

***Synechococcus* sp. PCC 7002: A Cyanobacterial Chassis for Advanced Biofuel Production**

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Goal of Sandia's Bioscience Research Foundation

We analyze, understand, and **control** the functions of biological systems to meet National Security challenges in biodefense and biofuels production.

- Analyze: Analyze samples to obtain the most relevant biological data for biodefense and biofuels.
- Understand: Integrate data sets and use them to build predictive models of biological systems.
 - Decipher molecular mechanisms relevant to biofuels or biodefense.
 - Recognize biological correlations and dependencies to identify unique signatures, markers and/or design new pathways.
- Control: **Rationally modify the functions of biological systems.**
 - Bioenergy: Demonstrate and optimize efficient, cost-effective production of **drop-in fuels** and co-products from the conversion of lignocellulose or algae.
 - BEID: Design and deploy countermeasures and materials which are safer and/or more effective than traditional solutions

Cyanobacterial-Based Biofuel Production

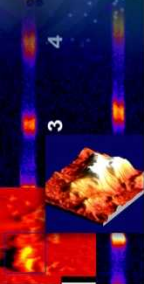
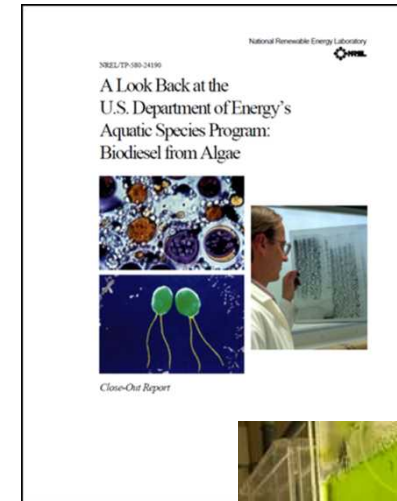
Desirable strain traits

- Easily transformed
- Homologous recombination – targeted genome integration
- Gene expression not complicated by RNAi
- Established genetic tools
- Fast growth rates and strain robustness

Process design advantages

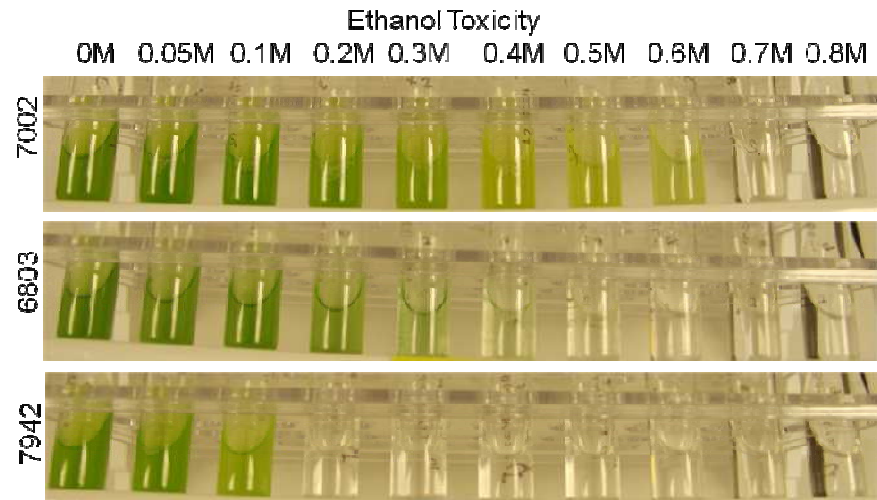
- Product excretion enables continuous production
- Biomass harvesting not required
- Lower nutrient requirements (N&P)

Cyanobacterial-based biofuel production has numerous advantages compared to traditional algal biofuels.



Synechococcus sp. PCC 7002

- Model cyanobacterium
 - Genetic tools available
 - Genome sequence
- Rapid doubling time (2.6 – 4 hrs)
- Temperature tolerance
 - Grows under 22-40°C
- Salt tolerance
 - Survives > 1.5 M NaCl (2.5x sea water)
- High light tolerance
 - Survives light intensities > 4.5 mE m⁻² s⁻¹ (2x peak sunlight)
- Biofuel tolerance
 - Grows in ethanol concentrations up to 0.6 M

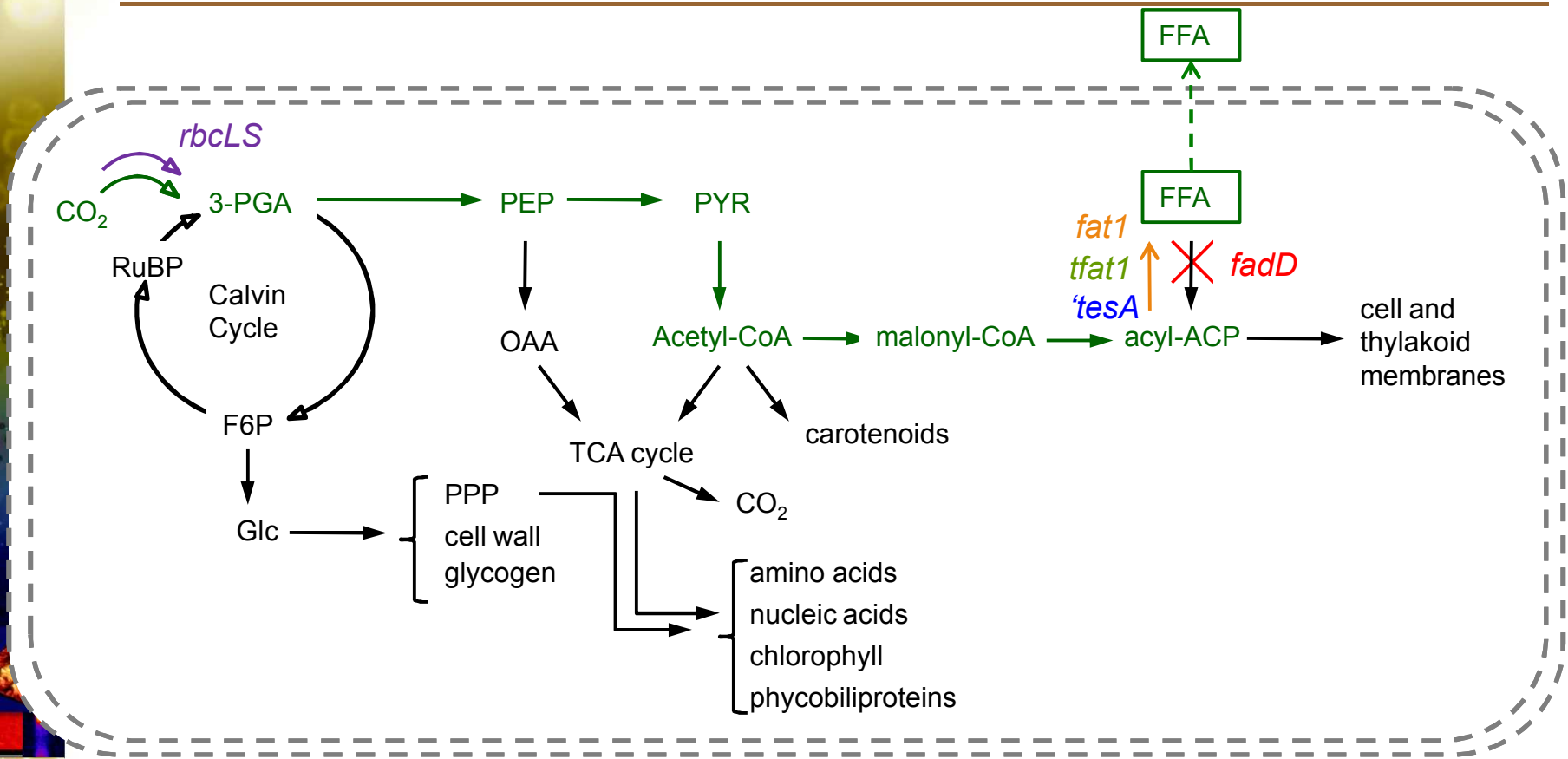


Synechococcus sp. PCC 7002 has many desirable traits for advanced biofuel production.

Engineering *Synechococcus* sp. PCC 7002 for Advanced Biofuel Production

- Traditional Metabolic Engineering of *Synechococcus* sp. PCC 7002 for Free Fatty Acid Production (Funding: Truman Fellowship)
- Systems-Level Metabolic Engineering of *Synechococcus* sp. PCC 7002 for Alkane Production (Funding: Early Career LDRD)

Traditional Metabolic Engineering of *Synechococcus* sp. PCC 7002 for FFA Production

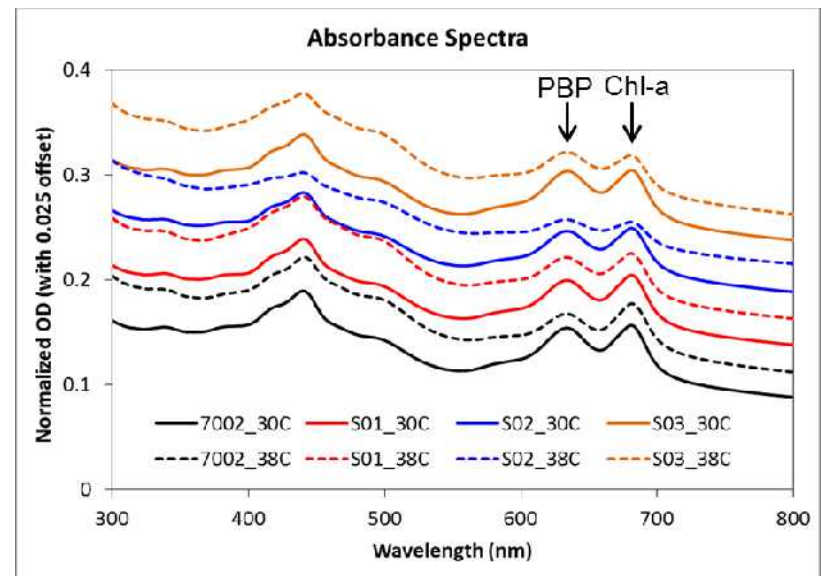
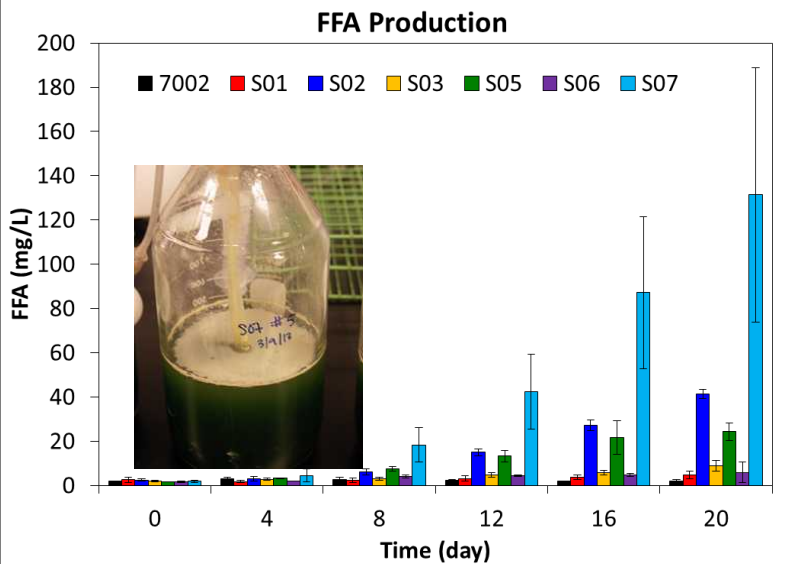
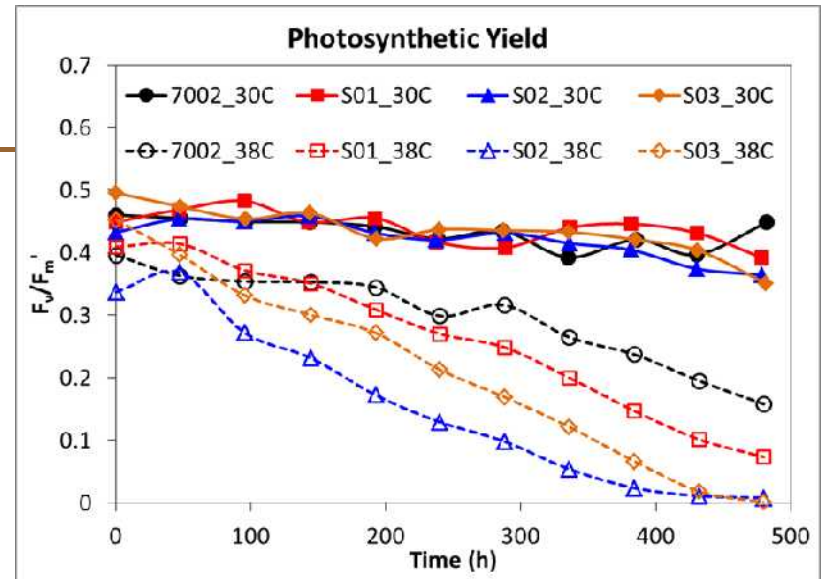


7002: wild type; S01: $\Delta fadD$; S02: $\Delta fadD$, *'tesA*; S03: $\Delta fadD$, *fat1*; S05: $\Delta fadD$, *tfat1*; S06: $\Delta fadD$, P_{trc}-*'tesA*, *rbcLS*; S07: $\Delta fadD$, *'tesA*, P_{psbA1}-*rbcLS*

Six engineered strains of *Synechococcus* sp. PCC 7002 were constructed for FFA production.

FFA Production

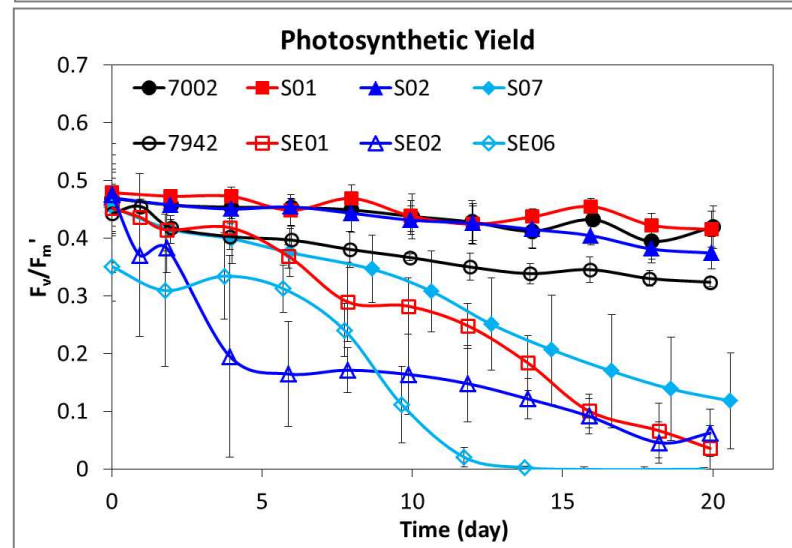
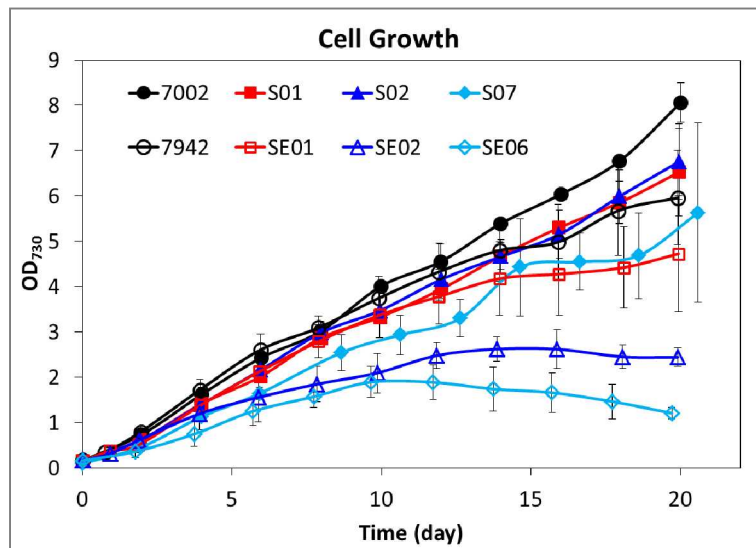
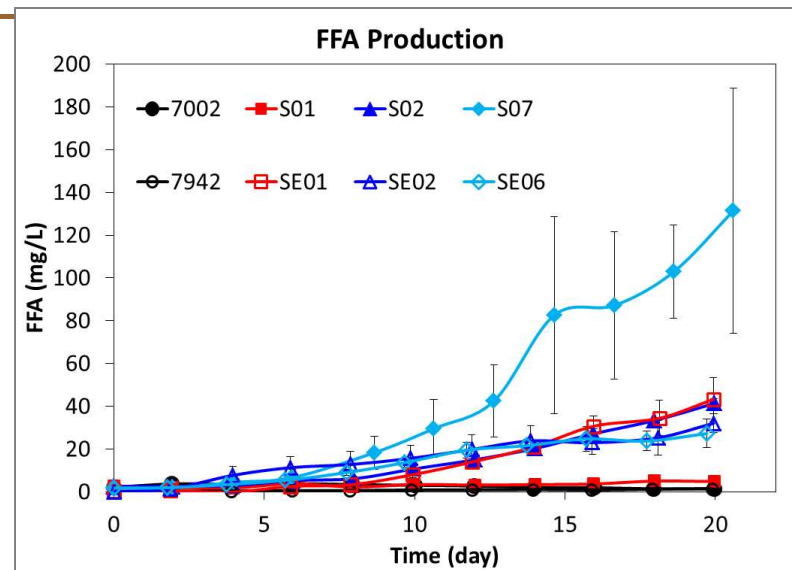
- Maintenance of high photosynthetic yields and photosynthetic pigments (PBP, Chl-a) is temperature dependent
- Significant increase in FFA production with overexpression of RuBisCO from *psbAI* promoter



High temperatures and high FFA production had negative impacts on the physiology of the host, *Synechococcus* sp. PCC 7002.

Host Selection: *Synechococcus* sp. PCC 7002 vs *Synechococcus elongatus* PCC 7942

- Highest FFA producing strains:
 - 7002: S07 (131 mg/L)
 - 7942: SE01 (43 mg/L)
- Improved growth and photosynthetic yields for 7002 strains

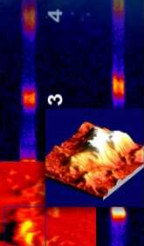
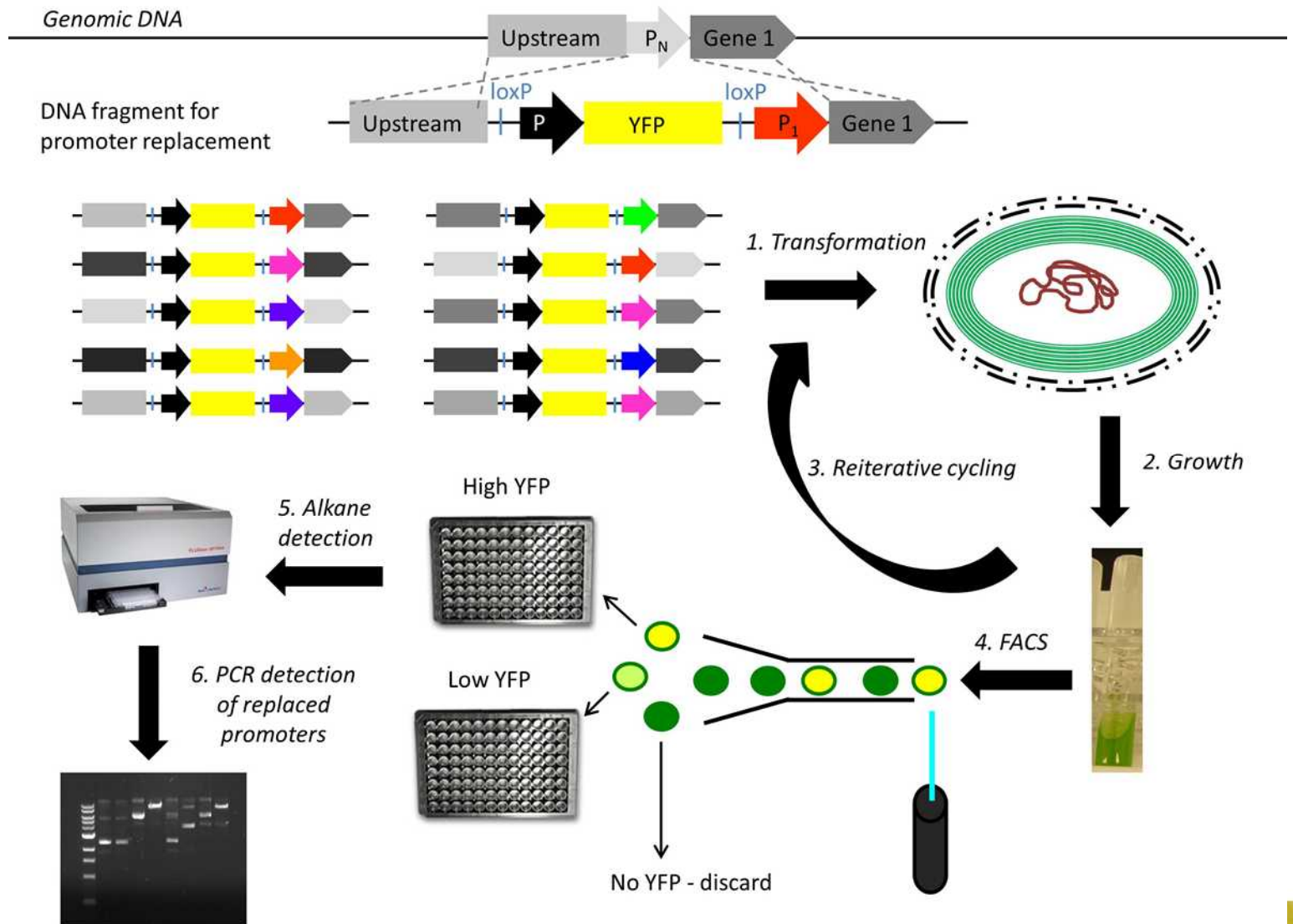


Synechococcus sp. PCC 7002 is an advantageous host for FFA production.

Traditional Metabolic Engineering: Conclusions

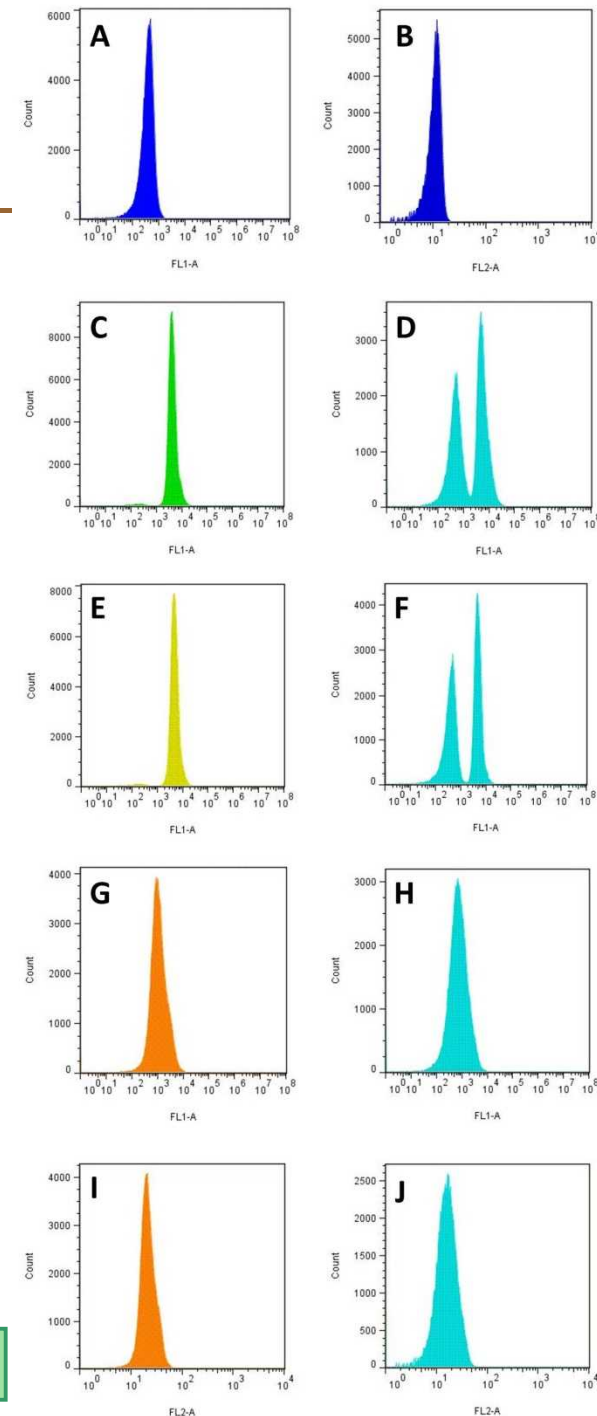
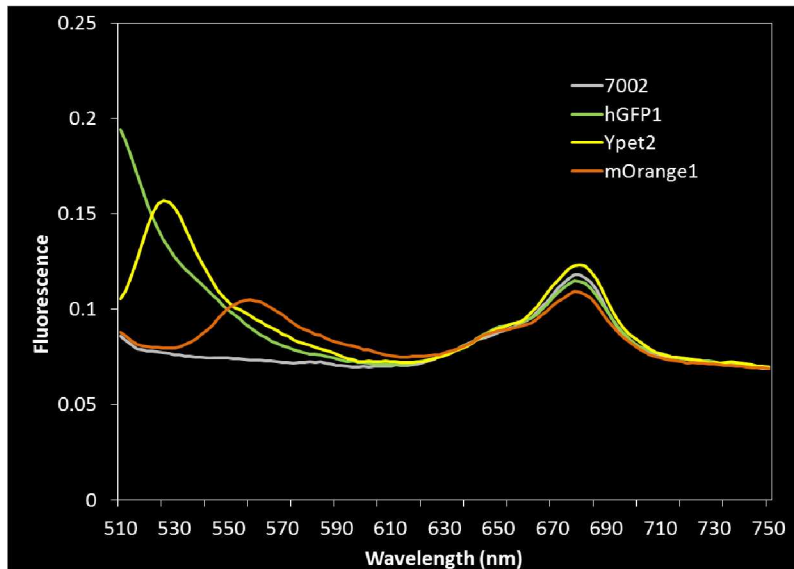
- *Synechococcus* sp. PCC 7002 is a better host for FFA production compared to *S. elongatus* PCC 7942
- Carbon fixation (RuBisCO) limits FFA biosynthesis in *Synechococcus* sp. PCC 7002
- Gene expression is critical for optimizing FFA production
- FFA production has complex, systems-level interactions (growth, photosynthetic yield, pigments) that cannot be optimized by traditional pathway engineering

Development of a Systems-Level Metabolic Engineering Technique for *Synechococcus* sp. PCC 7002

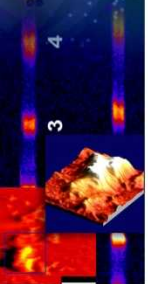


Fluorescent Protein Reporter Selection

- 3 fluorescent proteins investigated: GFP, YFP, and OFP
- YFP has the best combination of :
 - high signal intensity
 - minimal overlap with cyanobacterial photosynthetic pigments
 - good flow cytometer peak separation (488nm excitation laser)
 - excitation (514nm) and emission (527nm) peaks

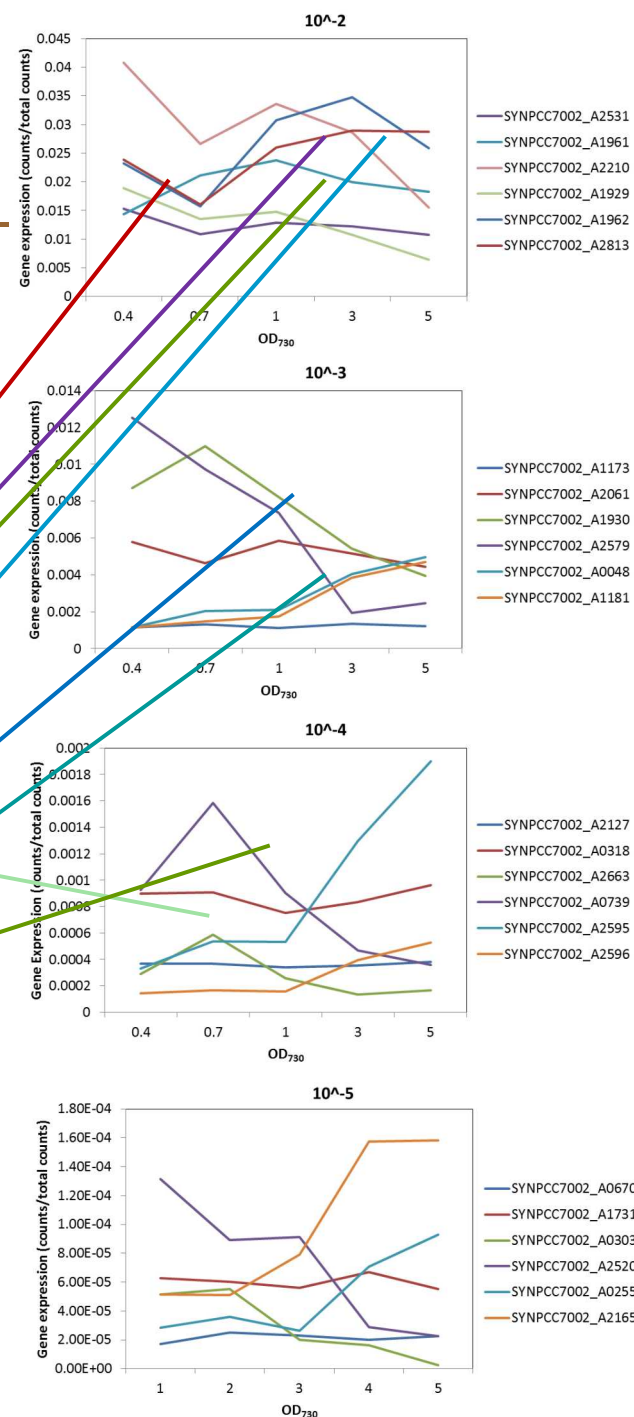
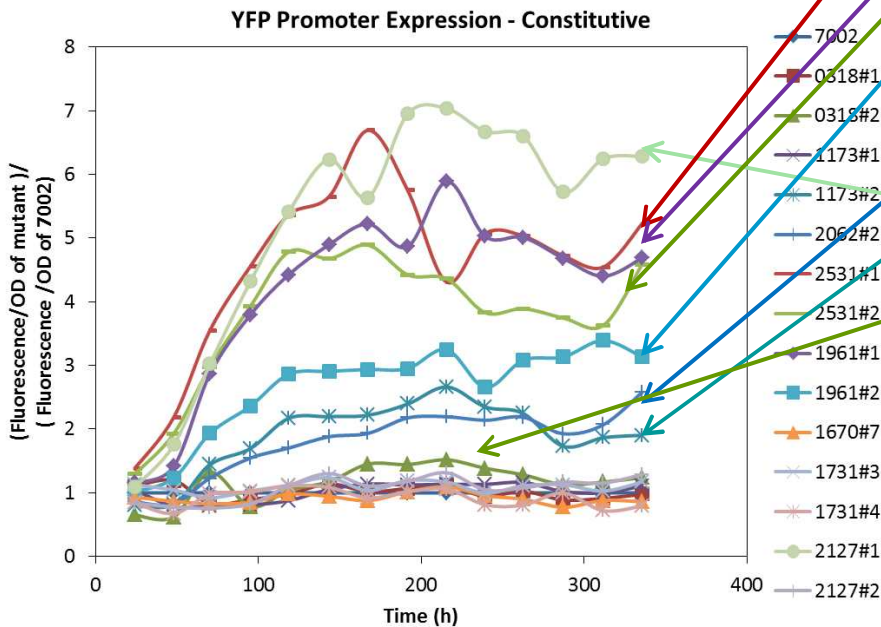


YFP is a good reporter in Synechococcus sp. PCC 7002.



Promoter Characterization in *Synechococcus* sp. PCC 7002

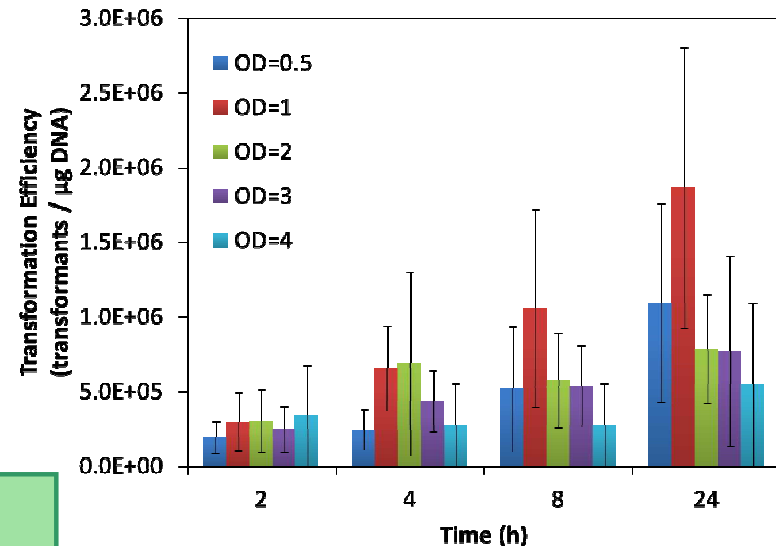
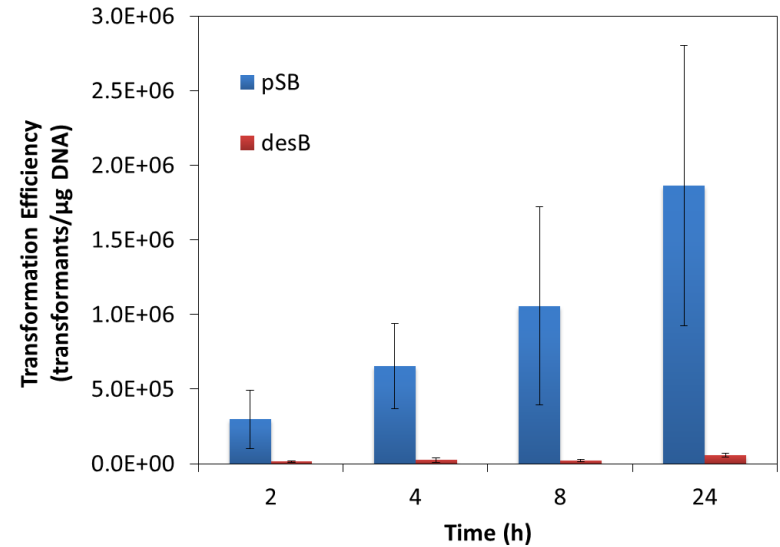
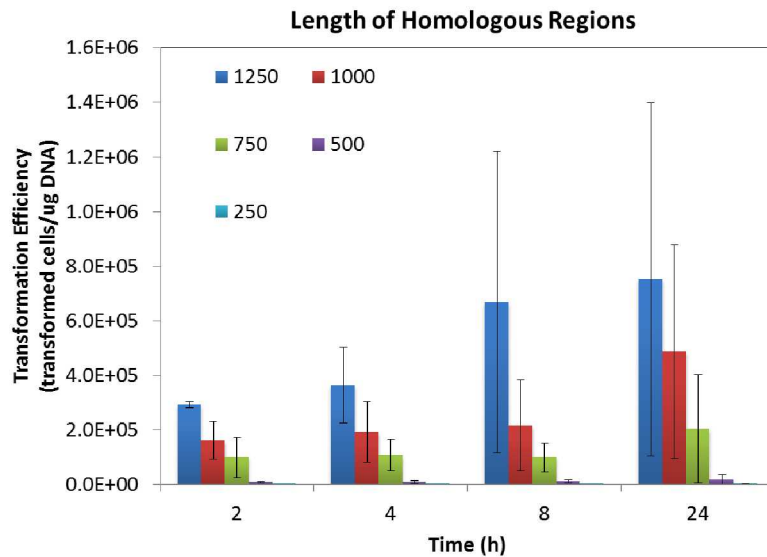
- Promoters selected from previously published RNAseq data (Ludwig, Bryant. 2011. *Frontiers in Microbiol.* 2:41.)
 - Expression levels: 10^{-5} to 10^{-2}
 - Regulatory patterns: constitutive, linear growth phase, and stationary phase
- Linked promoters to YFP for characterization under continuous light and diurnal conditions



Constitutive promoters were identified, cloned, and characterized in *Synechococcus* sp. PCC 7002.

Optimizing Transformation in *Synechococcus* sp. PCC 7002

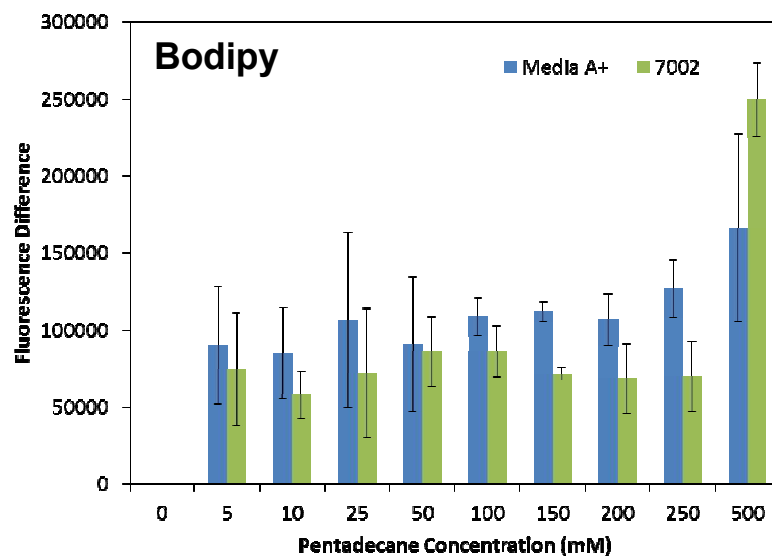
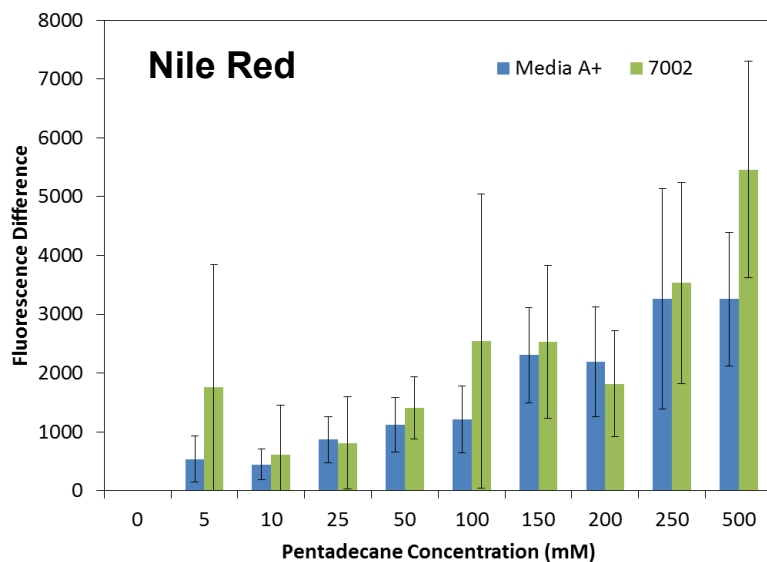
- Exonuclease activity present in *Synechococcus* sp. PCC 7002
- Cell concentration: $OD_{730} = 1.0$
- Length of homology for homologous recombination: 1kb
- Genome copy number (ploidy): lowest at 34°C , $60 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$, $OD_{730} > 3.0$



Factors for high transformation efficiency in *Synechococcus* sp. PCC 7002 were optimized.

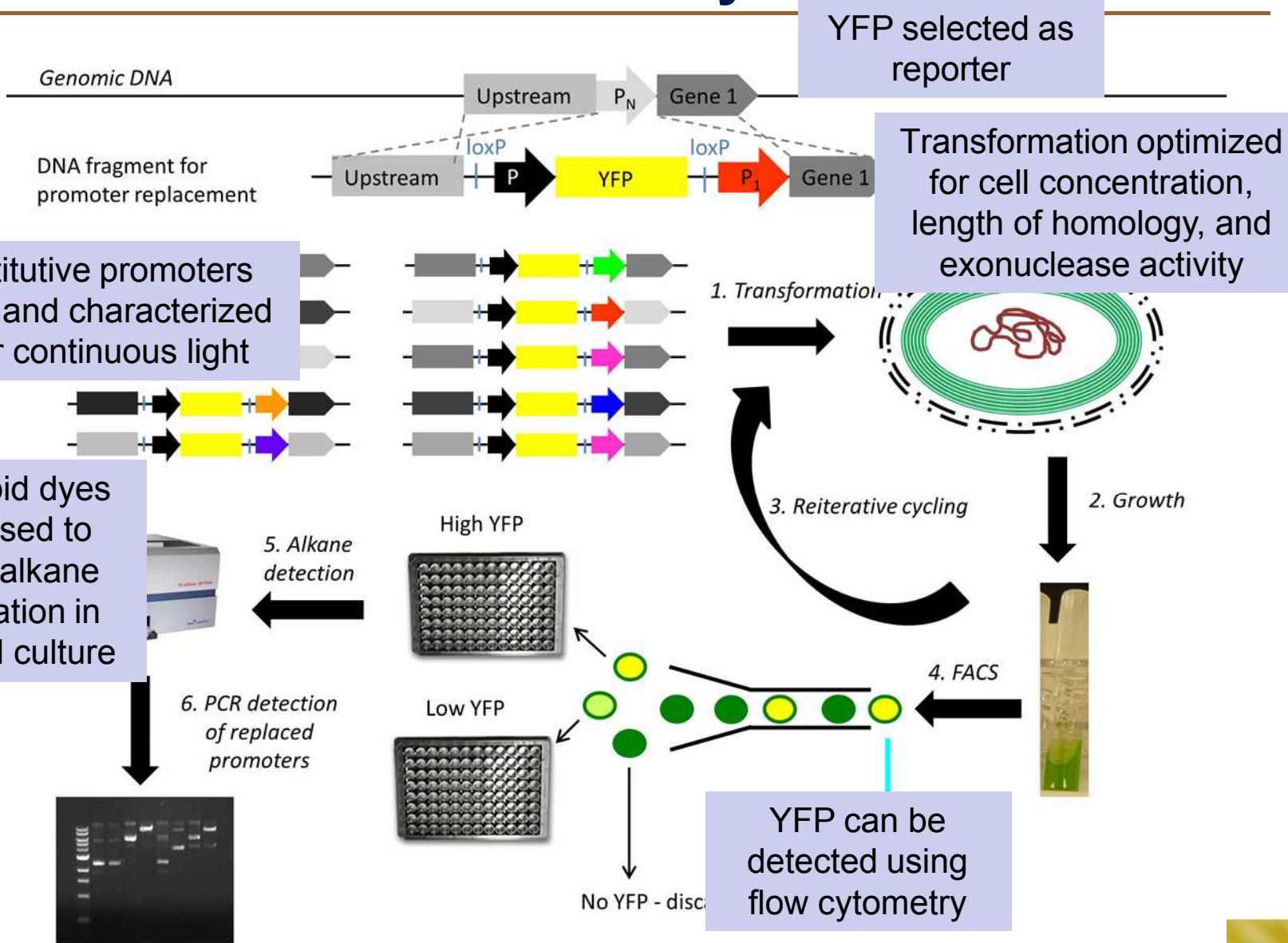
High-Throughput Alkane Screening

- 2 neutral lipid dyes: Nile red and bodipy
- Exogenous addition of pentadecane to media or dilute culture ($OD_{730} = 0.1$)
- Nile red is more quantitative
- Bodipy is more sensitive



Neutral lipid dyes can be used for high-throughput screening of alkane production.

Systems-Level Metabolic Engineering: Summary



Systems-Level Metabolic Engineering: Future Work

- Clone and characterize linear growth phase and stationary phase expression promoters
- Determine conditions for minimal genome copy number
- Analyze fluorescence detection of multiple YFP copies
- Characterize neutral lipid stains for intracellular alkane production
- Design synthetic DNA constructs for systems-level metabolic engineering (JGI DNA synthesis proposal pending)
- Demonstrate systems-level metabolic engineering for improving alkane production in *Synechococcus* sp. PCC 7002

Publications

Anne M Ruffing and Christine A Trahan. Biofuel Toxicity and Mechanisms of Biofuel Tolerance in Three Model Cyanobacterial Strains. To be submitted to *Algal Research* (June 2014).

Anne M Ruffing. Improved Free Fatty Acid Production in Cyanobacteria with *Synechococcus* sp. PCC 7002 as Host. To be submitted to *Frontiers in Bioengineering and Biotechnology* (March 2014) – Research Topic: Cyanobacteria, the green *E. coli*.

Anne M Ruffing. RNA-Seq analysis and targeted mutagenesis for improved free fatty acid production in an engineered cyanobacterium. *Biotechnology for Biofuels*. 2013. 6:113.

Anne M Ruffing. --Borrowing genes from *Chlamydomonas reinhardtii* for free fatty acid production in engineered cyanobacteria. *Journal of Applied Phycology*. 2013. 25(5): 1495-1507.

Anne M Ruffing. Metabolic Engineering of Hydrocarbon Biosynthesis for Biofuel Production. Book chapter in *Liquid, Gaseous and Solid Biofuels – Conversion Techniques*. Intech, Rijeka, Croatia. 2013.

Anne M Ruffing and Howland DT Jones. Physiological effects of free fatty acid production in genetically engineered *Synechococcus elongatus* PCC 7942. *Biotechnology and Bioengineering*. 2012. 100(9): 2190-2199. (Cover, Spotlight)

Anne M Ruffing. Engineered Cyanobacteria: Teaching an old bug new tricks. *Bioengineered Bugs*. 2011. 2(3).



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