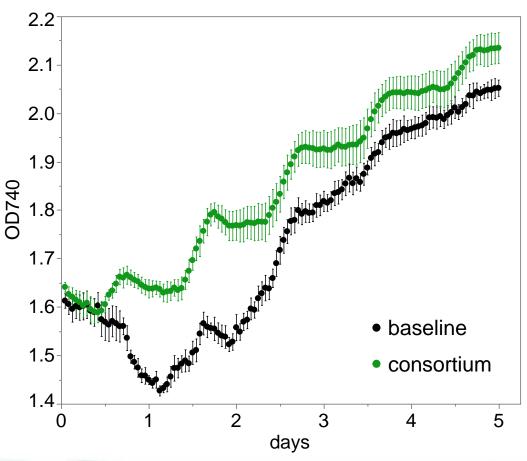


3. Accomplishments/Progress/Results

Task 4 (Verification & Optimization of Consortia Performance)

- intrageneric *Nannochloropsis* (algae only) consortia testing in progress
 - 1) high-throughput bioassays
 - 2) flask-scale bioassays
 - 3) bioreactor trials

Bioreactor trial under field-relevant light and temperature profiles showed lag in growth by baseline strain upon inoculation compared to 4strain consortium (n = 3).



3. Accomplishments/Progress/Results Summary of Milestones

Measure	Description	Due Date	Status
GNG.1	Technical and Cultivation Readiness Validated	30-Apr- 2018	\checkmark
2.1. ML	DNA molecular tracking tool developed to differentiate at least 4 species of <i>Nannochloropsis</i>	31-Jul- 2018	\checkmark
2.2. ML	At least two <i>Nannochloropsis</i> consortia designed based on two key functional characteristics (e.g. growth rate, temperature, salinity, pest tolerance)	31-Oct- 2018	\checkmark
3.1.ML	HiSCI system modified to achieve one <i>Nannochloropsis</i> cell per microdroplet and viable recovery of cells in Las Cruces, NM fall light and temperature conditions	31-Jan- 2019	\checkmark
3.2. ML	Bacteria collected for screening from at least 20 different sources	31-Jan- 2019	\checkmark
3.3. ML	At least one bacterium identified that improves biomass production rates $(g/m^2/day) > 30\%$ in fall cultivation simulations (Las Cruces temperature and light profiles in September, grown in field cultivation media) for one strain/species of <i>Nannochloropsis</i> over the monoculture baseline	30-Apr- 2019	In Progress
4.1. ML	Intrageneric consortia demonstrating a 30% productivity improvement over a <i>Nannochloropsis</i> monoculture grown in triplicate under fall Las Cruces environmental conditions in commercial cultivation media	31-Jan- 19	\checkmark

4. Relevance

Project Goal:

- double productivity of open, outdoor *Nannochloropsis* cultures via through intrageneric consortia & algae-bacteria consortia
- ➤ important in its utilization of ecological and high throughput approaches

Industry Relevance:

- project has potential to advance the state of technology and enhance the economic viability of algal biofuels by increasing yield & reducing losses in cultivation systems
- project supports BETO goals by advancing toolkits and technologies to meet 2020 DOE PEAK targets of 18 g/m²/d and 80 GGE ton⁻¹ biomass

Technology Transfer:

- \succ consortia will be available for use in industry
- ➤ information will be available to broader community via publications

5. Future Work

Task 3 (Screening & ID of Growth-Promoting Bacteria)

- Main Tasks
 - scale-up to larger volumes in flasks to verify phenotype(s)
 - > new HiSCI runs with field isolate and known growth-promoting taxa
- Stretch Goal
 - > assays with taxa described in literature or acquired from collaborators (Sandia)
- Milestones
 - 3.3.ML At least one bacterium identified that improves biomass production rates >30% in fall cultivation simulations for one strain/species of *Nannochloropsis* over the monoculture baseline
 - GNG.2 Bacteria-augmented *Nannochloropsis* cultures demonstrate a 30% productivity improvement over the monoculture baseline

mitigation: low-throughput screening with known beneficial taxa

5. Future Work

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Task 4 (Verification/Optimization of Consortia Performance)

- Main Tasks
 - scale-up to larger volumes to verify phenotype(s)
 - continue perturbations
- Stretch Goal
 - ➤ testing of intergeneric consortia
- Milestones
 - 4.1.ML Intrageneric consortia demonstrating a 30% productivity improvement over a *Nannochloropsis* monoculture grown in triplicate under fall Las Cruces environmental conditions in commercial cultivation media
 - GNG.3 Simulations or fall field trials demonstrating productivity improvements over the monoculture baseline to reach target of 14 g/m²/d

mitigation: UV mutagenesis, intergeneric consortia

5. Future Work *Task 5 (PEAK Field Challenge)*

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• Main Tasks (at NMSU)

- ➤ scale consortia members for field trials
- grow consortia and baseline control in miniponds (25 cm depth, 300-L) semicontinuously for >2 growth cycles and sample for metrics of biomass
- Stretch Goal
 - ➤ repeat with pest challenges or intergeneric consortia
- Milestones
 - 5.1 ML Algae and bacteria inocula scaled to >200L for 300-L PEAK field challenge
 - 5.2.ML Field trials demonstrating >30% productivity and stability improvements over monoculture baseline for a target of at least 14 g/m²/d in the fall growth season (Sept-October)

Summary

- <u>Overview</u>: project aimed to improve productivity and robustness of strains, resulting in a doubling of fall productivity
- <u>Approach</u>: development of intrageneric algal consortia via rational design and algal-bacteria consortia via high throughput screening
- <u>Accomplishments/Progress/Results</u>
 - Task 2: strain data compiled, additional screenings conducted, 9 consortia designed, molecular tracking tool developed and validated
 - Task 3: HiSCI system optimized, >30 environmental samples collected, 3 iterations conducted, additional screenings in progress
 - Task 4: high-throughput and flask-scale bioassays completed, additional trials in progress
- <u>Future Work</u>: continuation of Tasks 3 and 4 (consortia development and testing) and PEAK Field Challenge
- <u>Relevance</u>: **direct relevance to the bioenergy industry** because project addresses challenges associated with open ponds