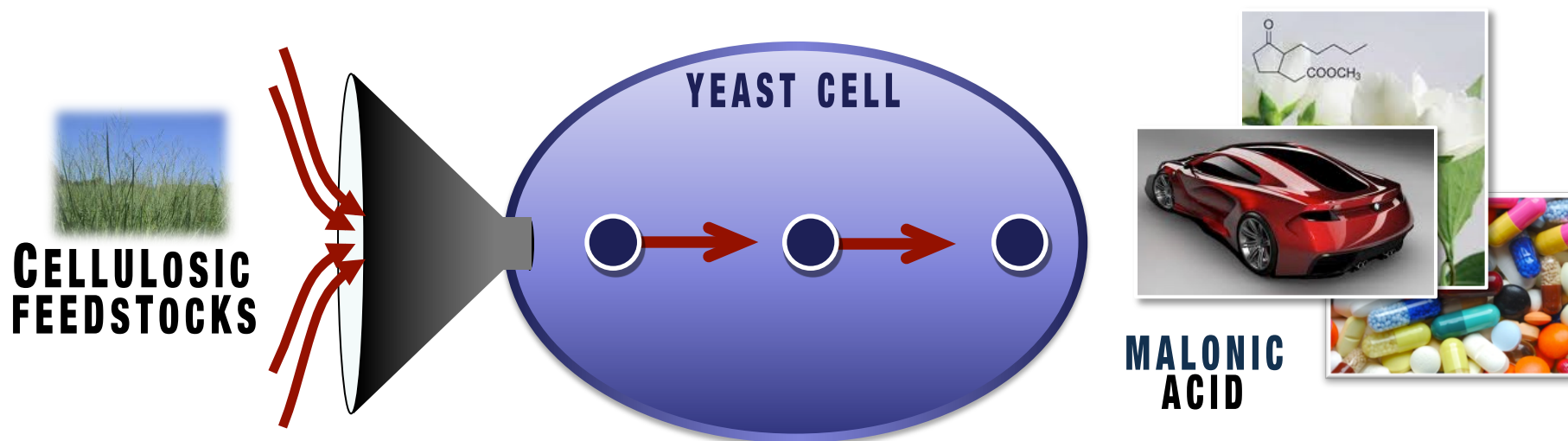


LYGOS

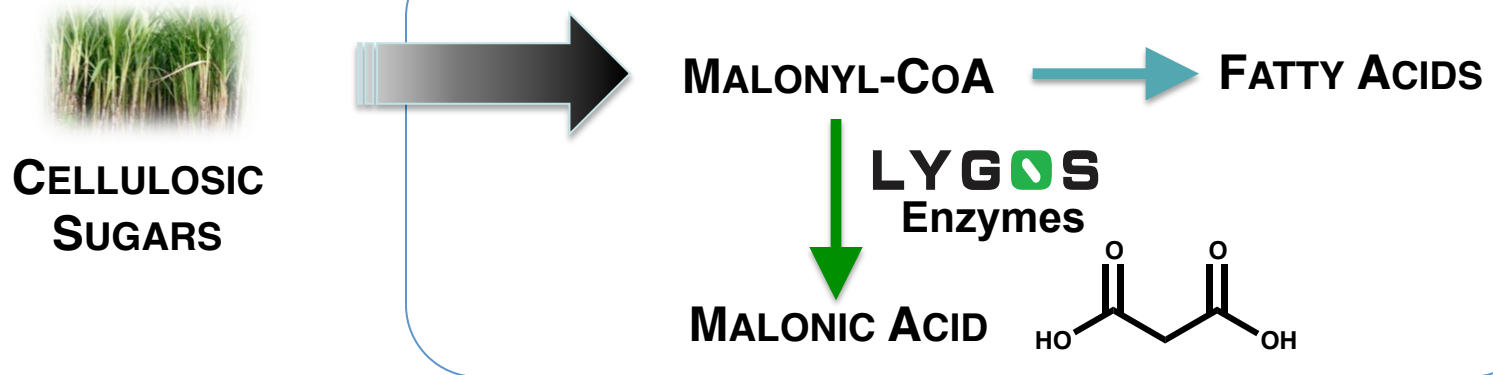
DESIGN & OPTIMIZATION OF A BIOCHEMICAL PRODUCTION PLATFORM WITH BIOSENSOR-GUIDED SYNTHETIC EVOLUTION



March 23, 2015
DOE BETO Conversion Review

Eric J. Steen, Ph.D.
Lygos, Inc.

Malonic Acid Is An Ideal Biological Product



- **Malonyl-CoA lies on the carbon superhighway in biology**
 - Pathway is compatible with all available, low-cost feedstocks
 - Malonyl-CoA is basis of fatty acid production (90%+ yields reported)
- **Malonic acid is an ideal molecule to produce biologically**

Theoretical Yield (lb-product / lb-sugar): 1.73



Goal Statement

Develop an integrated approach to biochemical pathway optimization for production of malonic acid & demonstrate path toward commercially-relevant fermentation metrics.

Relevance to US & BETO mission:

- **Reduce dependence on foreign oil**
 - **130 million pounds produced in foreign countries**
- **Increase production of fuels & chemicals from lignocellulosic feedstocks**
- **Decrease greenhouse gas (GHG) emissions**
 - **100 million pounds of CO₂ sequestered**
 - **Eliminate 34 million pounds of sodium cyanide use**
- **Identified as Top 30 bioproduct by DOE**

Quad Chart Overview

Timeline

- Start: Aug 31, 2013
- End: January 1, 2016
- Project is 47% complete by spend (as of 12/31/15)

Budget, \$2.06MM

	FY 13 Costs	FY 14 Costs	Planned FY 15
DOE Funded	\$310,481	\$561,458	\$996,468
Lygos Cost Share	\$34,498	\$62,387	\$99,648

Barriers

- Bt-J: Cost effective production of bioproducts
- Productivity, yield
- Reduce cycle time for biocatalyst development
 - Design tools
 - Construction tools
- Bt-B: biomass variability

Partners

- Cellulosic sugars
- Fermentation Scaling
 - DOE-funded ABPDU

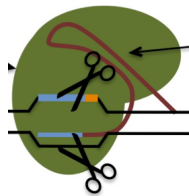
Project Overview & Technical Approach

Overarching Goal: Develop an integrated approach to biochemical pathway optimization for production of malonic acid

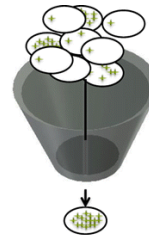
- **Objective #1: Genome-scale metabolic node perturbation**
 - *In silico* biocatalyst design & management software
 - Multiplexed biocatalyst engineering, diversity creation, & targeting tools
- **Objective #2: Deploy biosensor screening method to enable faster identification of improved biocatalysts**
 - Biocatalyst evaluation via biosensor screen
- **Objective #3: Use statistical approach to guide biocatalyst design**
- **Objective #4: Translate benchtop fermentation metrics to pilot scale**



in silico
design



biocatalyst
construction tools



high throughput
screening

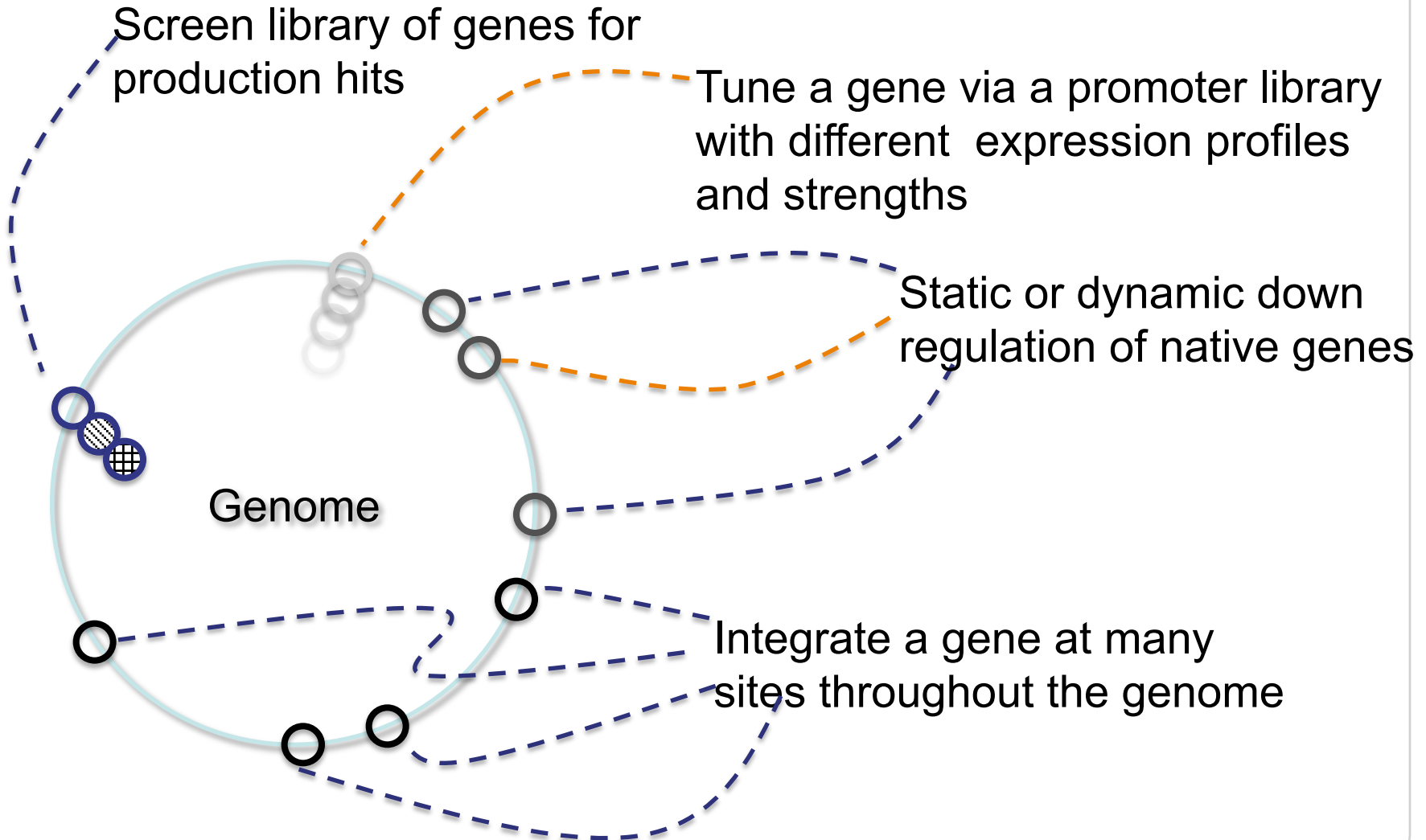


scale up

Project Approach - Management

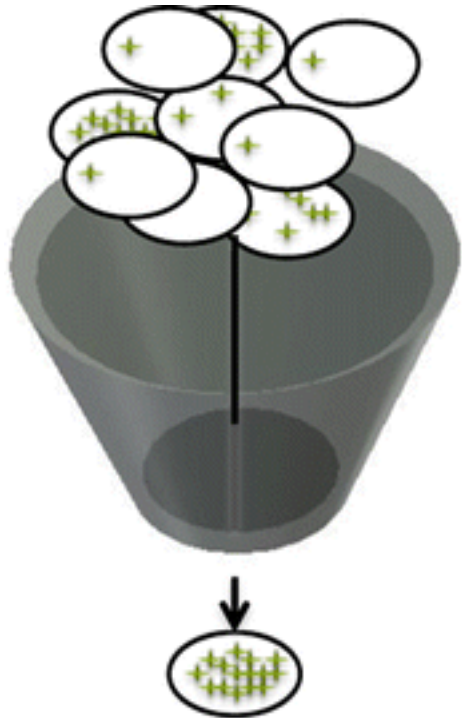
- **Lygos is responsible for biocatalyst design, screening, and fermentation process development**
- **Leveraging expertise of partners**
 - Lignocellulosic feedstocks – commercial providers
 - Fermentations scale up – eg., ABPDU
- **Materials & facility scheduling are planned in advance**
- **Go/no-go decision points support objectives**
 - DNA construction capacity
 - Screening capacity
 - Fermentation metrics (yield, productivity, scale)

Task A: Software tools for designing & constructing biocatalysts



Task B: Biosensor-based screening

Genetic diversity is created which results in individual biocatalysts that produce different levels of malonic acid - but we don't know which ones.



Large, diverse population



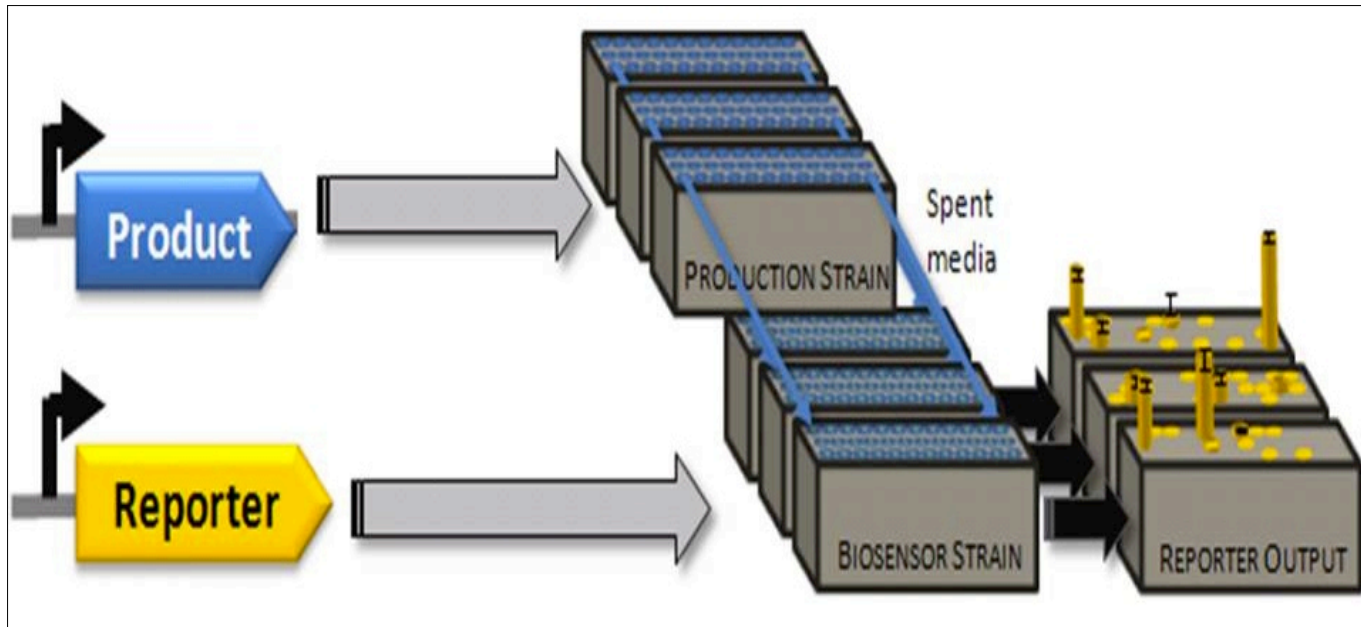
**Identify improved
production strain(s)**

Biosensor screens can offer the most cost effective and direct measurement.

Dietrich JA et al. ACS Synthetic Biology 2013

Task B: Biosensor-based screening

Malonic acid concentration is detected via biosensor strain



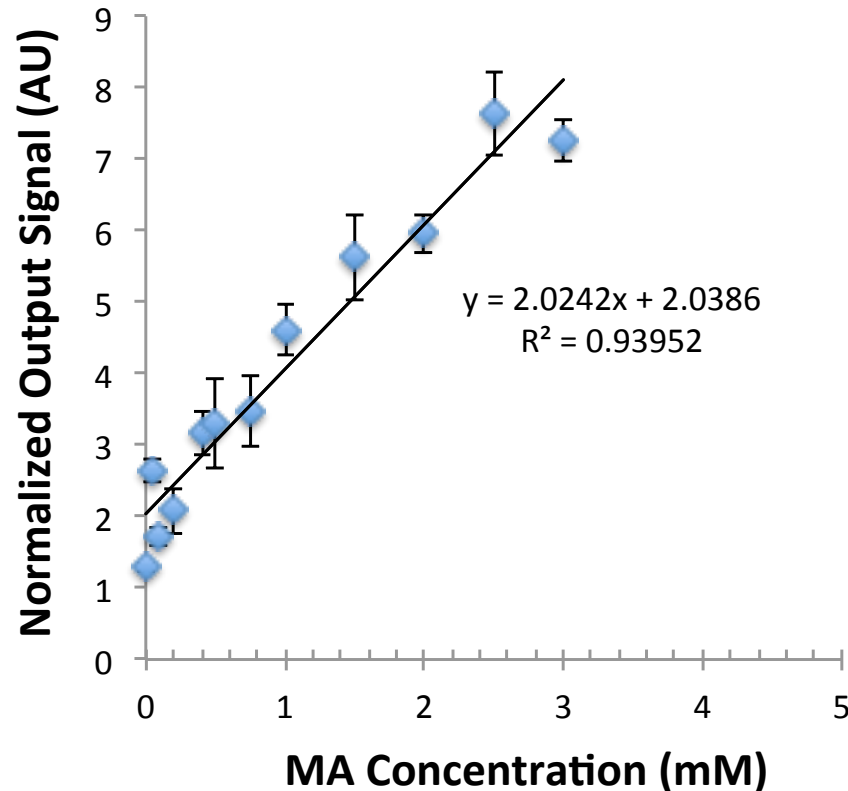
Step 1: Malonic acid (MA) producing strains are grown

Step 2: Spent media is transferred to a standalone biosensor strain culture

Step 3: Biosensor strain grows proportionally to MA concentration

Task B: Biosensor-based screening

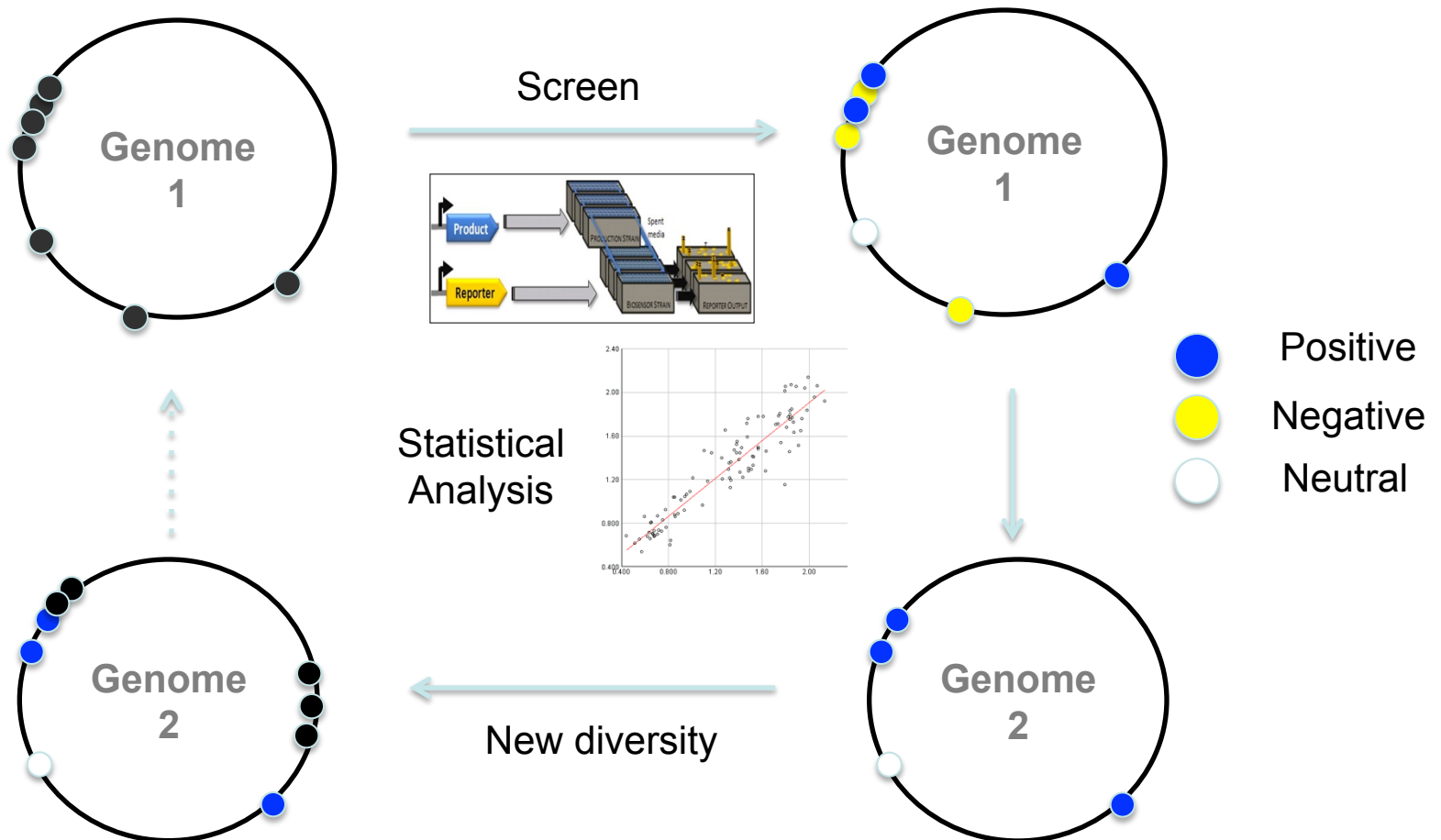
Output of the biosensor screen demonstrates a linear range of malonic acid detection



Improved biocatalysts are identified

Task C: Metabolic hotspot ranking

Metabolic hotspots are ID'd & ranked via statistical analysis in order to more efficiently identify subsequent engineering targets



Task C: Metabolic hotspot ranking

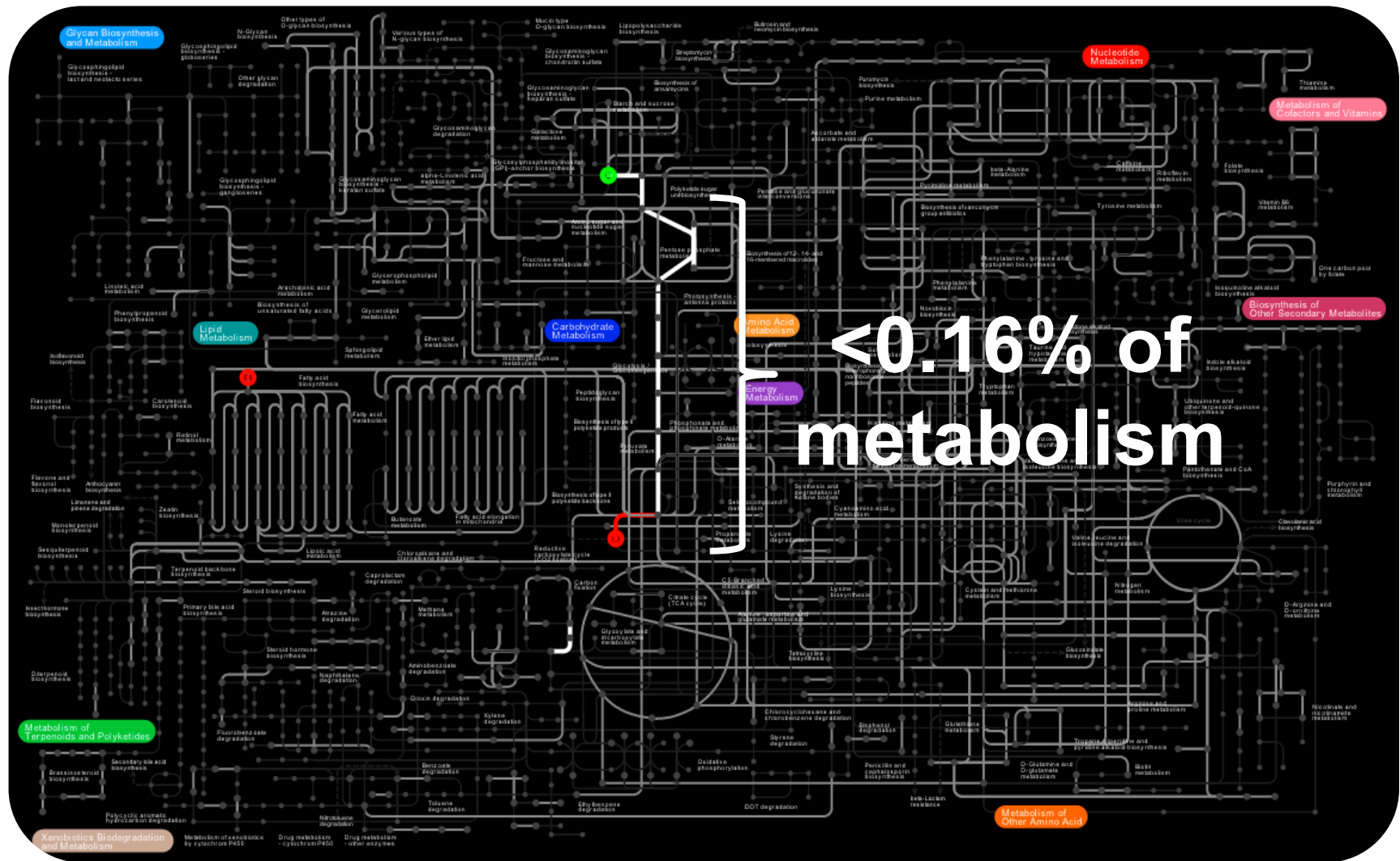
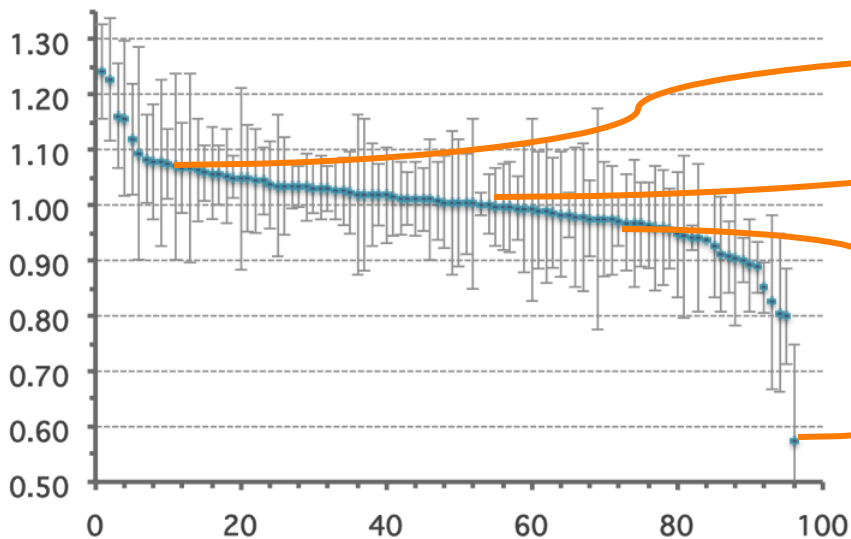


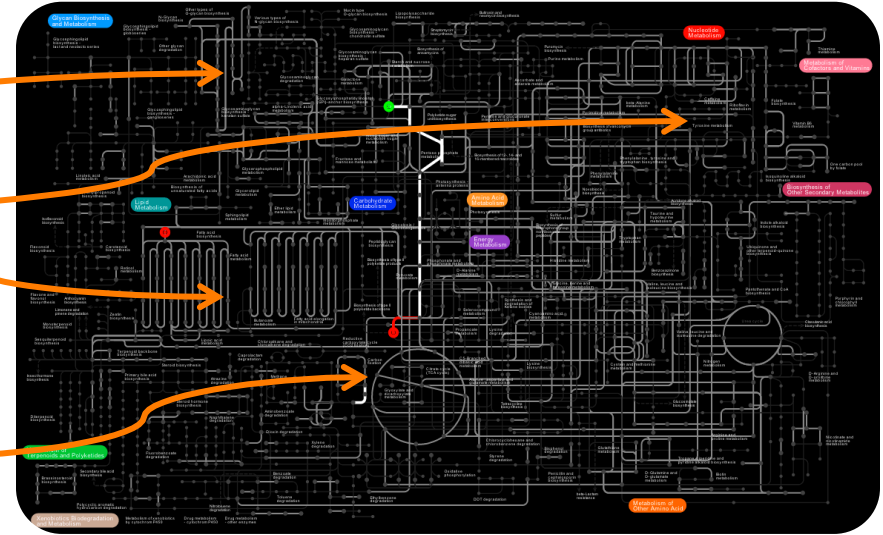
Photo credit: Arkin et al. <http://glamm.lbl.gov/>

Task C: Metabolic hotspot ranking – transcription factors

Fold-Increase



No. of strains

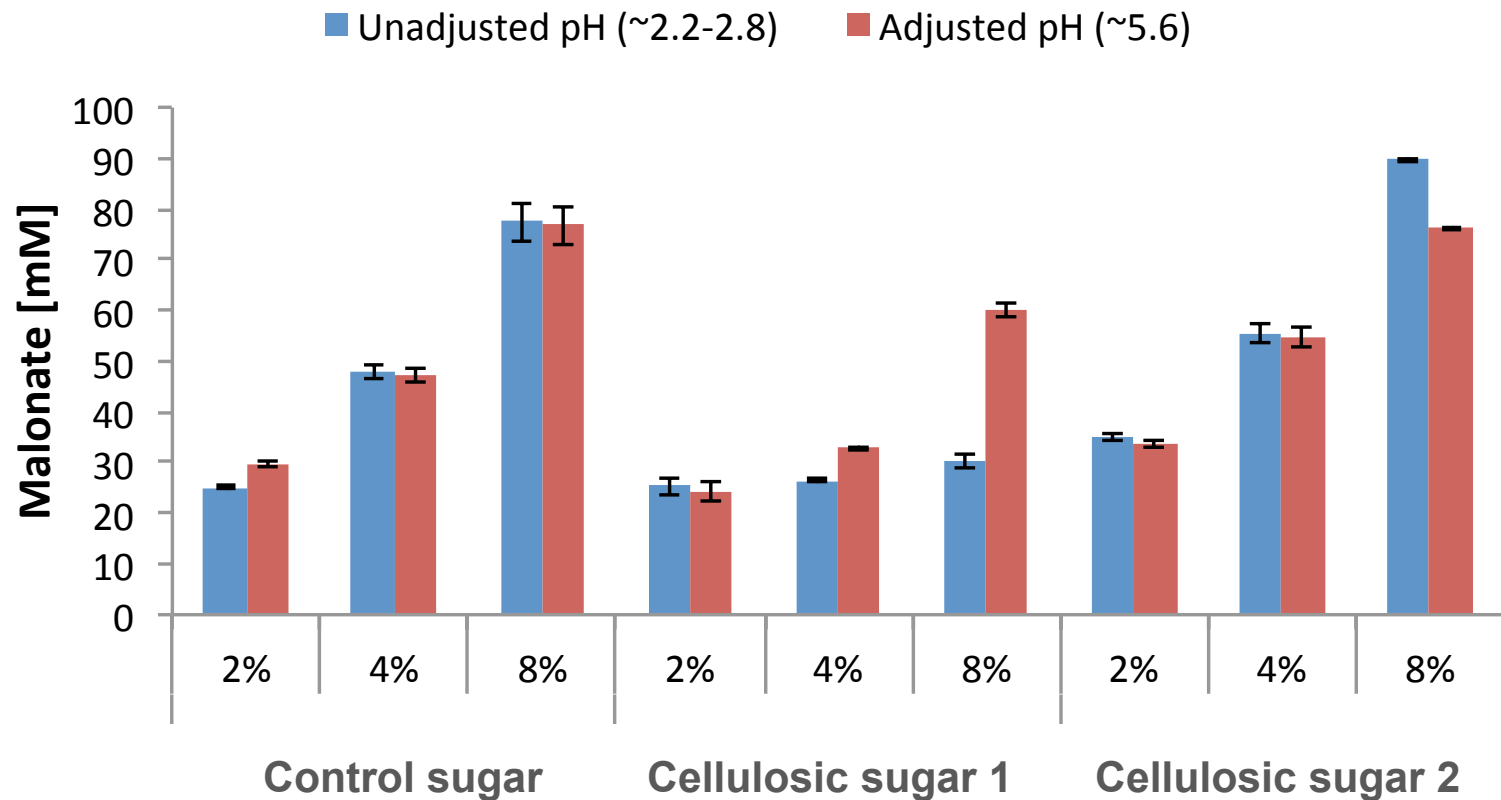


Metabolic Map

- New target identification
- Perturbation refinement
- Metabolic hotspot ID & rank

Guiding iterative metabolic engineering

Task C: Plate Performance Using Cellulosic Sugars



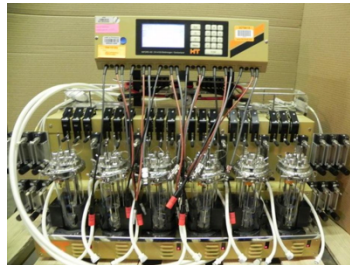
- Obtained samples from Commercial Suppliers
- Similar strain performance between glucose and cellulosic sugars
- Model organisms do not grow (data not shown)

Task D: Translate plate screening metrics to pilot scale

Addressing barrier to translating small scale experiments to relevant fermentation process



0.5 mL



500 mL



300 L

Successful initiation of process development & scaling to 300-L; Achieved 90% of metrics across scales

Relevance

The work in this project promotes:

- Tools for cost effectively building and validating biocatalysts
- Replacing petro products with lignocellulosic-derived products
- Technoeconomic analysis suggests that upon successful commercialization could result in production cost of \$1.60/lb; down from project initiation of >\$600K/lb
- The potential to reduce carbon dioxide emissions related to petroleum produced malonic acid
- Identification of pathway targets may aid in development of other bio-products & fuels
- Synthetic evolution process could be applied to other products

Future Work

- **Package & distribute automated design tools**
- **Further expansion of biocatalyst construction & screening capacity & workflow**
 - Key milestone (10-20 plates up to 100 plates)
- **Achieve plate and tank fermentation metrics**
 - Key milestone in plates (yield, titer, productivity)
 - Key milestone in 300-L (yield, titer, productivity)
- **Downstream purification process** (beyond scope of project)
- **Benchtop Process Integration** (beyond scope of project)
 - Scale up integration risks
 - Feed & Process variation effects
 - Integration of biomass process
- **Derivative Product Development** (beyond scope of project)

Summary

- **Overview:** designing & constructing biocatalysts to consume cellulosic sugars to produce bioproducts (malonic acid)
- **Approach:** deploying synthetic biology techniques to accelerate the path toward commercialization (eg., design tools, screening tools, and validation).
- **Technical Accomplishments:**
 - Design software & tools; Biosensor screen
 - Continuously improved strains are being developed (Y,T,P)
 - Successful initiation of pilot scale fermentations
- **Relevance**
 - Bt-J Cost effective production of bioproducts (\$1.60/lb)
- **Future Work**
 - Attain YTP milestones; expand workflow capacity

Acknowledgements

- **DOE BETO**

- **Jeffrey Dietrich**
- **Jay Keasling (SAB)**
- **Leonard Katz (SAB)**
- **Azadeh Alikhani**
- **Mario Ouellet**
- **Kristy Hawkins**
- **Will Holtz**
- **David Melis**
- **Tina Mahatdejkul-Meadows**
- **Clayton McSpadden**
- **Eric Gates**
- **Karl Fisher**
- **Clem Fortman**
- **Jacinto Chen**
- **Paul Bryan**
- **Neil Renninger**
- **ABPDU**
- **Todd Pray**

Additional Slides

Responses to Previous Reviewers' Comments

- Project has not previously been reviewed

Publications, Patents, Presentations, Awards, and Commercialization

Patents:

- None published relating to this grant, but we are developing inventions previously described in PCT Pub. No. [2013/134424](#) in work supported by the grant.

Presentations:

- Steen EJ. *Synthetic biology for brewing*. Synbiobeta Lecture: Synthetic Biology for Computer Programmers. October 8, 2013, San Francisco, CA.
- Steen EJ. *An industrial perspective on synthetic biology*. Synbiobeta Lecture: Synthetic Biology for Computer Programmers. November 31, 2013 San Francisco, CA.

Commercialization:

- **Financing** – seeking capital for reaching next TRL
- **Offtake** – early discussions for use in a number of markets
- **Internal product development effort**