

Purified Cell-Free Systems as a Metabolic Engineering and Biochemicals Production Platform

SIMB Annual Meeting

Joseph A. Rollin, Christopher W. Johnson, Peter St. John, Gregg T. Beckham

National Renewable Energy Laboratory

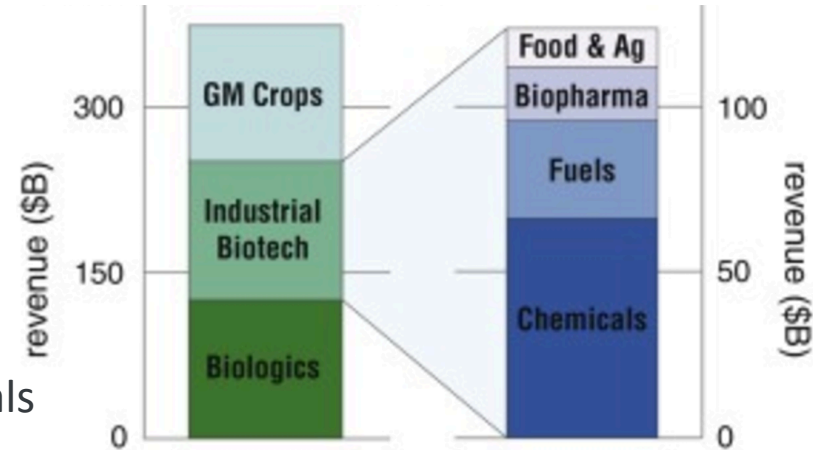
30 July 2017

- The Promise of Industrial Synthetic Biology
- Cell-Free as a Bioconversion Platform
 - Lysate, Purified, and Hybrid Systems
 - Example Application: Biomass Sugars to H₂
 - Key Metrics
 - Challenges and Opportunities
- Cell-Free Metabolic Engineering

Promise of Industrial Synthetic Biology

Industrial applications

- Current:
 - Energy- and carbon-efficient conversions
 - Novel platform molecules
 - Waste stream utilization
- “vast difference: a mixture of valuable chemicals and a valuable mixture of chemicals”



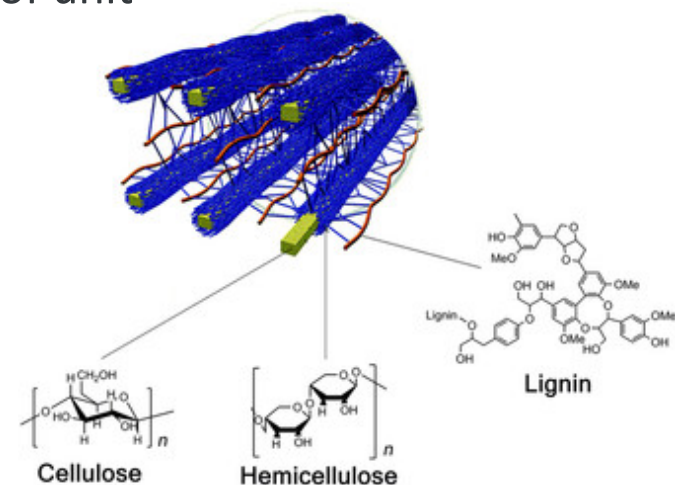
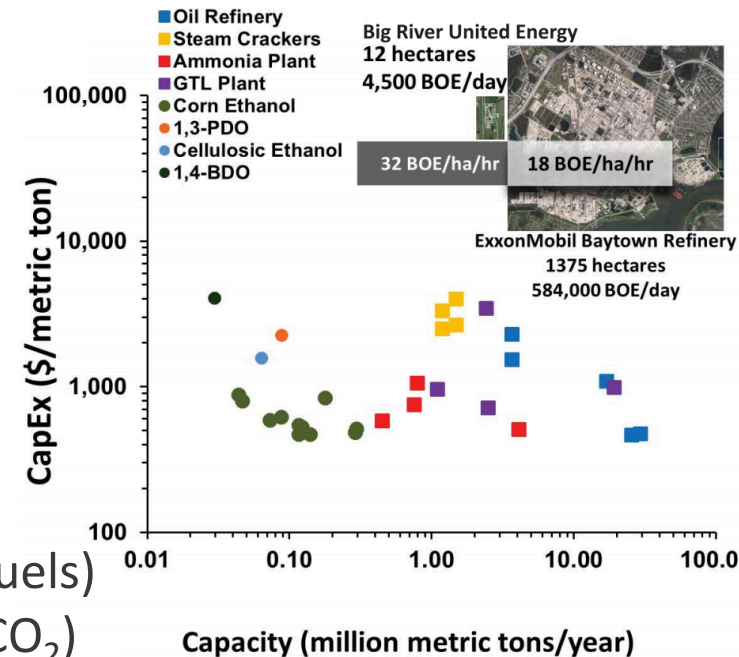
Rogers et al. (2016) Curr Op Biotechnol

Promise of Industrial Synthetic Biology

Industrial applications

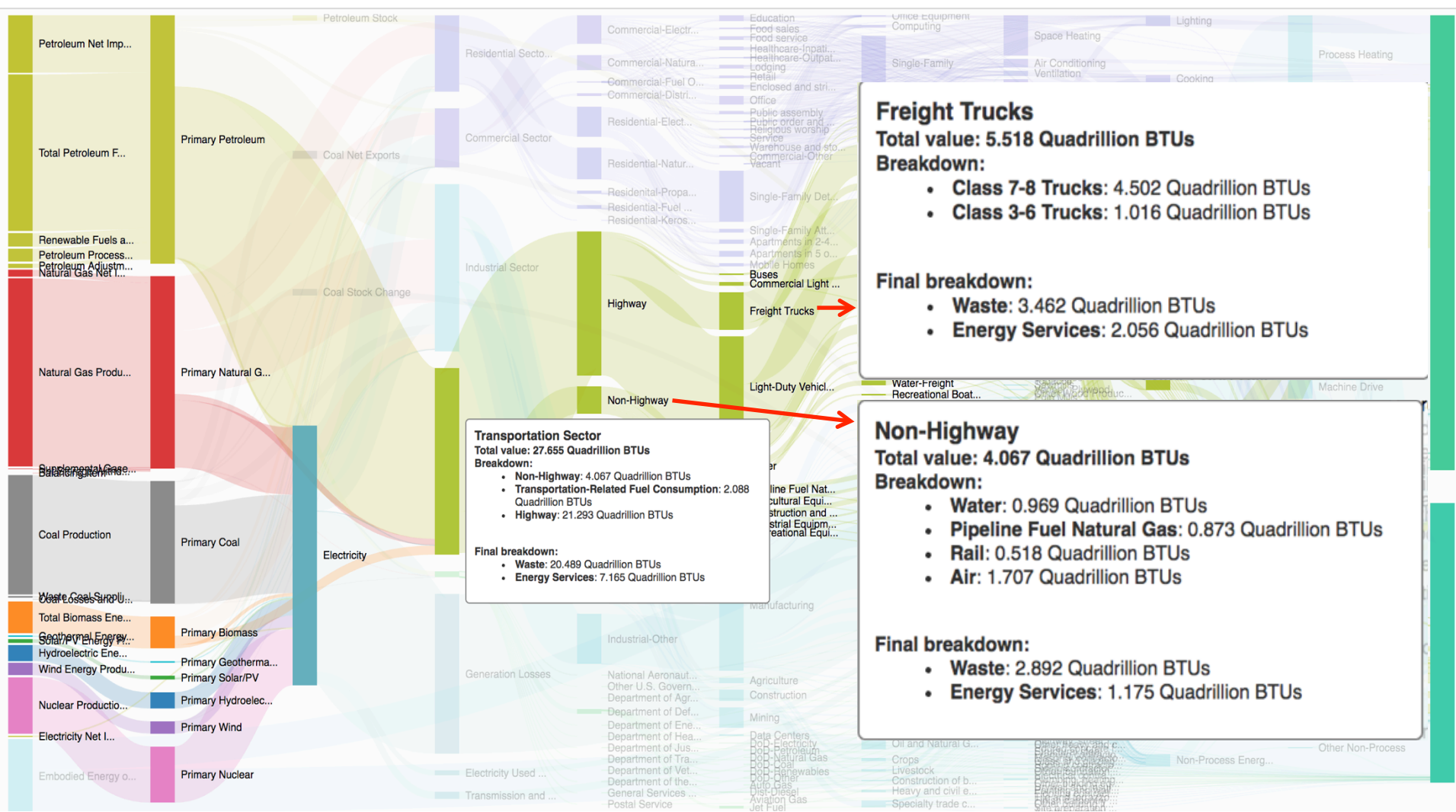
- Current:
 - Energy- and carbon-efficient conversions
 - Novel platform molecules
 - Waste stream utilization
“vast difference: a mixture of valuable chemicals and a valuable mixture of chemicals”
- Long-term:
 - All C-based synthetic materials (including biofuels)
 - Renewable C is distributed (biomass, biogas, CO₂)
 - Biological conversions benefit from economy of unit number: CapEx efficient at small scale
 - Heterologous feedstocks
 - Selective removal of functionality required
 - Average oxidation state ~0

Enormous possible impact – potential to mitigate >15% of US emissions



Xu et al. (2016) Energy Technol

Transportation (Non-light duty): ~9% of US Energy Use



Freight Trucks

Total value: 5.518 Quadrillion BTUs

Breakdown:

- **Class 7-8 Trucks:** 4.502 Quadrillion BTUs
- **Class 3-6 Trucks:** 1.016 Quadrillion BTUs

Final breakdown:

- **Waste:** 3.462 Quadrillion BTUs
- **Energy Services:** 2.056 Quadrillion BTUs

Transportation Sector

Total value: 27.655 Quadrillion BTUs

Breakdown:

- **Non-Highway:** 4.067 Quadrillion BTUs
- **Transportation-Related Fuel Consumption:** 2.088 Quadrillion BTUs
- **Highway:** 21.293 Quadrillion BTUs

Final breakdown:

- **Waste:** 20.489 Quadrillion BTUs
- **Energy Services:** 7.165 Quadrillion BTUs

Non-Highway

Total value: 4.067 Quadrillion BTUs

Breakdown:

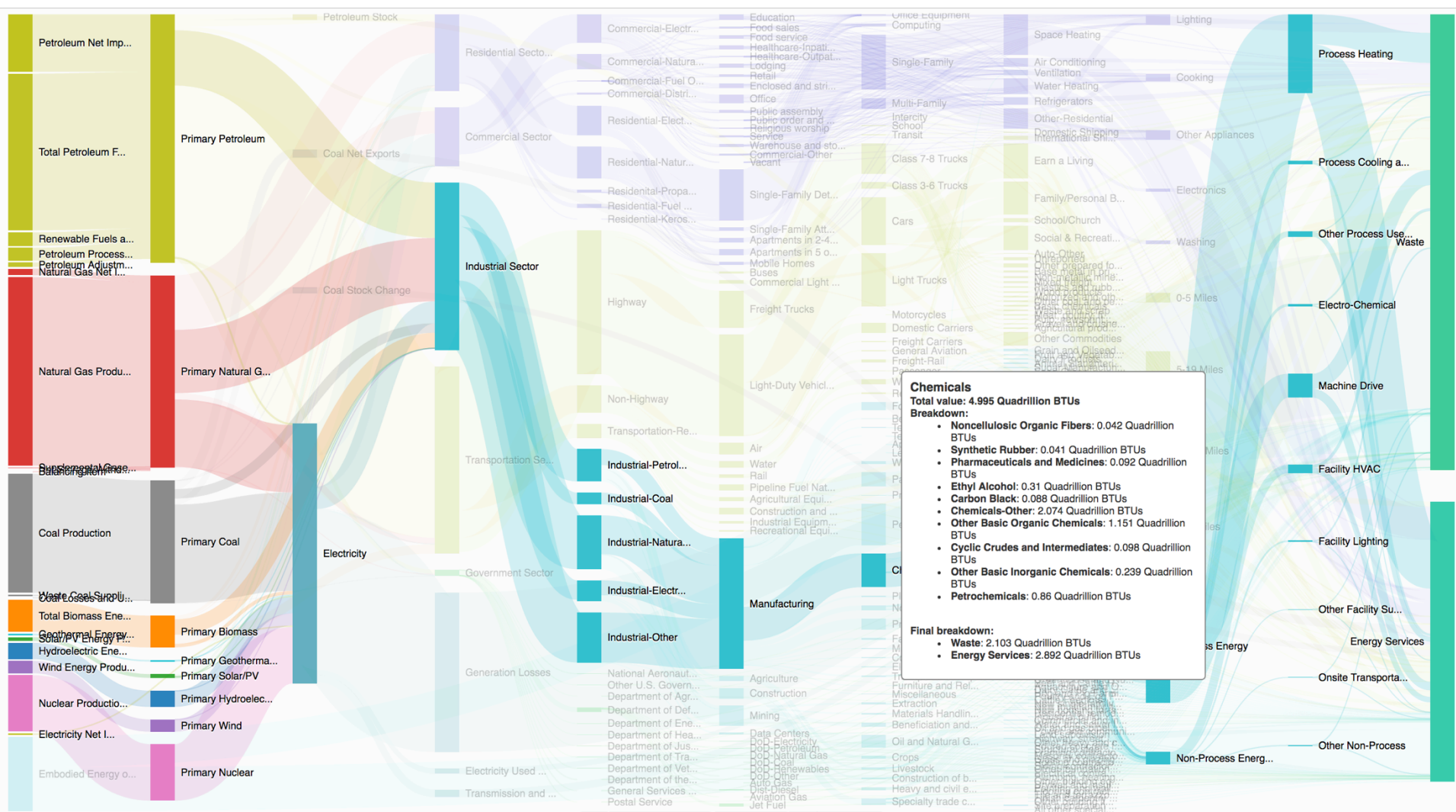
- **Water:** 0.969 Quadrillion BTUs
- **Pipeline Fuel Natural Gas:** 0.873 Quadrillion BTUs
- **Rail:** 0.518 Quadrillion BTUs
- **Air:** 1.707 Quadrillion BTUs

Final breakdown:

- **Waste:** 2.892 Quadrillion BTUs
- **Energy Services:** 1.175 Quadrillion BTUs

Chemical Production: 5% of US Energy Use

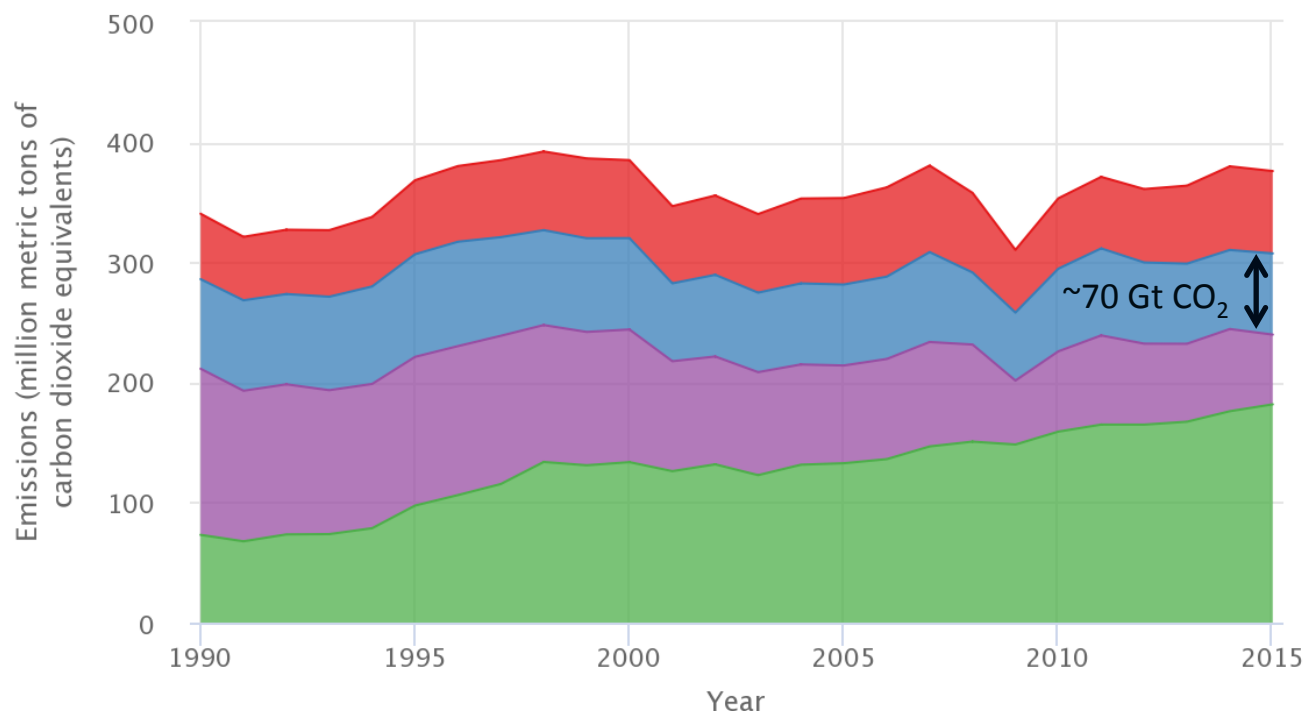
...plus 1% of additional emissions



Chemical Production: 5% of US Energy Use

...plus 1% of additional emissions

U.S. Greenhouse Gas Emissions from Industrial Processes, 1990-2015



● Mineral products ● Chemical production and use ● Metal production
● Production and use of fluorinated gases

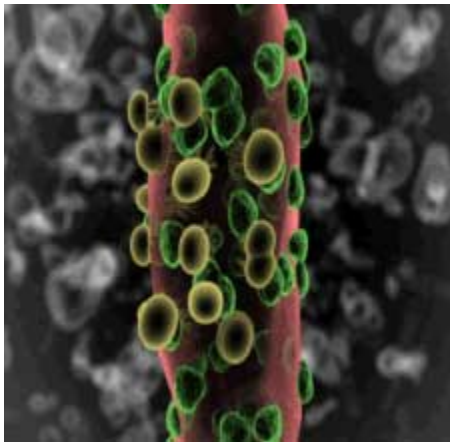
Source: US EPA's Inventory of US Greenhouse Gas Emissions and Sinks: 1999-2015
<https://www.epa.gov/ghgemissions/inventory-us-greenhouse-gas-emissions-and-sinks>

Cell-Free Bioconversion

Bioconversions – production of value-added compounds using biocatalysts



Microbial



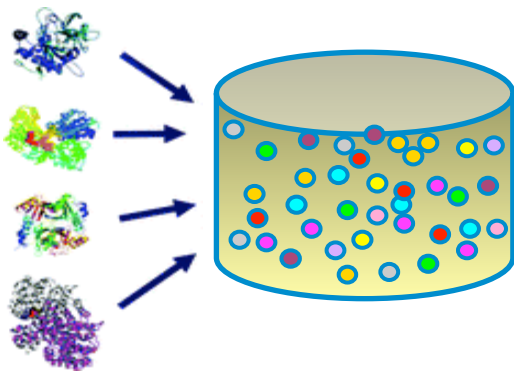
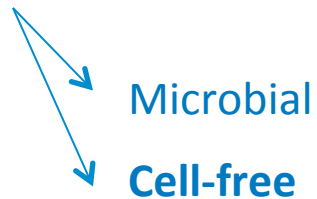
Guedes et al. (2007) *Anim. Feed. Sci. Technol.*

Microbial Fermentation

- Dominant biomanufacturing platform
- High engineering complexity
 - Self-replicating / self-repair
 - Natural system – evolves for maximum growth rate (opposed to engineering objective)
- Membrane-enclosed
 - Transport required
 - Solvent toxicity

Cell-Free Bioconversion

Bioconversions – production of value-added compounds using biocatalysts



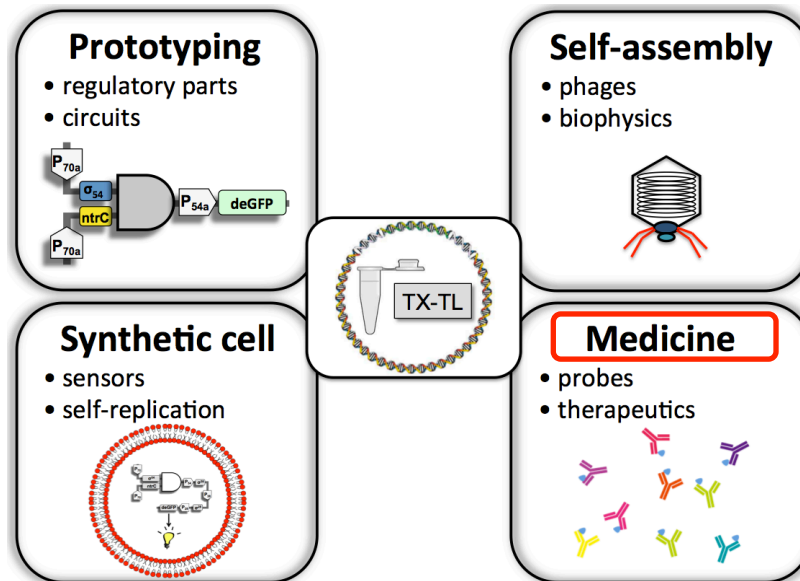
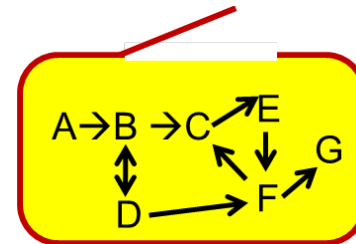
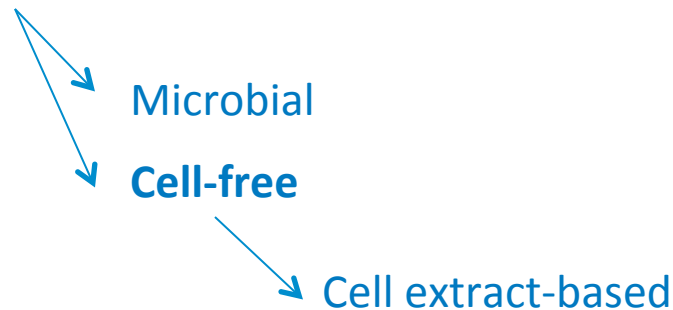
Zhang. (2011) ACS Catalysis 1: 998

Cell-Free Bioconversion

- Fast reaction rate
- Broad reaction conditions (solvent tolerance, reduced effect of toxins, temp., pH, etc.)
- Complete orthogonality
→ high yield
- Challenge: unstable, expensive co-factors
→ Pathway design
→ Cofactor engineering

Cell-Free Bioconversion

Bioconversions – production of value-added compounds using biocatalysts

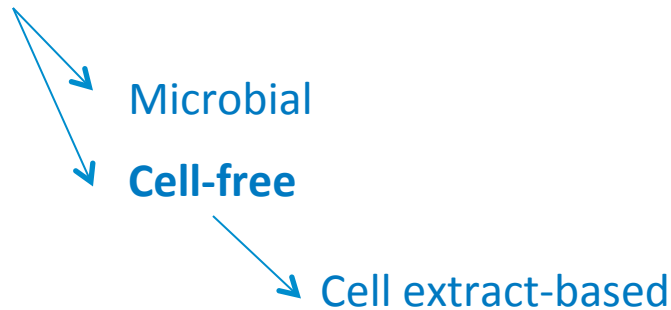


SUTRO
BIOPHARMA

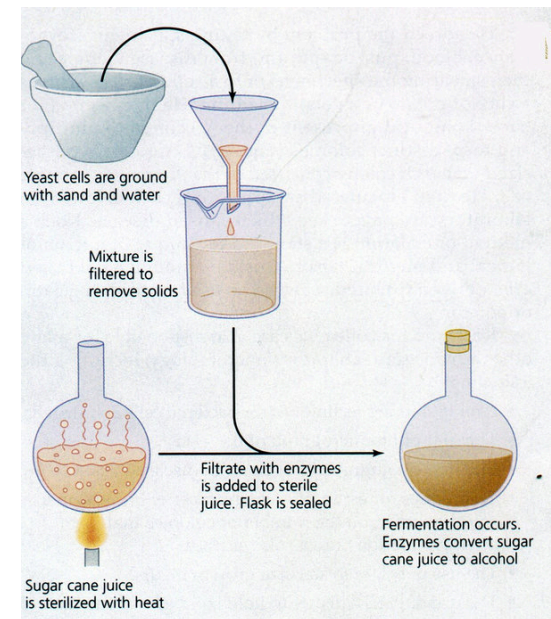
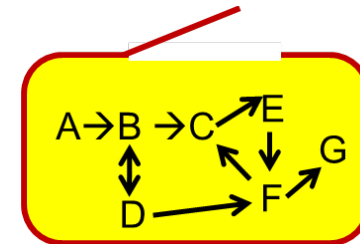
www.noireauxlab.org

Cell-Free Bioconversion

Bioconversions – production of value-added compounds using biocatalysts

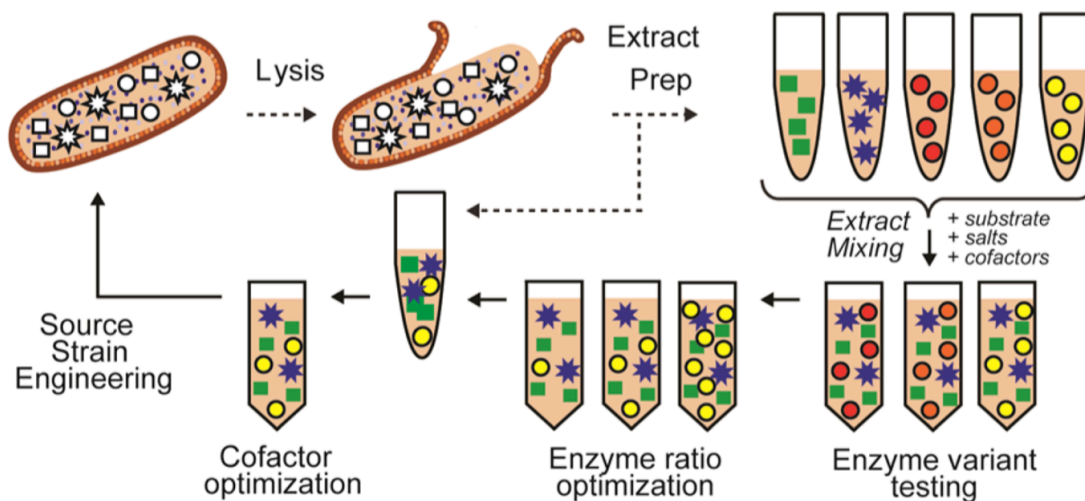
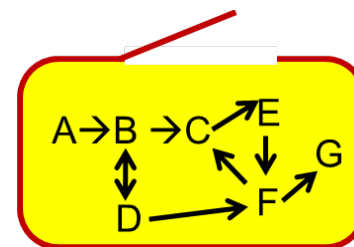
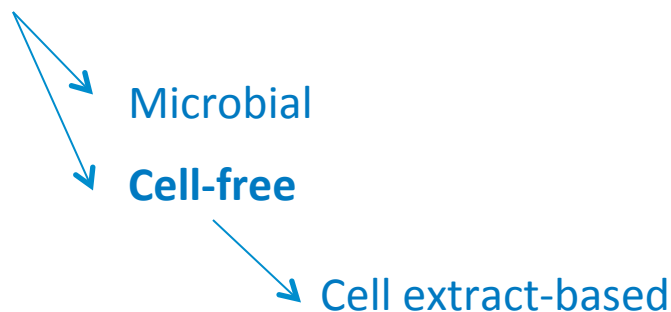


Cell-Free Ethanol Fermentation
Eduard Buchner
Nobel Prize in Chemistry, 1907



Cell-Free Bioconversion

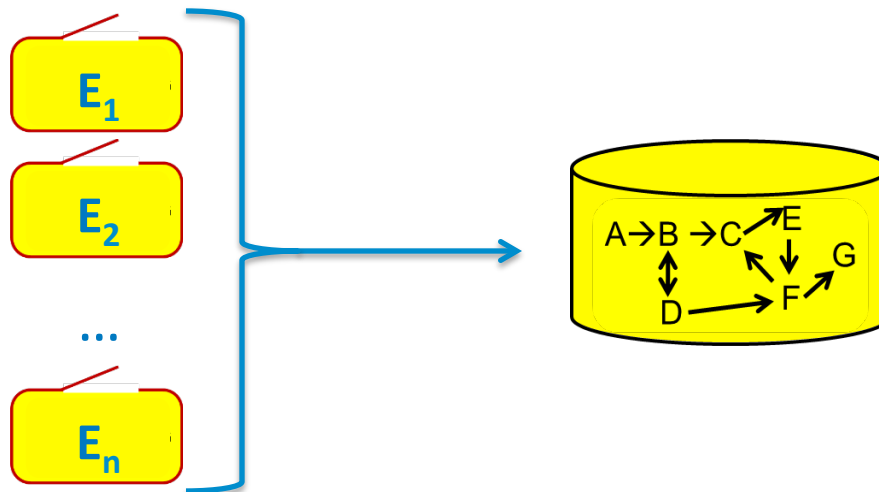
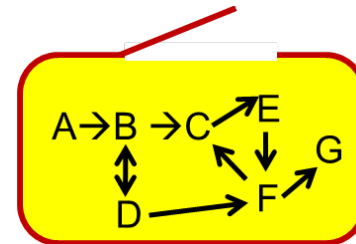
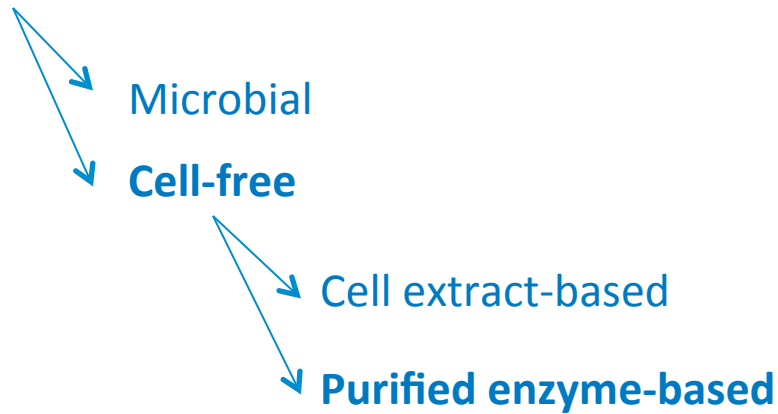
Bioconversions – production of value-added compounds using biocatalysts



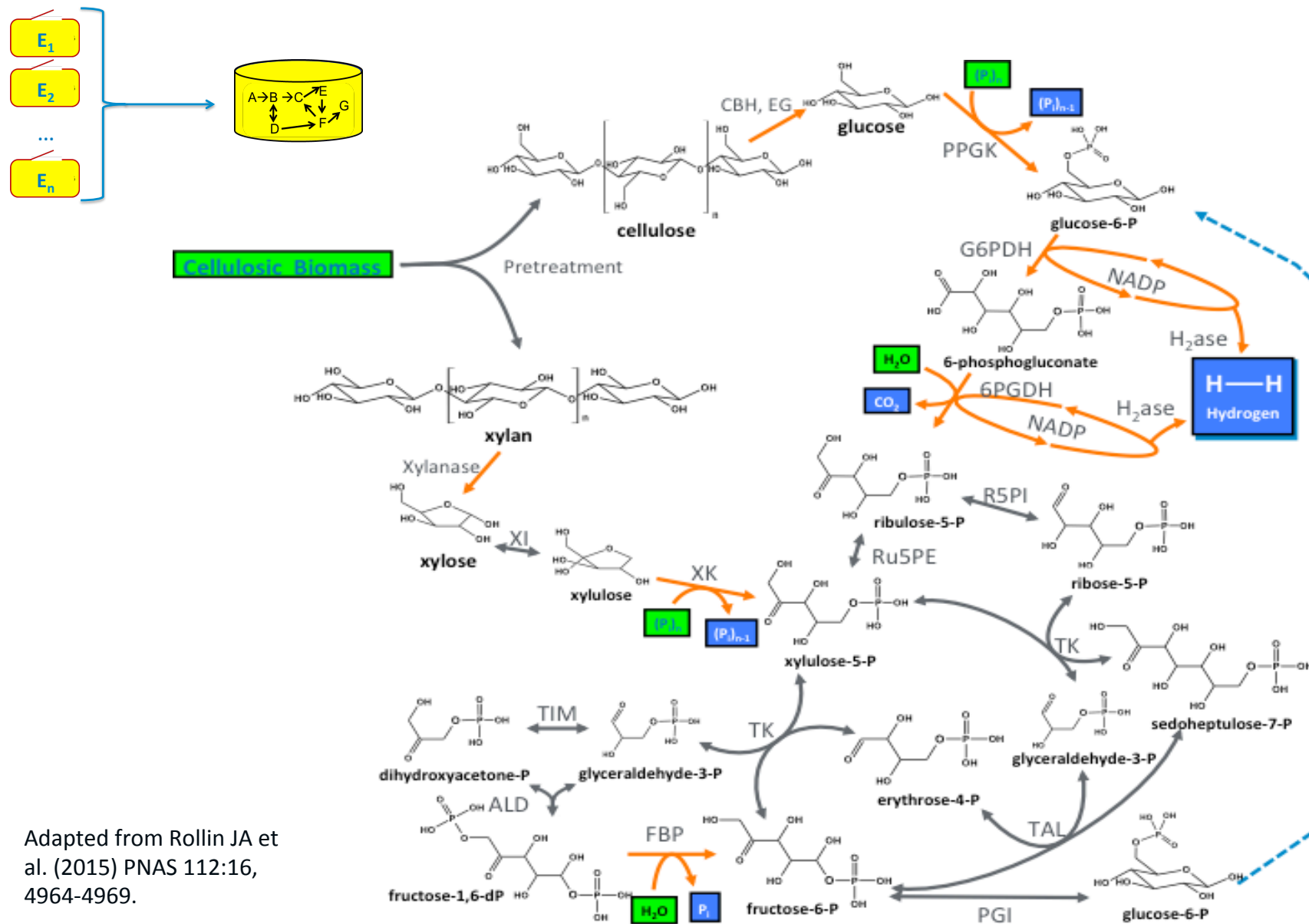
Dudley QM et al.
(2016) ACS Synth Biol
5, 1578-1588.

Cell-Free Bioconversion

Bioconversions – production of value-added compounds using biocatalysts

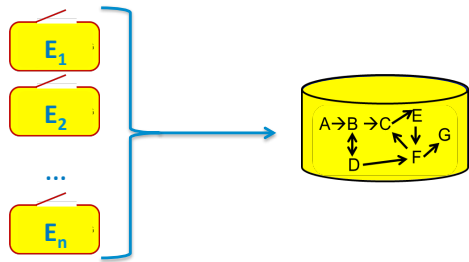


Purified Cell-Free Example: Biomass Sugars to H₂

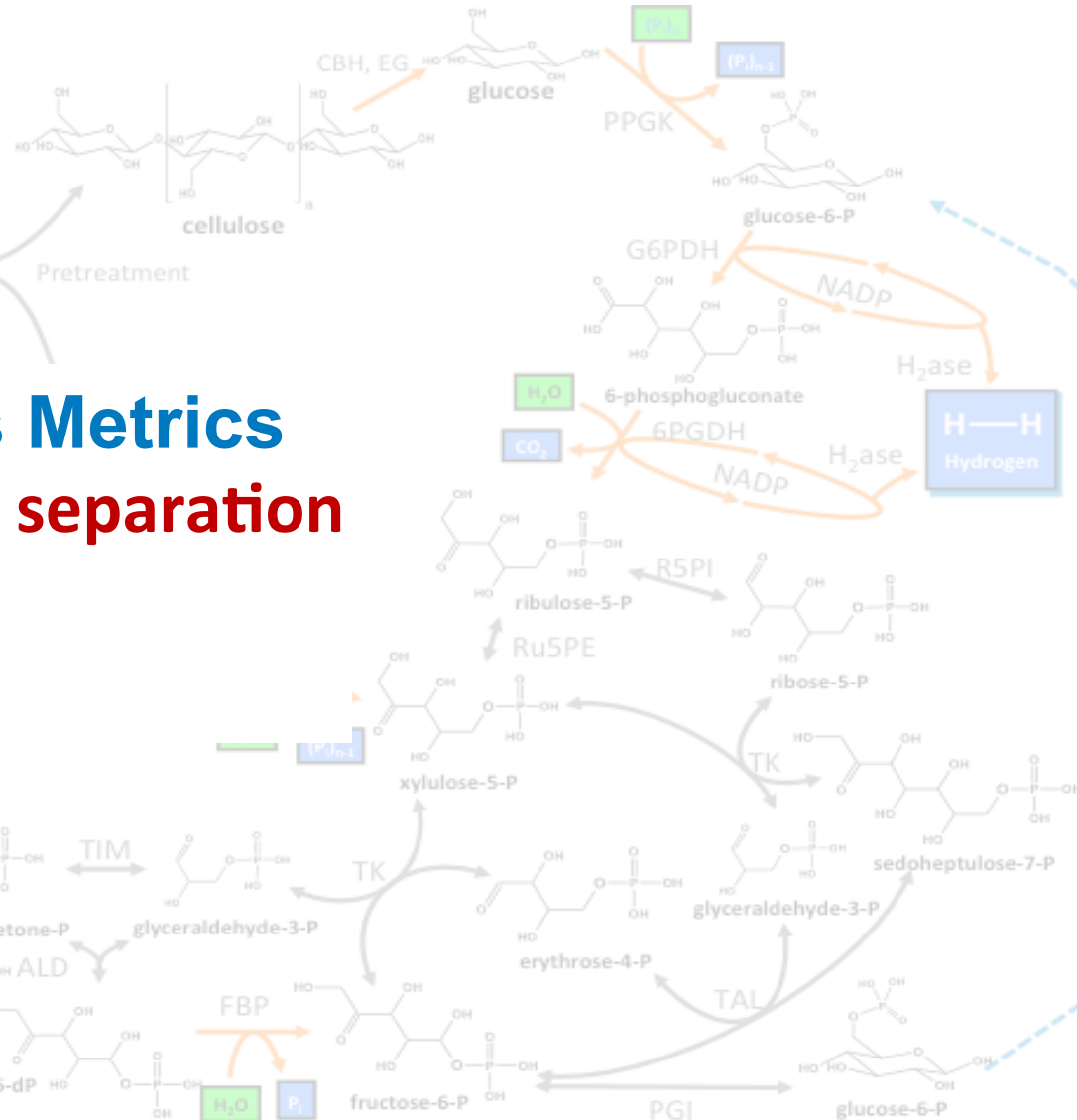


Adapted from Rollin JA et al. (2015) PNAS 112:16, 4964-4969.

Purified Cell-Free Example: Biomass Sugars to H₂



Cellulosic Biomass

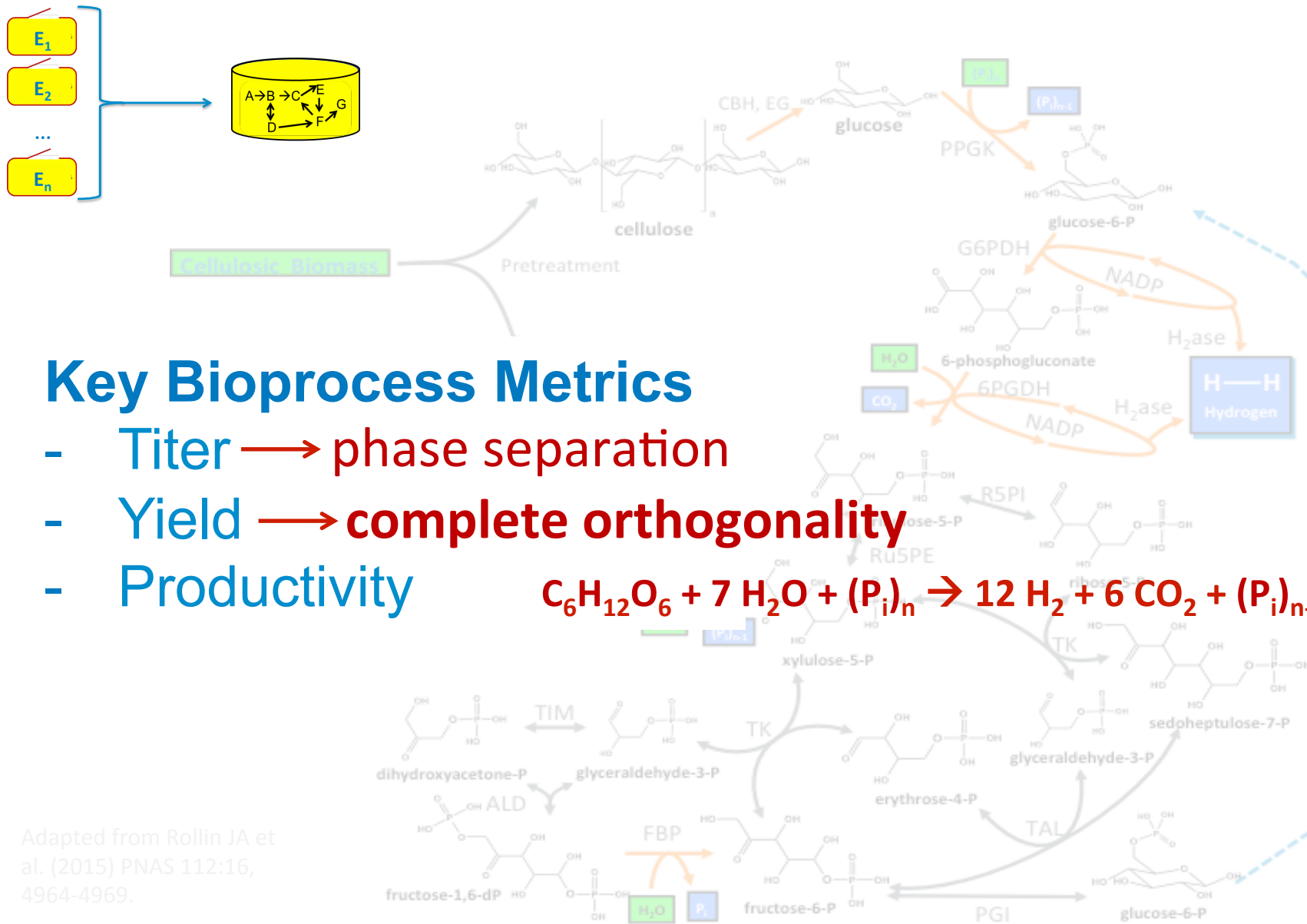


Key Bioprocess Metrics

- Titer → phase separation
- Yield
- Productivity

Adapted from Rollin JA et al. (2015) PNAS 112:16, 4964-4969.

Purified Cell-Free Example: Biomass Sugars to H₂

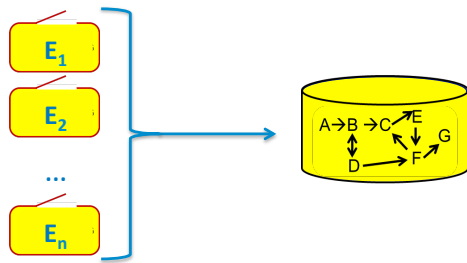


Key Bioprocess Metrics

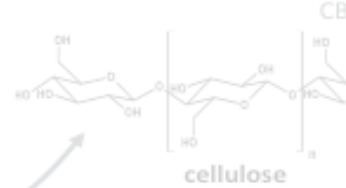
- Titer → phase separation
- Yield → **complete orthogonality**
- Productivity

Adapted from Rollin JA et al. (2015) PNAS 112:16, 4964-4969.

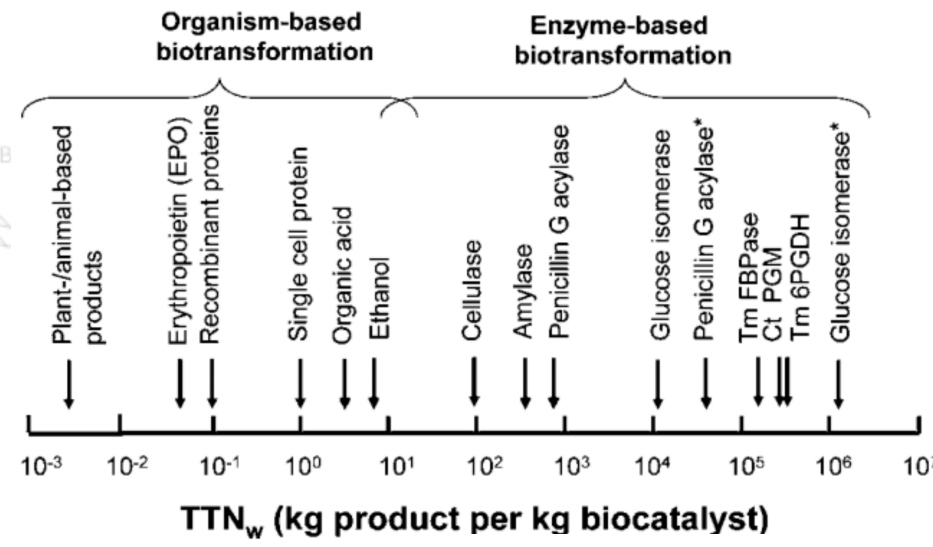
Purified Cell-Free Example: Biomass Sugars to H₂



Cellulosic Biomass

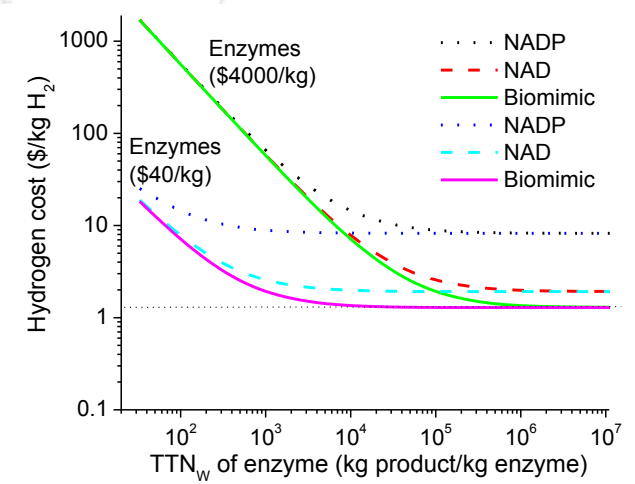
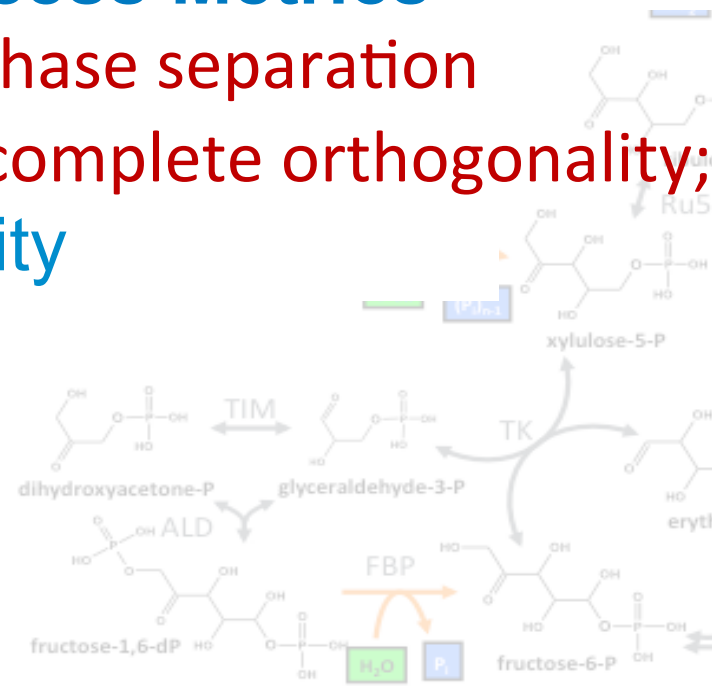


Pretreatment



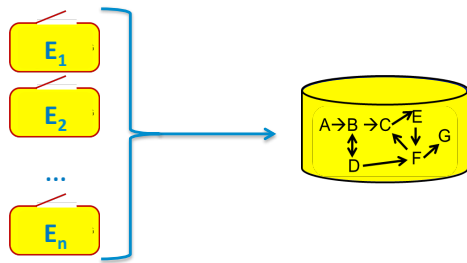
Key Bioprocess Metrics

- Titer → phase separation
- Yield → complete orthogonality; **TTN_w**
- Productivity

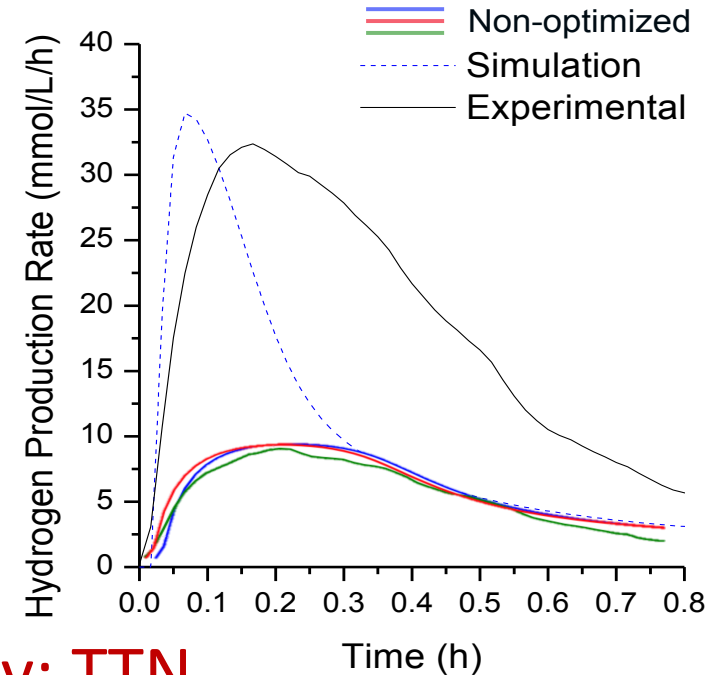


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Purified Cell-Free Example: Biomass Sugars to H₂

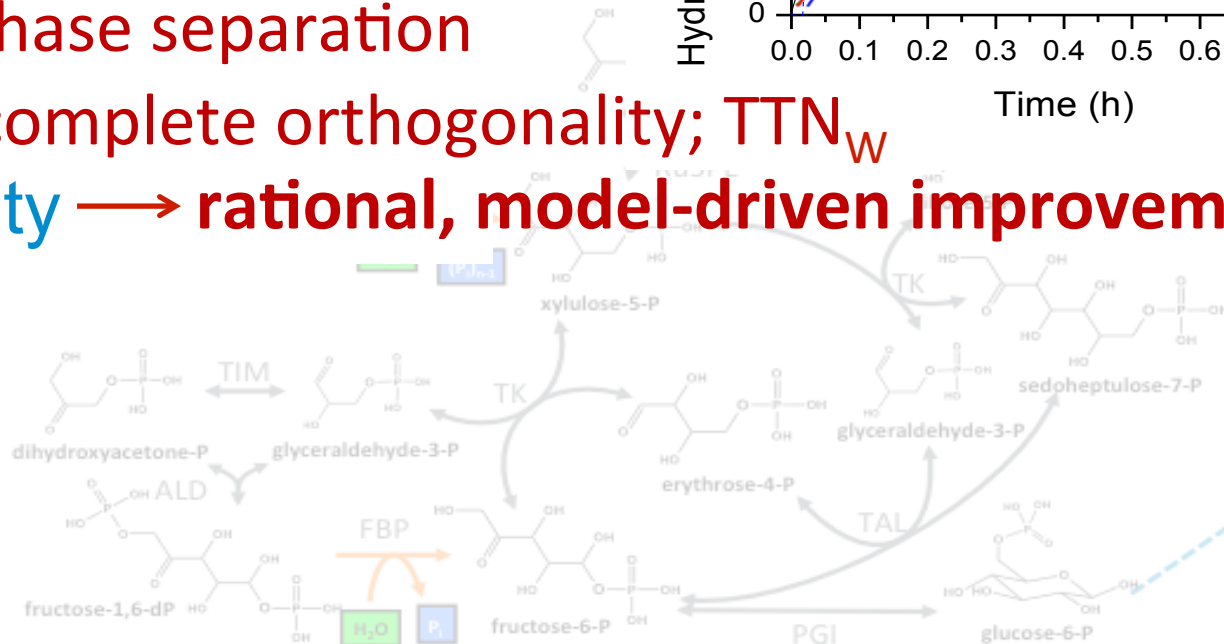


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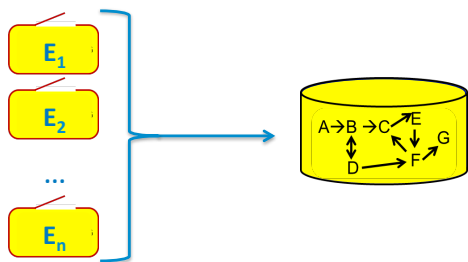
Key Bioprocess Metrics

- Titer → phase separation
- Yield → complete orthogonality; TTN_W
- Productivity → rational, model-driven improvement

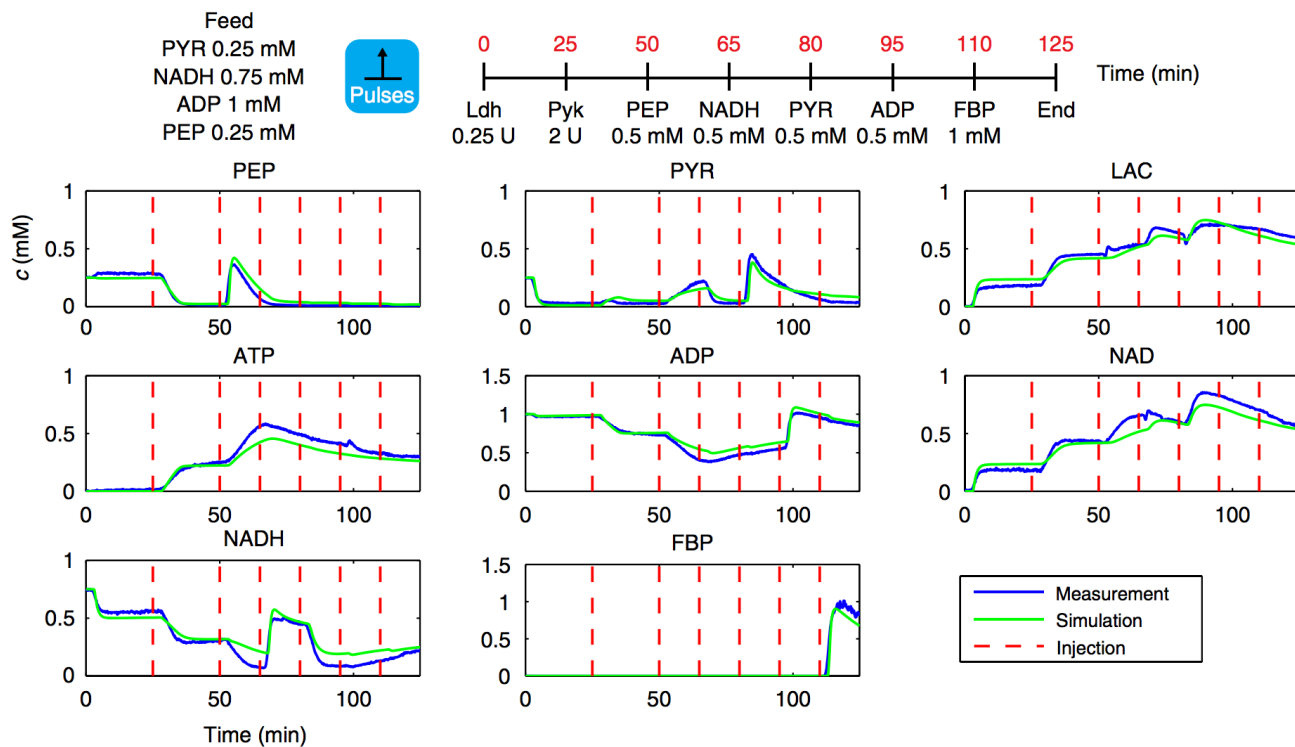


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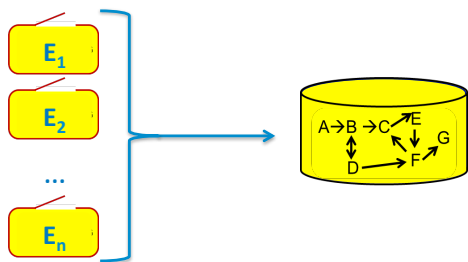
Purified Cell-Free Conversion – Challenges and Opportunities



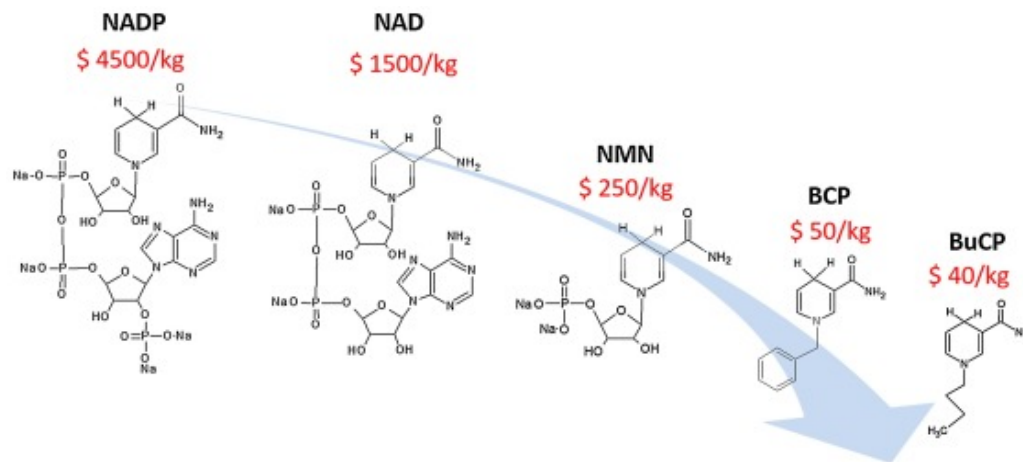
- Advanced modeling
 - Ensemble approaches
 - Perturbations for robust parameterization
 - Alternative objectives (min [NADH], reduce inhibitory intermediates, modulation of ATP, PP_i recycle rates etc.)



Purified Cell-Free Conversion – Challenges and Opportunities

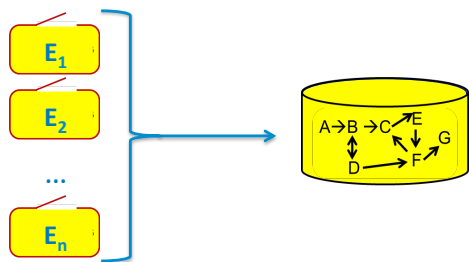


- Advanced modeling
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- Cofactor engineering
 - Replacement with biomimetics
 - Opportunities to harness external reducing power

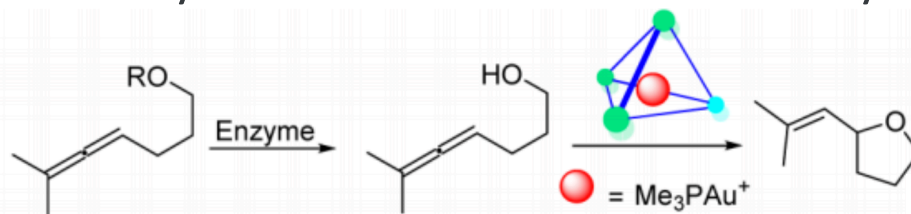


You and Zhang (2017) Proc Biochem

Purified Cell-Free Conversion – Challenges and Opportunities



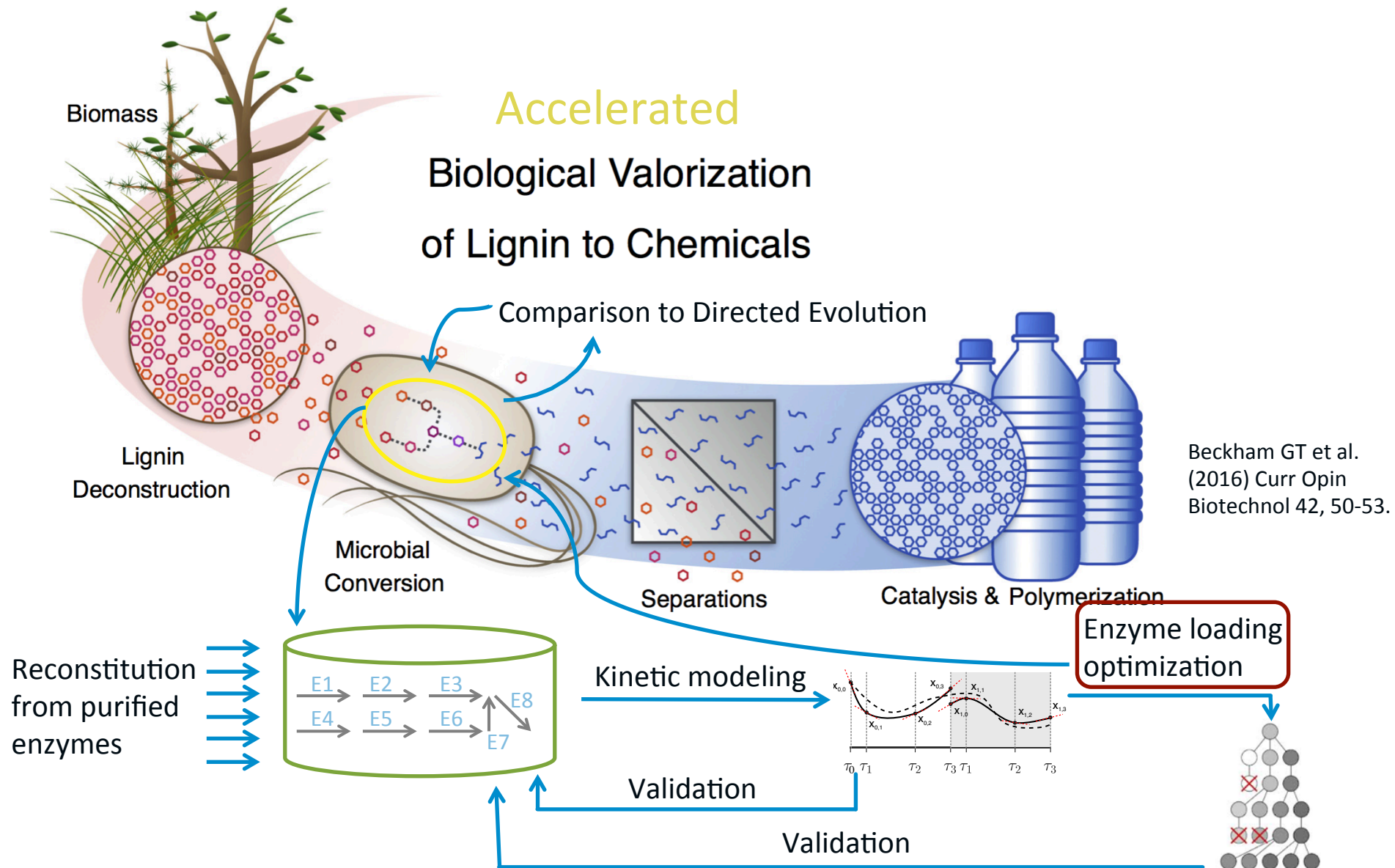
- Advanced modeling
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 - Alternative objectives (min [NADH], reduce inhibitory intermediates, modulation of ATP, PP_i recycle rates etc.)
- Cofactor engineering
 - Replacement with biomimetics
 - Opportunities to harness external reducing power
- Tunable pathway properties
 - Artificial actuators
 - Engineered allostery
- Compartmentalization / spatial control enabling for:
 - Substrate channeling
 - Hybridization with chemical catalysis



Denard et al. (2013) ACS Cat

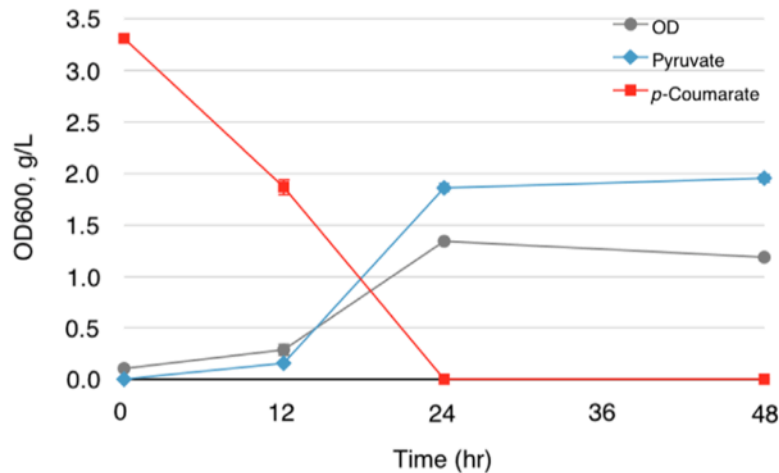
Cell-Free Metabolic Engineering

Accelerated Biological Valorization of Lignin to Chemicals

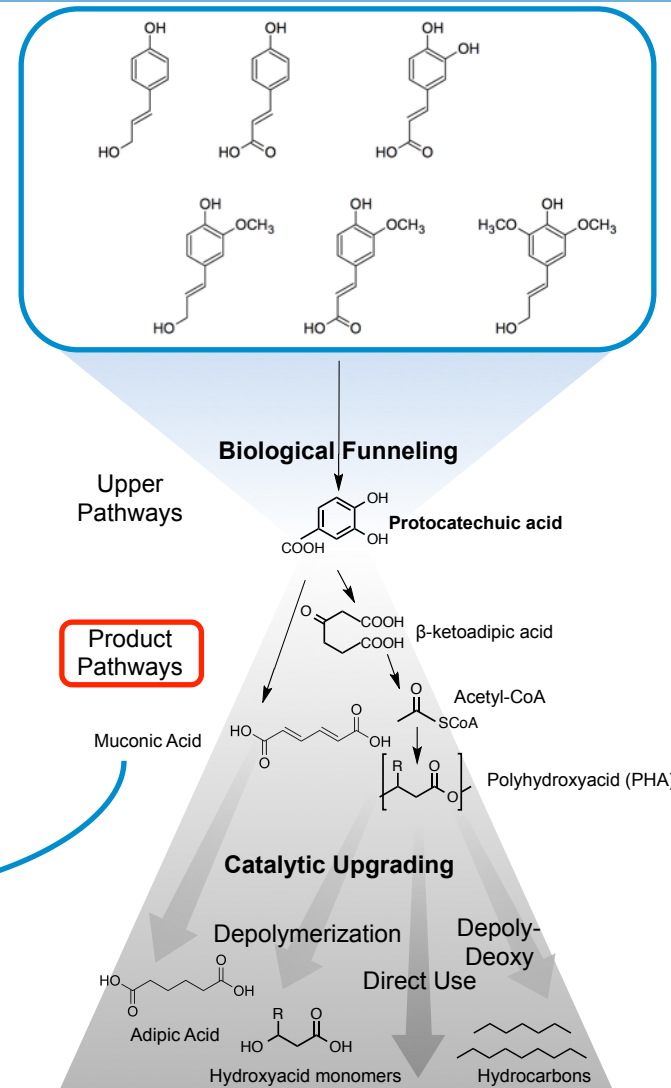


Beckham GT et al. (2016) Curr Opin Biotechnol 42, 50-53.

Pseudomonas putida KT2440 – Lignin Biological Funneling

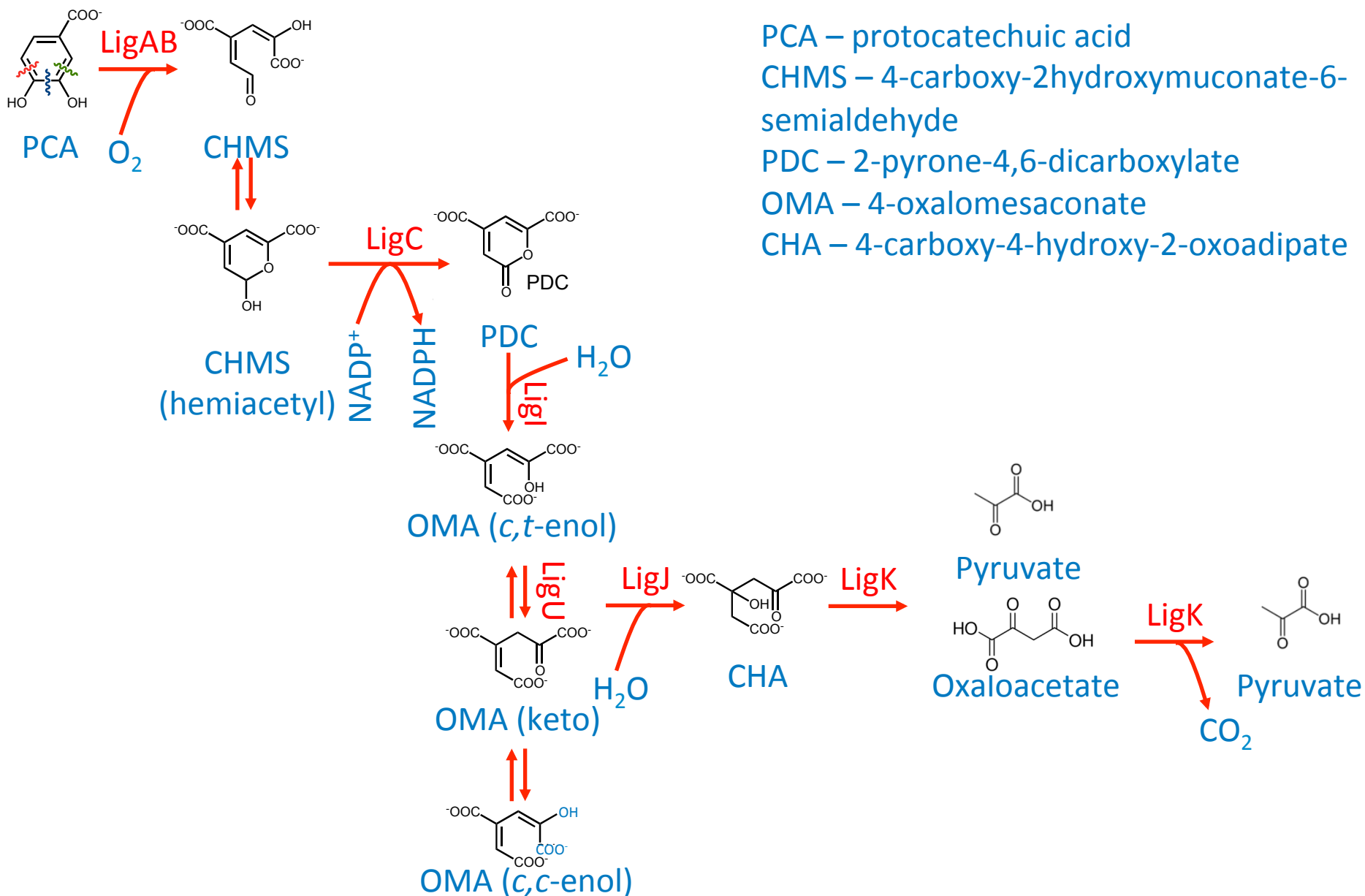


Johnson CW and Beckham GT. (2015) Metab Eng 28, 240-247.

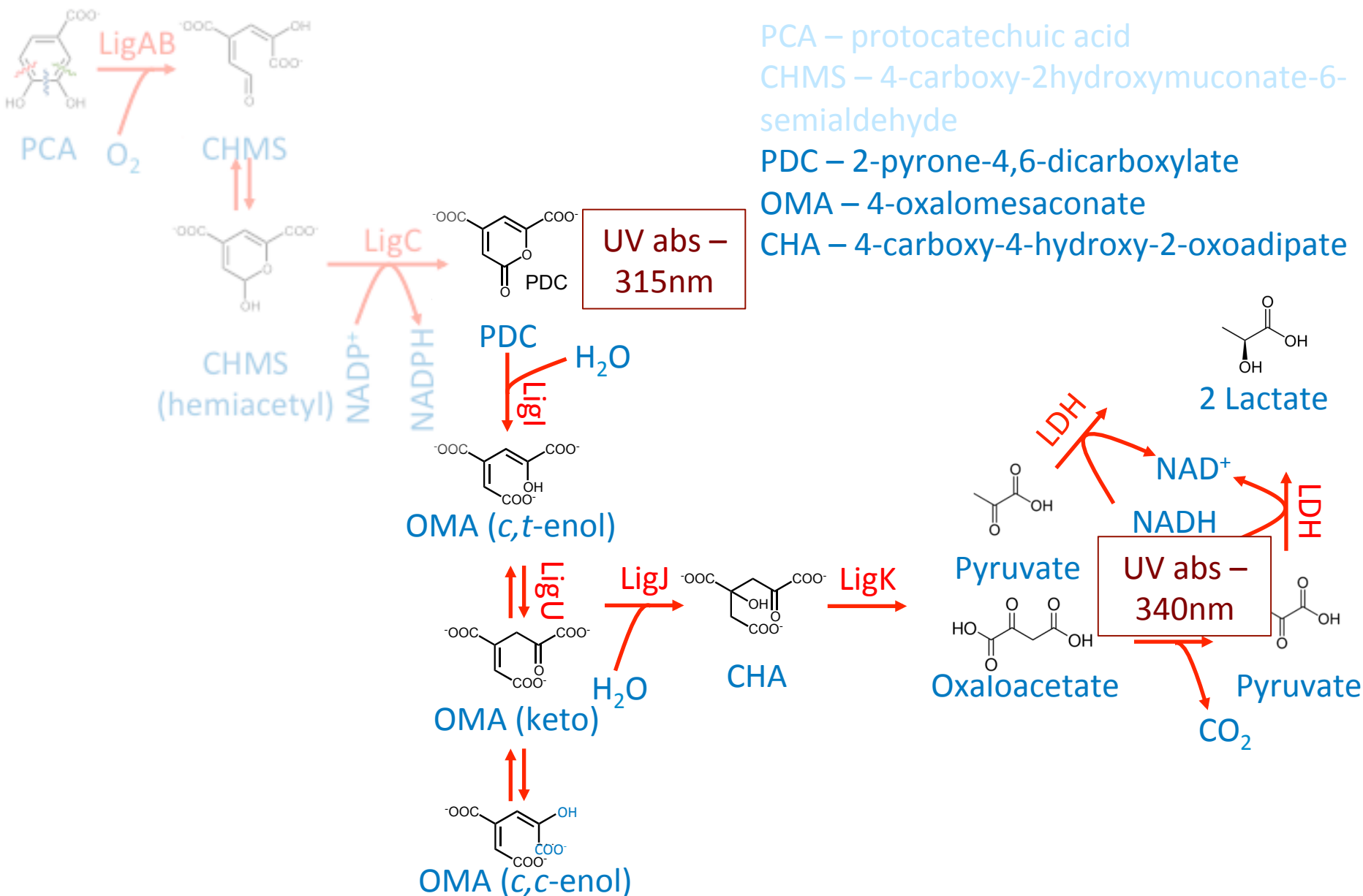


Linger JG et al. (2014) PNAS 111:33, 12013-12018.

Pathway Selection: PCA 4,5-meta Cleavage

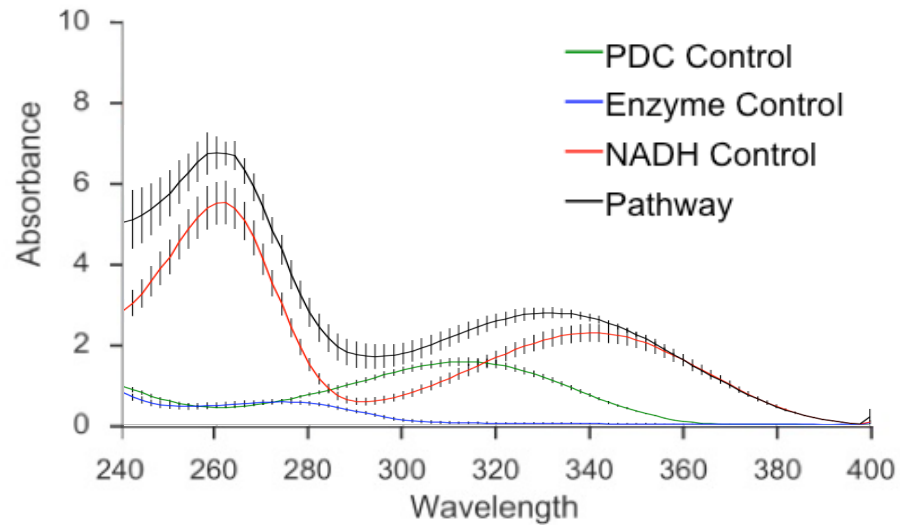


Pathway Selection: PCA 4,5-meta Cleavage

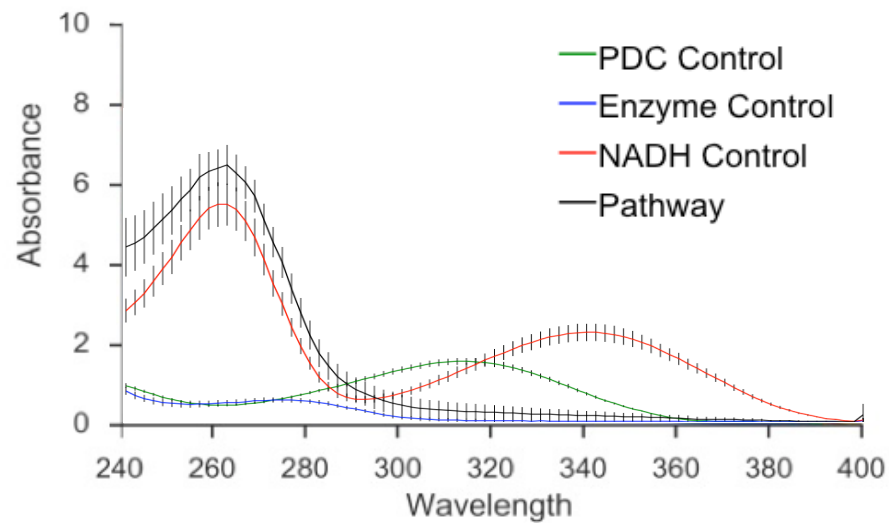


Confirmation of Pathway Activity

0 min

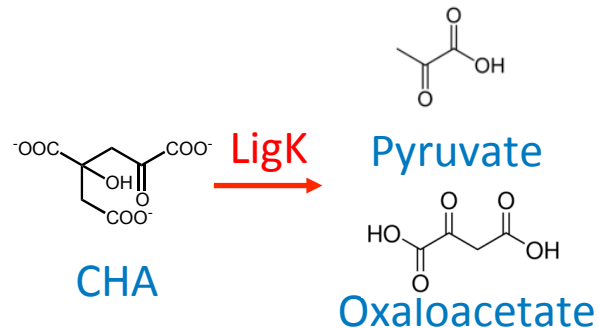


10 min



Pathway Kinetic Modeling

- Michaelis-menten kinetics rate equations built for each reaction



$$r_{LigK1} = \frac{d[OAA]}{dt} = \frac{d[Pyr1]}{dt} = \frac{k_{cat}^{LigK1} [LigK][CHA]}{K_{M,CHA}^{LigK} + [CHA]}$$

Pathway Kinetic Modeling

- Michaelis-menten kinetics rate equations built for each reaction
- Parameters determined for enzymes collectively
 - Orthogonal collocation for rapid curve-fitting
 - Bootstrap method to determine probability density for each variable
- Additional possible parameters: inhibition constants, inactivation terms

$$r_{LigI} = \frac{d[OMA_{enol}]}{dt} = \frac{k_{cat}^{LigI} [LigI][PDC]}{K_M^{LigI} + [PDC]}$$

$$r_{LigU} = \frac{d[OMA_{keto}]}{dt} = \frac{k_{cat}^{LigU} [LigU][OMA_{enol}]}{K_M^{LigU} + [OMA_{enol}]}$$

$$r_{LigJ} = \frac{d[CHA]}{dt} = \frac{k_{cat}^{LigJ} [LigJ][OMA_{keto}]}{K_M^{LigJ} + [OMA_{keto}]}$$

$$r_{LigK1} = \frac{d[OAA]}{dt} = \frac{d[Pyr1]}{dt} = \frac{k_{cat}^{LigK1} [LigK][CHA]}{K_{M,CHA}^{LigK} + [CHA]}$$

$$r_{LigK2} = \frac{d[Pyr2]}{dt} = \frac{k_{cat}^{LigK2} [LigK][OAA]}{K_{M,OAA}^{LigK} + [OAA]}$$

$$P = \begin{bmatrix} K_M^{LigI} \\ k_{cat}^{LigU} \\ K_M^{LigU} \\ k_{cat}^{LigJ} \\ K_M^{LigJ} \\ k_{cat}^{LigK1} \\ K_{M,CHA}^{LigK} \\ k_{cat}^{LigK2} \\ K_{M,OAA}^{LigK} \end{bmatrix}$$

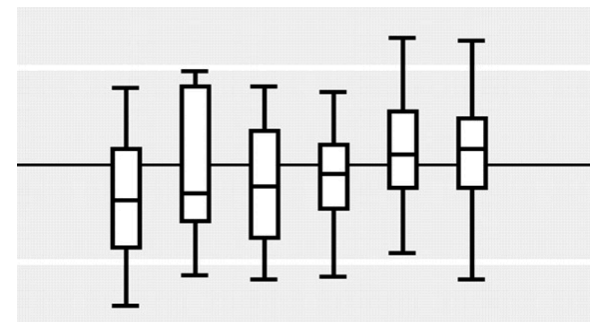
Pathway Kinetic Modeling - Experimental Validation and Iteration

- Input at least 10 different conditions to inform 9-parameter model

[E] (μM)

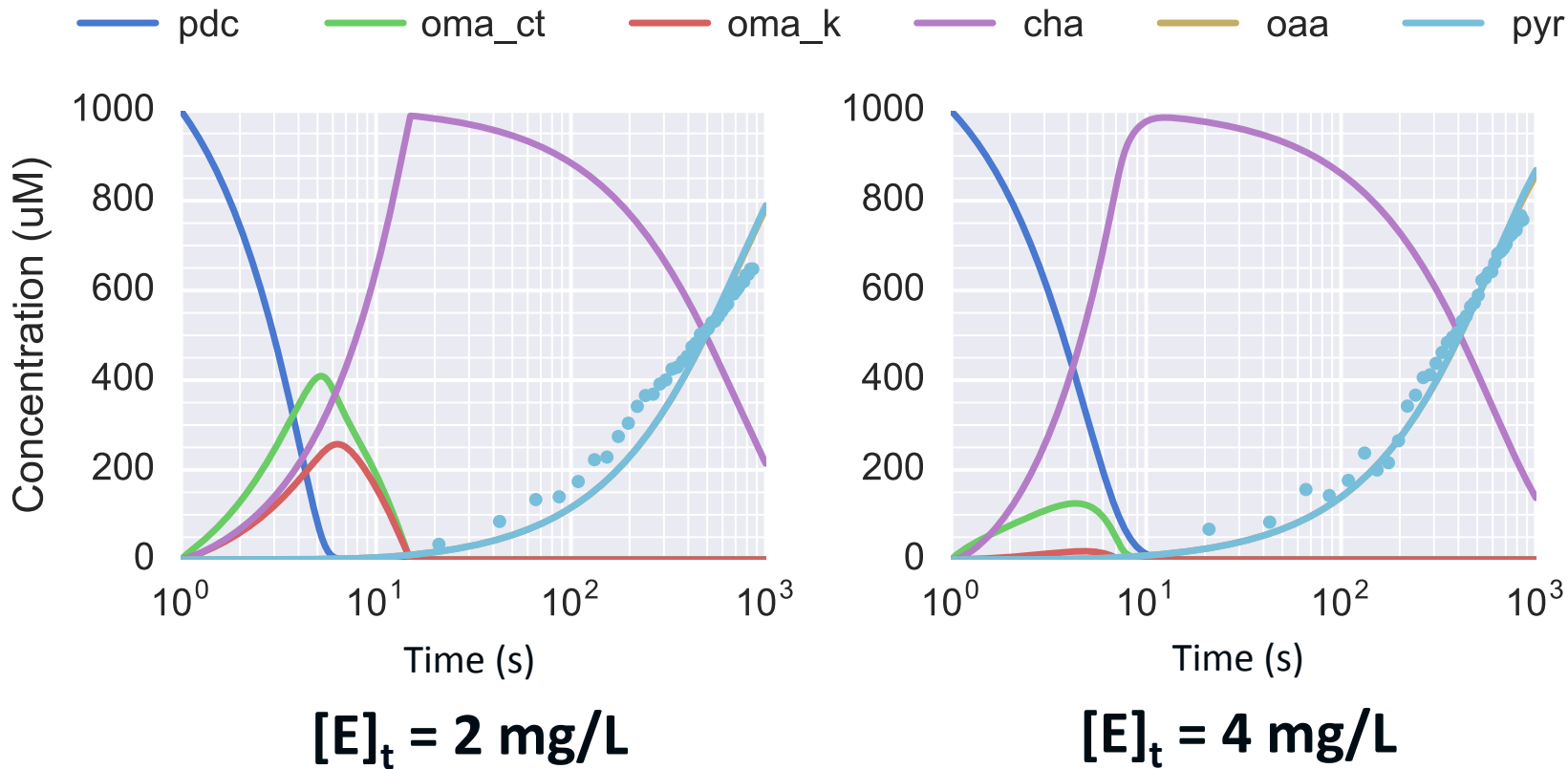
$$\begin{bmatrix} \text{E1} \\ \text{E2} \\ \text{E3} \\ \text{E4} \end{bmatrix} = \begin{bmatrix} 0.1 \\ 0.1 \\ 0.1 \\ 0.1 \end{bmatrix}, \begin{bmatrix} 0.5 \\ 0.5 \\ 0.5 \\ 0.5 \end{bmatrix}, \begin{bmatrix} 1 \\ 1 \\ 1 \\ 1 \end{bmatrix}, \begin{bmatrix} 0.1 \\ 1 \\ 1 \\ 1 \end{bmatrix}, \begin{bmatrix} 1 \\ 0.1 \\ 1 \\ 1 \end{bmatrix}, \begin{bmatrix} 1 \\ 1 \\ 0.1 \\ 1 \end{bmatrix}, \begin{bmatrix} 1 \\ 1 \\ 1 \\ 0.1 \end{bmatrix}, \dots$$

- Output parameter range estimation



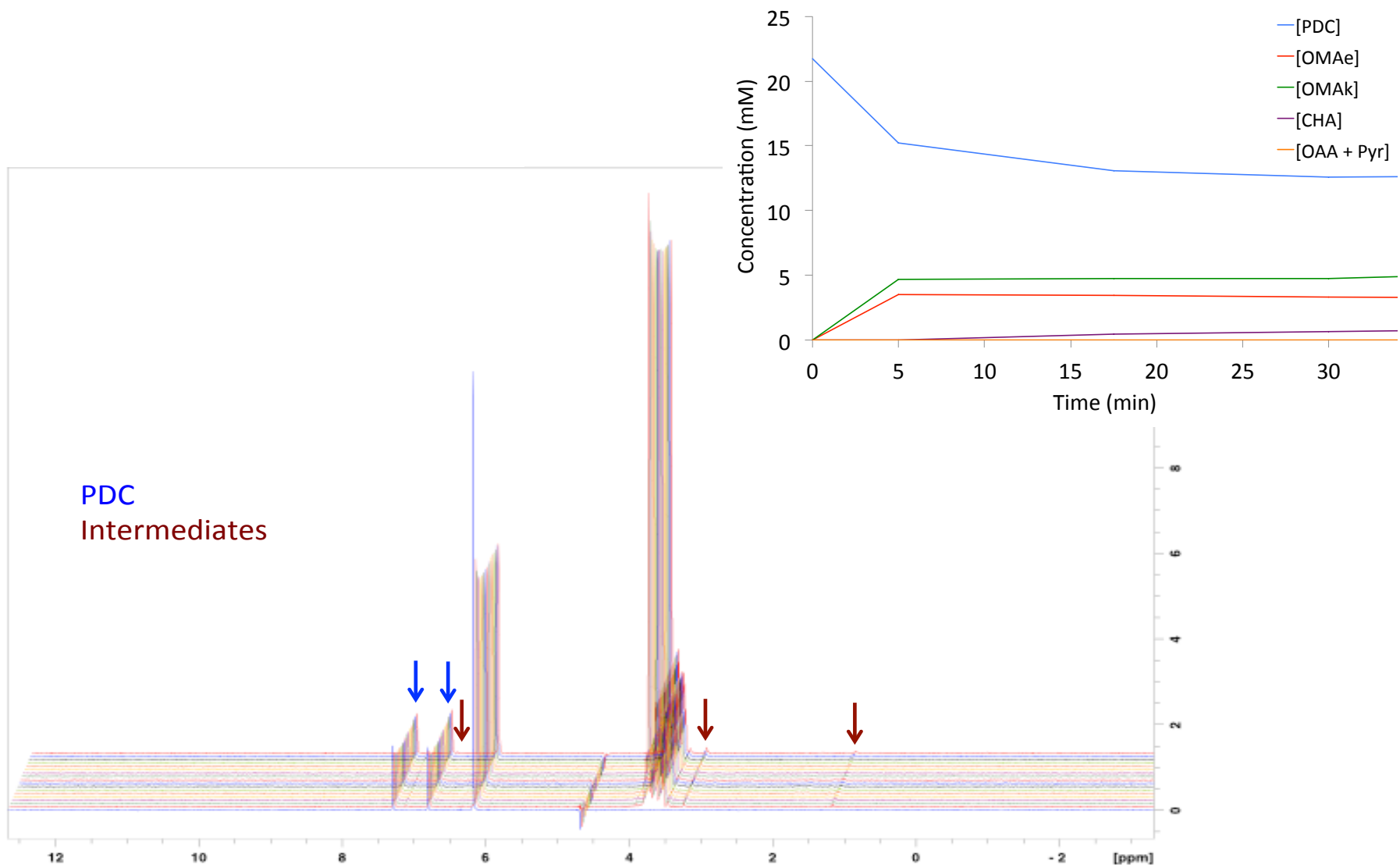
- Iterative improvement by varying enzyme ratio, substrate loading

Preliminary Model Results (Peter St. John)



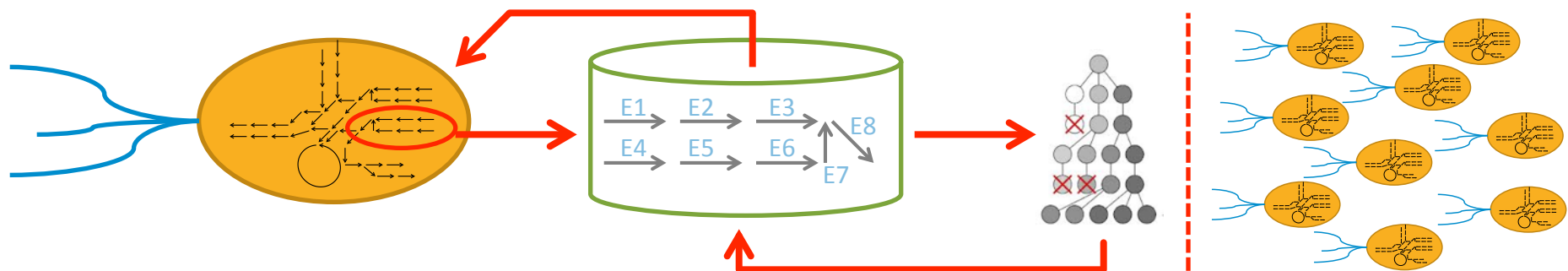
- Good initial fit of [Pyr] based on literature kinetic parameters
- **Intermediates detection required for robust model**

NMR Metabolomics – Initial Data



Cell-Free Metabolic Engineering Project Summary

1. Cell-free replication of lignin biological funneling pathways of interest
2. Characterization of the enzymes in this system
 - Kinetic parameters for each reaction: K_M , k_{cat} , K_I
 - Simultaneous system parameter estimation
3. Predict improvements that may be used in the production strain
 - Expression levels
 - Enzyme engineering targets / variant candidates
4. Use RBS calculator (or alternative methods) to translate optimal enzyme loadings into genetic elements
5. Directed evolution of this pathway to compare this approach to genetic, whole-cell optimization methods



Thanks to my mentors and collaborators:

Gregg Beckham and Chris Johnson

Peter St. John, Renee Happs, and many other group members

This work is supported by the NREL Director's Fellowship



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