

Engineering Green Algae for High Biomass Productivity

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Problem Statement

Algae are excellent candidates for providing a renewable source of liquid fuels with an overall reduction in carbon emissions compared to conventional petroleum fuels. However, the economic viability of algal biofuel production at industrial scales is limited by the low biomass productivities of natural algal species. While algae rely on photosynthesis to fix CO₂ into metabolites for biomass generation, several other pathways act in a reverse manner, leading to the loss of biomass and release of CO₂. At night, algae activate the process of dark respiration, which allows the algal cells to continue the process of cell division but also leads to the loss of fixed carbon (Figure 3). In the light, photorespiration also results in the loss of fixed carbon and other cellular resources (Figure 2). Together, these two metabolic processes account for a 35-115% reduction in biomass productivity.

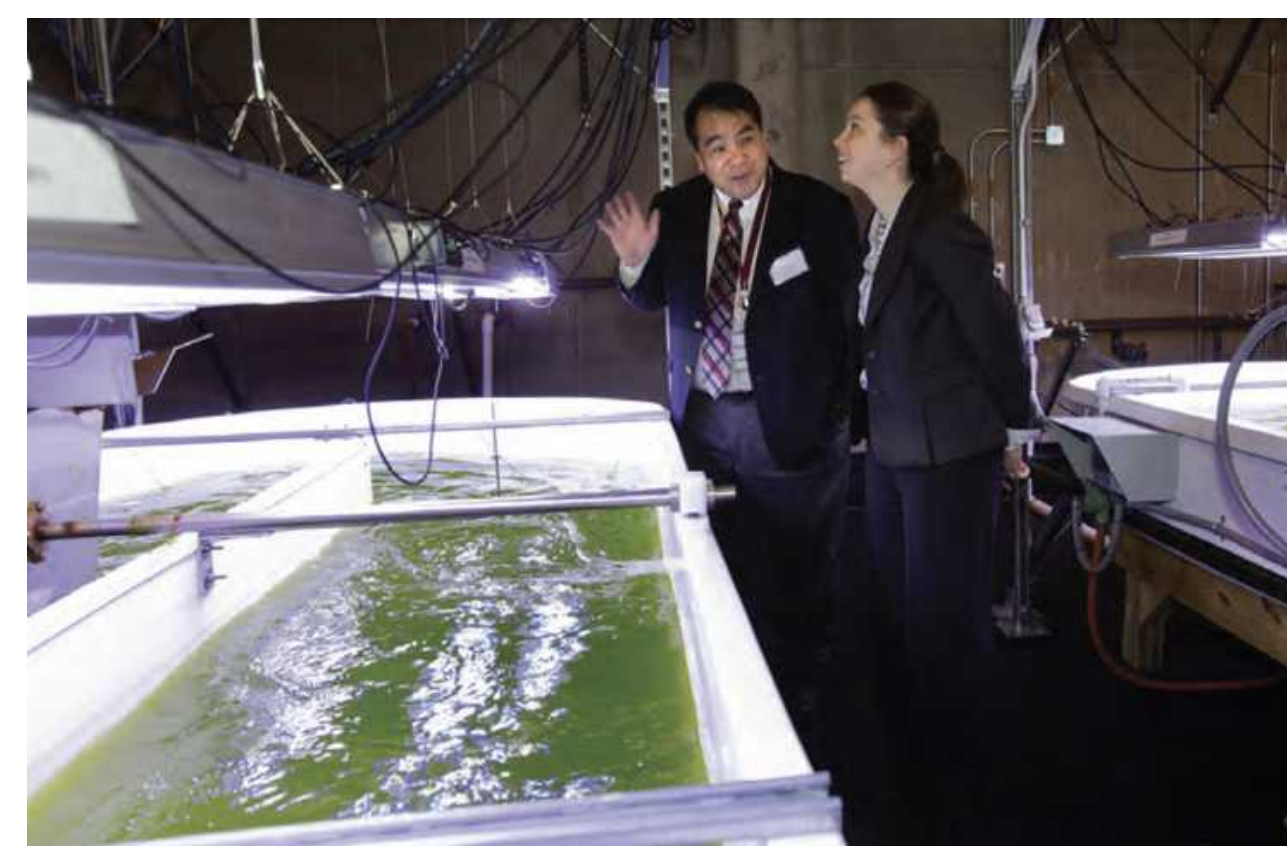


Figure 1. Algal Testbed at SNL, includes three 1,000 L raceway ponds with LED lights and temperature control to simulate outdoor conditions.

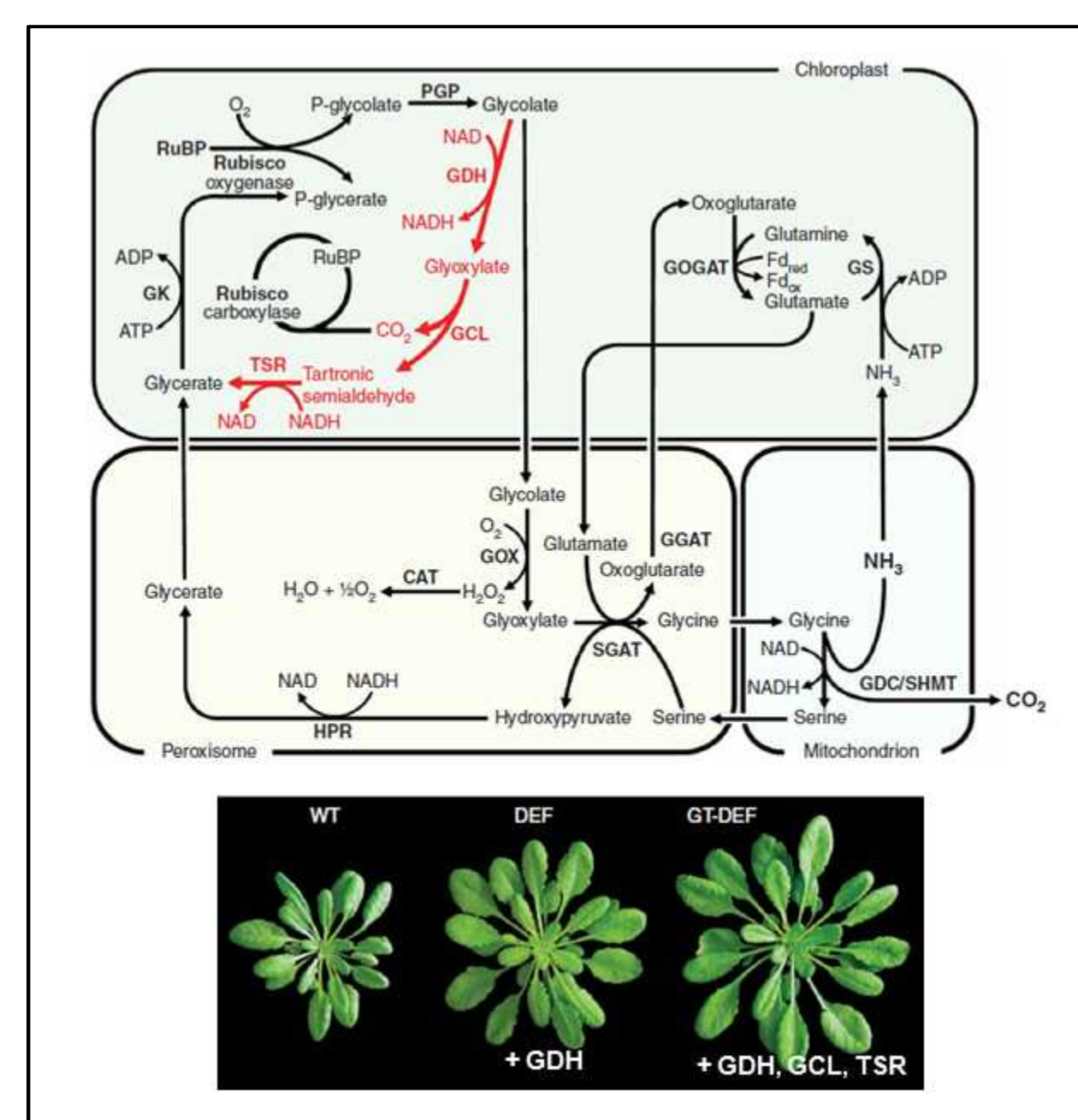


Figure 2. Genetic modification of plants with bacterial photosynthetic bypass pathway to reduce carbon loss associated with photorespiration.¹

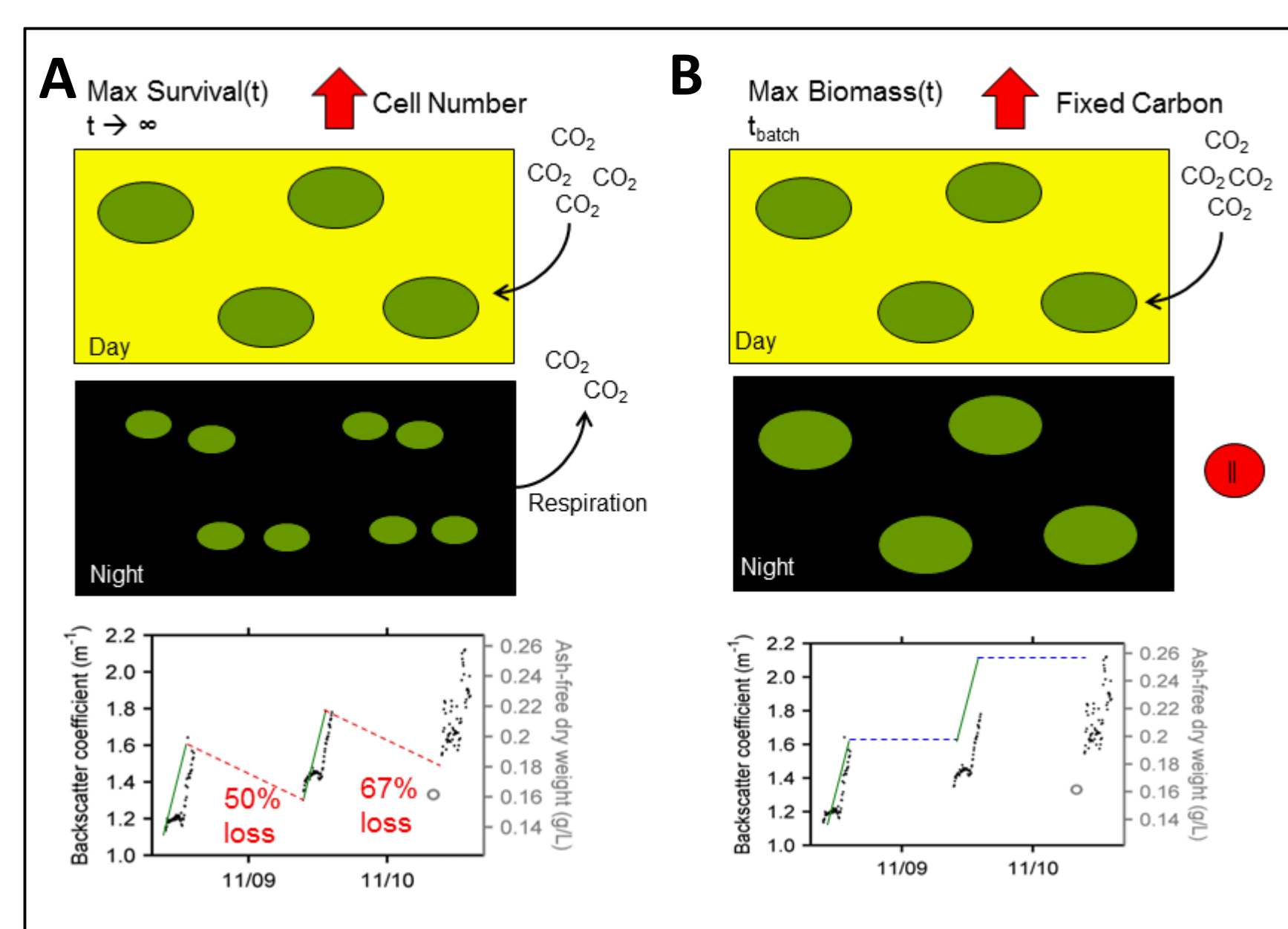


Figure 3. Carbon loss due to dark respiration in *Nannochloropsis salina* (A) and proposed *Nannochloropsis* strains with genetic modification of dark respiration (B).²

Hypothesis and Objectives

We hypothesize that reducing carbon loss from dark respiration and photorespiration through the genetic modification of algae will result in improved algal biomass productivities. We will test this hypothesis through the following objectives:

1. To develop CRISPR/Cas9 genome editing tools for *Nannochloropsis* species (Figure 4)
2. To investigate dark respiration using chemical inhibitors
3. To investigate glycolate production and excretion in *Nannochloropsis* species
4. To develop modified strains of *Nannochloropsis* with reduced dark respiration using CRISPR/Cas9 tools (Table 1)
5. To develop modified strains of *Nannochloropsis* with reduced photorespiration using CRISPR/Cas9 tools (Table 1)
6. To demonstrate scale-up potential of modified *Nannochloropsis* using SNL Algal Testbed (Figure 1)

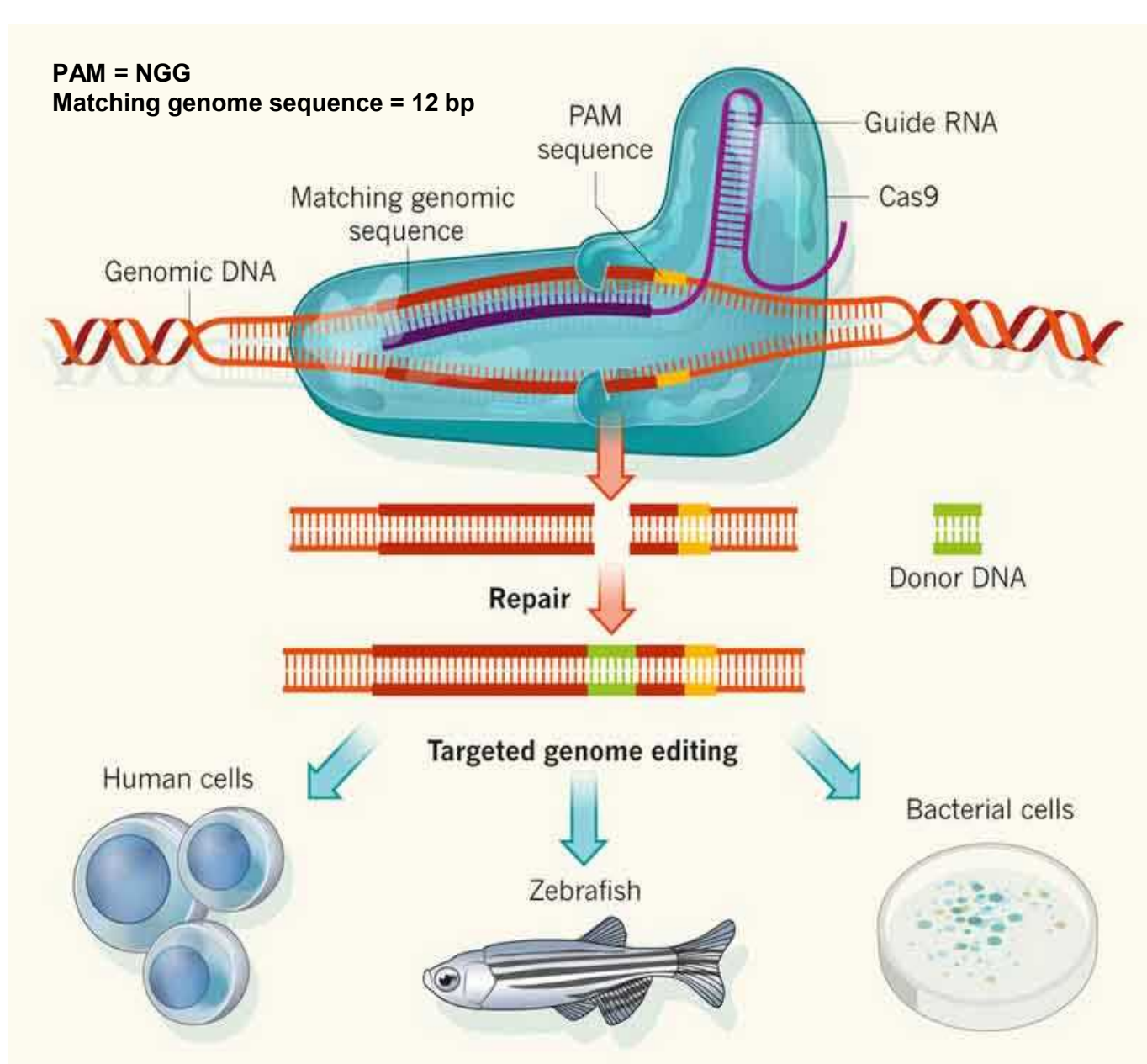


Figure 4. Schematic of CRISPR/Cas9 genome editing in eukaryotes.³

Table 1. List of genetic targets in *Nannochloropsis gaditana*

Pathway	Target Description	Gene Target
Laminarin degradation	Laminarinase	Nga02655
TAG degradation	TAG lipase CrLIP1 (<i>Chlamydomonas reinhardtii</i>)	Nga01367
	TGL3/TGL4/TGL5 (yeast) SDP1 (<i>Arabidopsis thaliana</i>)	Nga03028
Dark respiration	Cytochrome c oxidase COX1	Nga50030, Nga50029
	Alternative oxidase AOX1	Nga03289
Photorespiration	Glycolate dehydrogenase	glcD (<i>E. coli</i>)
	Glycolate carboxylase	glcE (<i>E. coli</i>)
	Tartronic semialdehyde reductase	glcF (<i>E. coli</i>)

Results: Genetic Tool Development

- Transformation (transfer of DNA across the cell wall) is challenging for algae due their tough algal cell walls.
- High voltage electroporation has been optimized for delivery of DNA into *Nannochloropsis* species (Figure 5).
- However, transformation across different species of *Nannochloropsis* is highly variable (Figure 6).

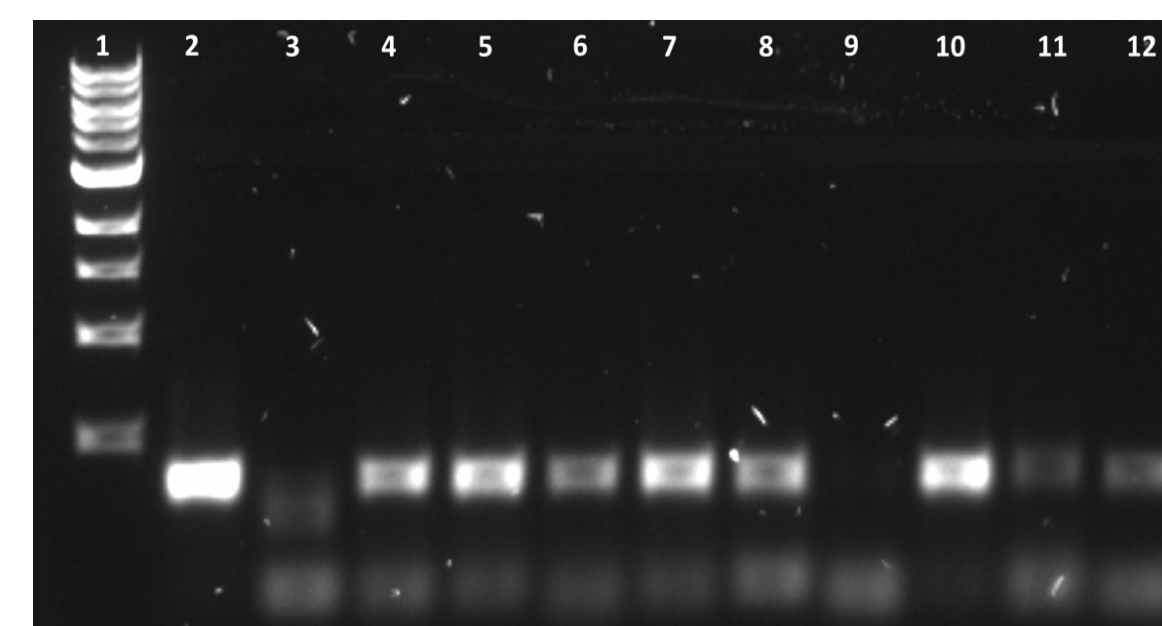


Figure 5. PCR screening of *ble* integrated into the *N. gaditana* genome via high voltage electroporation. Lane 1 = 1kb ladder, Lane 2 = pJTM008 (positive control), Lane 3 = *N. gaditana* gDNA, Lanes 4-12 = zeocin resistant colonies. Expected *ble* fragment = 368 bp.

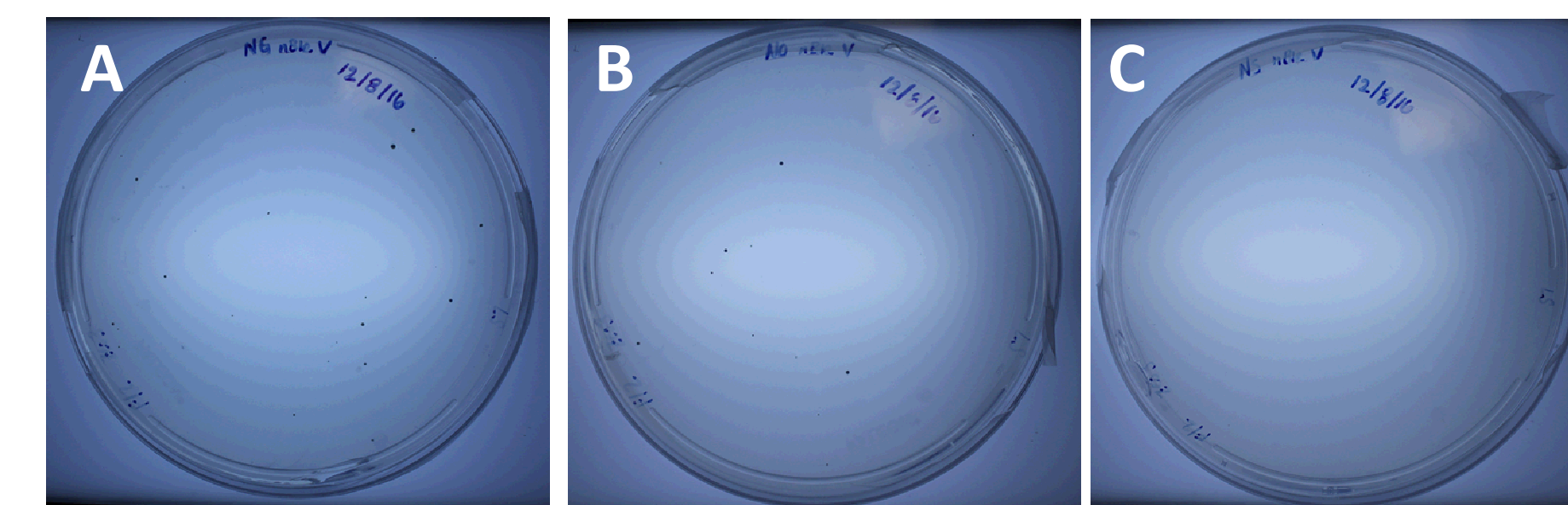


Figure 6. Zeocin resistant colonies of *N. gaditana* (A), *N. oceanica* (B), and *N. salina* (C) after high voltage electroporation of plasmid containing *ble*.

Table 2. Transformation of linearized plasmids into *N. gaditana*

Plasmid Construct	Number of Transformed Colonies
pBleT	19
pBleV	23
pBleT-Cas9	6
pBleV-Cas9	9

- Introduction of Cas9 leads to reduced transformation efficiencies, suggesting Cas9 expression is toxic in *Nannochloropsis* (Table 2).

Results: Physiological Measurements of Dark Respiration and Photorespiration

- Addition of SHAM (salicylhydroxamic acid), inhibitor of dark respiration, resulted in reduced growth (Figure 7).
- SHAM may also inhibit glycolate metabolism, as demonstrated in other algae.⁴

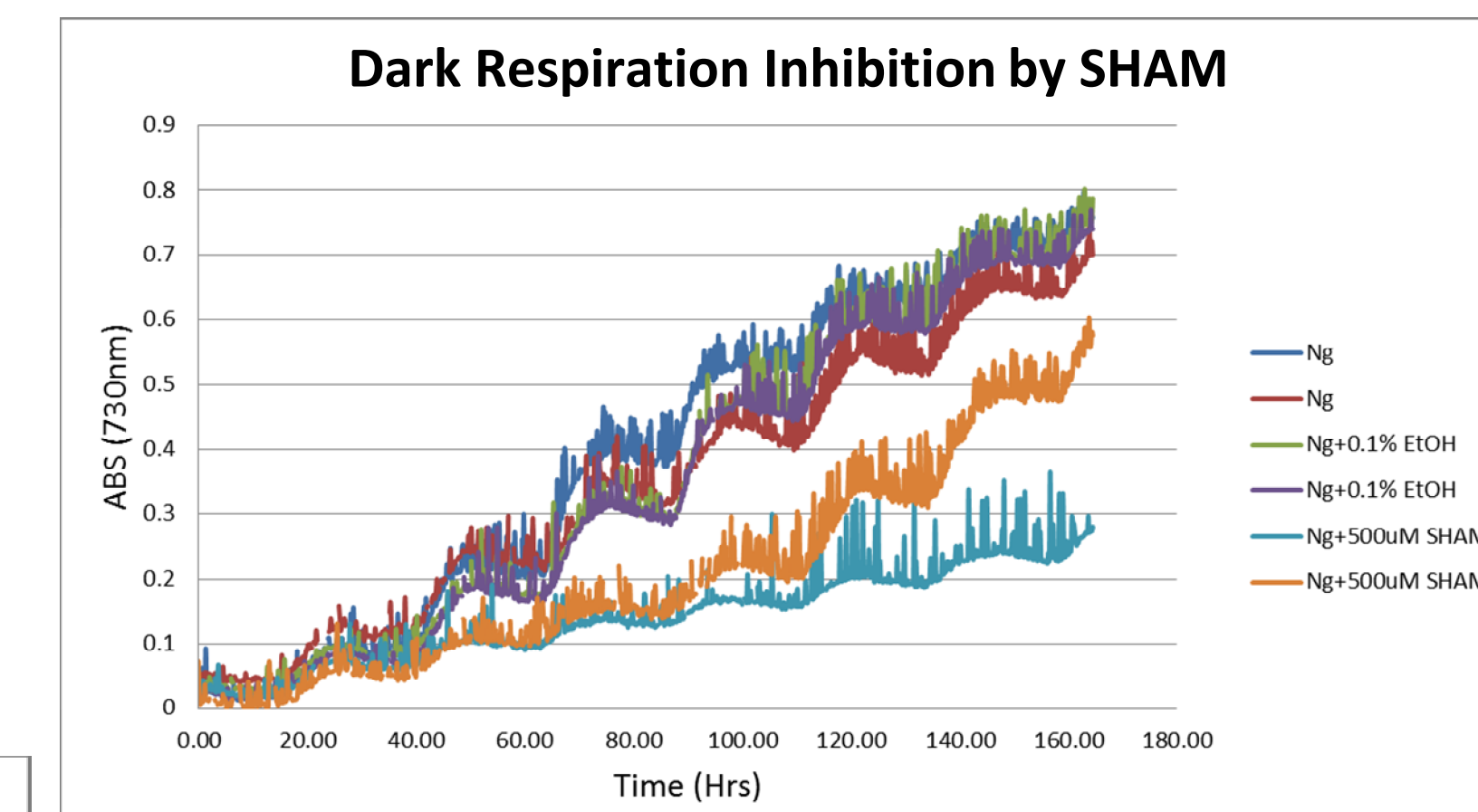


Figure 7. Growth of *N. gaditana* with addition of 500 μM of SHAM. Growth temperature = 25°C; Illumination = 1000 μE; 12h:12h Light/Dark.

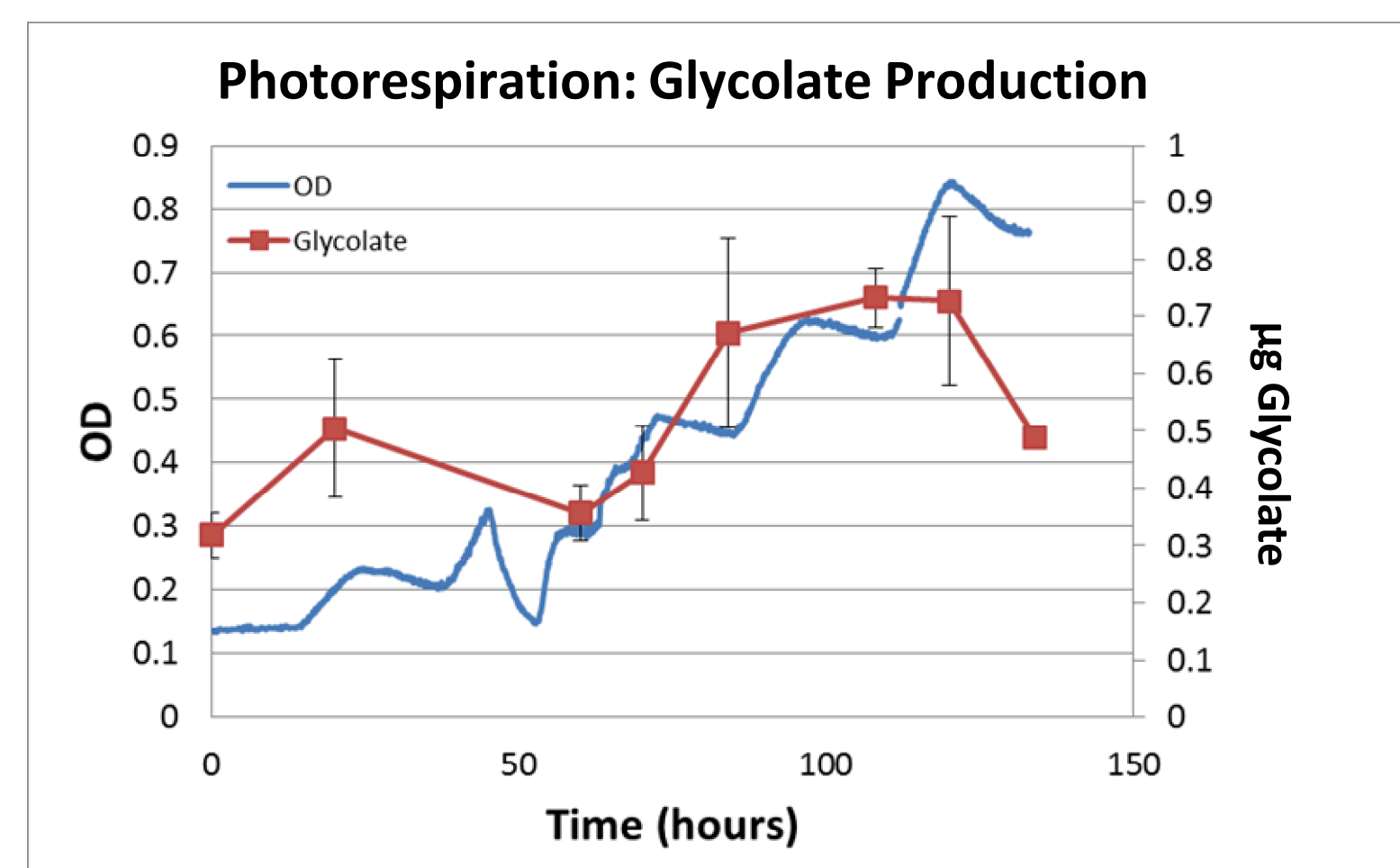


Figure 8. Growth of *N. gaditana* (blue) and glycolate excretion (red). Growth temperature = 25°C; Illumination = 1000 μE; 12h:12h Light/Dark.

Conclusions and Future Work

- Cas9 appears to be toxic to *Nannochloropsis* species.
- Various promoter constructs will be tested to optimize Cas9 expression (Joint Genome Institute DNA Synthesis project).
- Genetic modification of dark respiration genes will determine the effect of dark respiration on carbon loss.
- Photorespiration losses in *N. gaditana* may be low.

References and Acknowledgements

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3. Charpentier and Doudna. (2013) Nature. 495: 50-1.
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