

# BERAC Update

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**February 27, 2015**

# Systems Analysis of Biomass & Biofuel Production

## Cropping Systems



## Pretreated Biomass



## Hydrolysate



## Biofuels



Measurables

Site/Soil Type  
Crop/Seed/Row  
Plant/Harvest Date  
Fertilizer/Herbicide  
Season/Weather

Cellulose  
Hemicelluloses  
Lignin  
Plant Cell Residue

Total CHO  
C-5 & C-6 Sugars  
Amino Acids  
Organic Acids/Amides  
Ammonia/Phosphate  
~30 Metals/Inorganic Ions

Hydrolysate Inputs  
Transcripts  
Targeted Metabolites  
Excreted Products  
Fuel  
Input/Output COD  
Microbial Growth

## Data Management & Analysis

## Scientific Discoveries

# Collaboration Enables Discoveries

## GLBRC



Sustainability



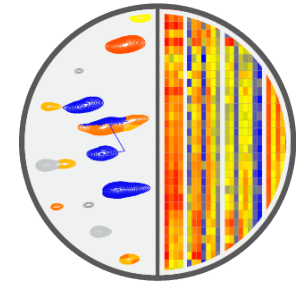
Plants



Deconstruction



Conversion



Enabling Technologies

## DOE BER

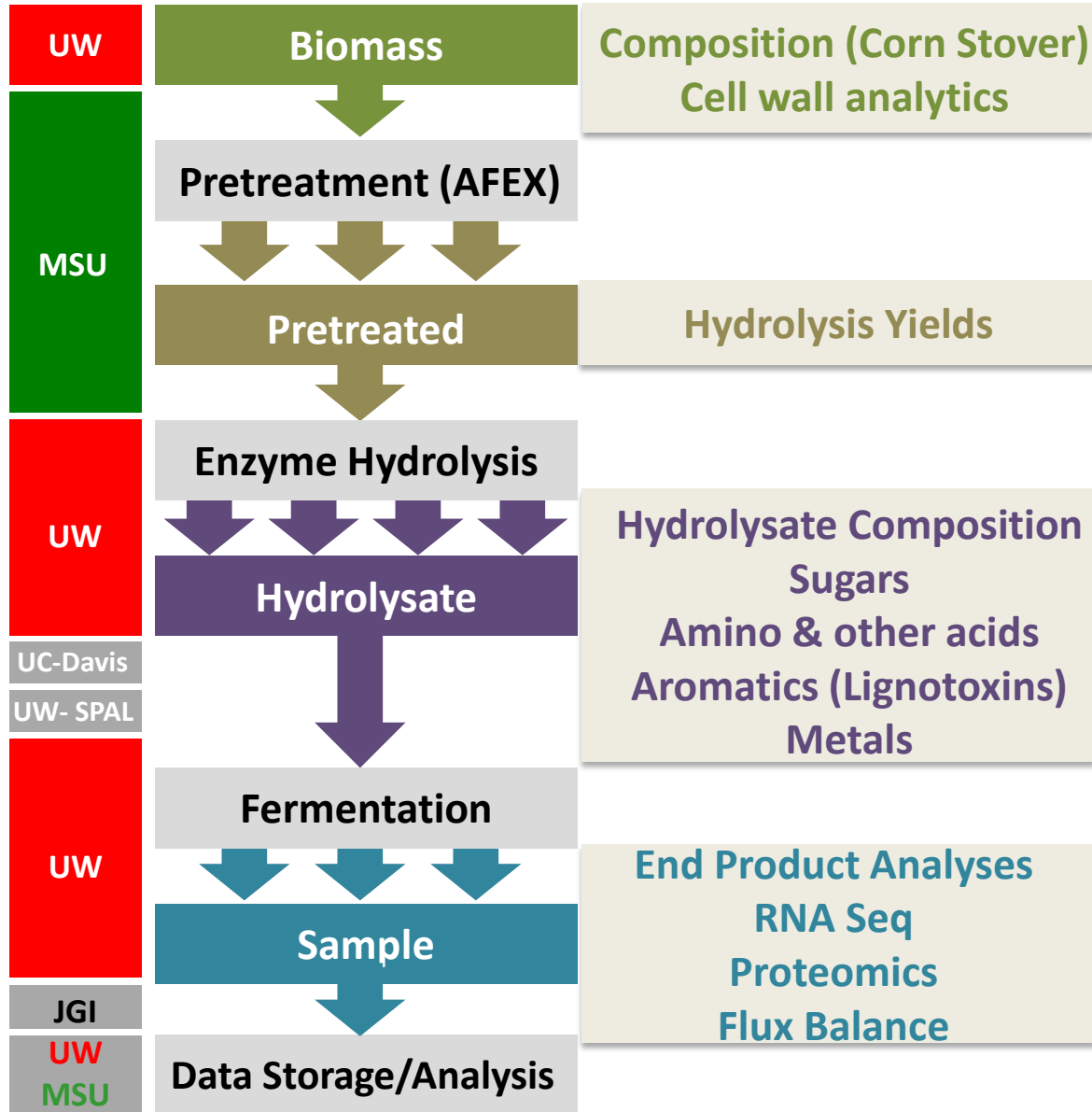


## “External”



Farmers  
Landowners  
Universities

# Biomass Field to Fuel Analysis





# Chemical composition of hydrolysates

Carbohydrates  
(Glc, Xyl, Ara,  
Gal, Man, Rha,  
Fru, Fuc)

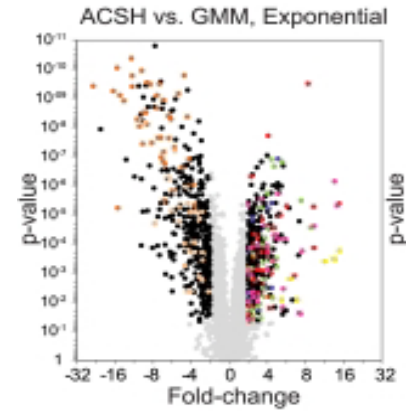
AAs  
(-C,M,W)

Nucleosides,  
Organic Acids,  
Aldehydes,  
Ketones,  
Amides  
Aromatics

Metals,  
Minerals  
& Anions



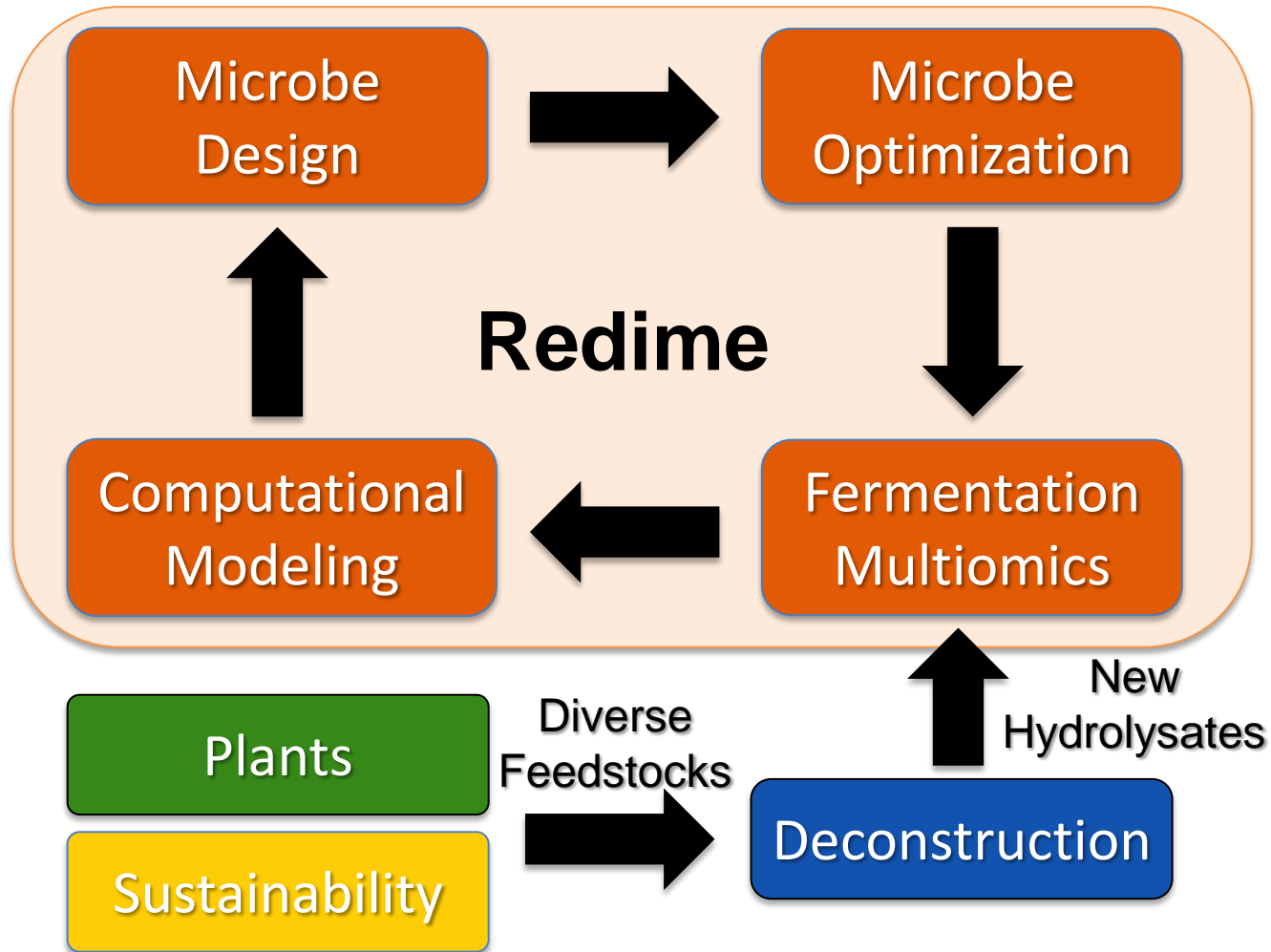
Media component	ACSH	SynH <sup>1</sup>	SynH <sup>2</sup>	SynH <sup>3</sup>
<b>CARBOHYDRATES (mM)</b>				
D-Glucose	343	333	333	333
D-Xylose	266	250	250	269
L-Arabinose <sup>†</sup>	81	-	-	-
D-Galactose <sup>‡</sup>	41	-	34	34
D-Galactose	35	-	34	34
D-Mannose	3.8	-	2.8	2.8
L-Rhamnose	0.8	-	-	-
L-Fucose	0.2	-	-	-
D-Fructose	8.2	-	8.2	8.2
<b>NBC COMPOUNDS (μM)</b>				
Lactate	0.5	-	0.5	0.5
Pyruvate	-	-	30	30
Citrate	-	-	10	10
Malate	-	-	0.1	0.1
Formate	11.2	-	10	10
Miscel	9.3	-	10	10
Succinate	0.8	-	0.5	0.5
Acetate	30	-	23	23
Isocitrate	76	-	60	60
Glucose	5.5	-	3	3
Glycine betaine	0.7	-	0.7	0.7
Choline	0.7	-	0.7	0.7
Carnitine	0.2	-	0.2	0.2
<b>SALT (mM)</b>				
MgCl <sub>2</sub>	-	22	3.4	3.4
K <sub>2</sub> HPO <sub>4</sub>	-	42	0.5	0.5
KCl	-	46	46	46
NaCl	-	24	24	24
KH <sub>2</sub> PO <sub>4</sub>	-	58	58	58
MgCl <sub>2</sub>	-	1	1	1
CaCl <sub>2</sub>	-	0.09	0.09	0.09
<b>AMINO ACIDS (μM)</b>				
Alanine	200	200	200	200
Arginine	270	400	400	400
Asparagine	163	200	200	200
Aspartate	270	200	200	200
Cysteine	n.d. <sup>§</sup>	60	60	60
Glutamine	302	100	100	100
Glutamate	469	460	460	460
Glycine	342	400	400	400
Histidine	50	66	66	66
Isoleucine	8	200	200	200
Leucine	460	380	380	380
Lysine	167	200	200	200
Methionine <sup>¶</sup>	n.d.	100	100	100
Phenylalanine	169	200	200	200
Proline	234	225	225	225
Serine	272	275	275	275
Threonine <sup>¶</sup>	276	228	228	228
Tryptophan	n.d.	58	58	58
Tyrosine	171	175	175	175
Valine	270	225	225	225
<b>NUCLEOTIDES (μM)</b>				
Adenine	-	100	100	50
Cytidine	-	100	100	50
Uridyl	-	100	100	50
Guanine	-	100	100	50
<b>TRACE COMPONENTS (μM)</b>				
Thiamine <sup>¶¶</sup>	-	10	10	10
Parabonate	-	10	10	10
β-methylcrotonic acid	-	10	10	10
p-Hydroxybenzoic acid	-	10	10	10
2,3-Dihydroxybenzoic acid	-	10	10	10
CoCl <sub>2</sub>	-	0.200	0.002	0.002
CuCl <sub>2</sub> ·2H <sub>2</sub> O	-	0.025	0.002	0.002
H <sub>2</sub> SO <sub>4</sub>	-	0.020	10	10
MnSO <sub>4</sub> ·H <sub>2</sub> O	-	0.002	0.002	0.002
FeCl <sub>3</sub>	-	18.6	17	17
ZnCl <sub>2</sub>	-	12	12	12
MnCl <sub>2</sub> ·4H <sub>2</sub> O	-	800	100	100
<b>LC-derived inhibitors</b>				
Penicillinamide	n.d. <sup>§</sup>	n.d. <sup>§</sup>	n.d. <sup>§</sup>	n.d. <sup>§</sup>
Penicillinamide	5.5	3.5 ± 0.6	-	2.75
Clonidineamide	8.8	7.1 ± 1.2	-	2.75
Hydroxyethylthiouracil	1.1	0.63	-	0.55
p-Chloroacetic acid	2.1	1.4 ± 0.2	-	1.05
Resorcinol	0.71	0.70 ± 0.02	-	3.85
Benzoic acid	0.48	0.32 ± 0.01	-	0.48
Synepic acid	0.02	0.02 ± 0.004	-	0.02
Cinnamic acid	0.08	0.08	-	0.09
Vanillic acid	0.08	0.15 ± 0.02	-	0.09
Caffeic acid	0.08	0.00 ± 0.001	-	0.09
Salicylic acid	0.122	0.21 ± 0.04	-	0.122
Synepic acid	0.162	0.07 ± 0.002	-	0.162
β-Hydroxybenzoic acid	0.187	0.11 ± 0.02	-	0.187
4-Hydroxyacetophenone	0.129	0.07 ± 0.002	-	0.129
Osmolite (mM)	110 ± 0.03	0.87	1.17 ± 0.01	1.19 ± 0.01



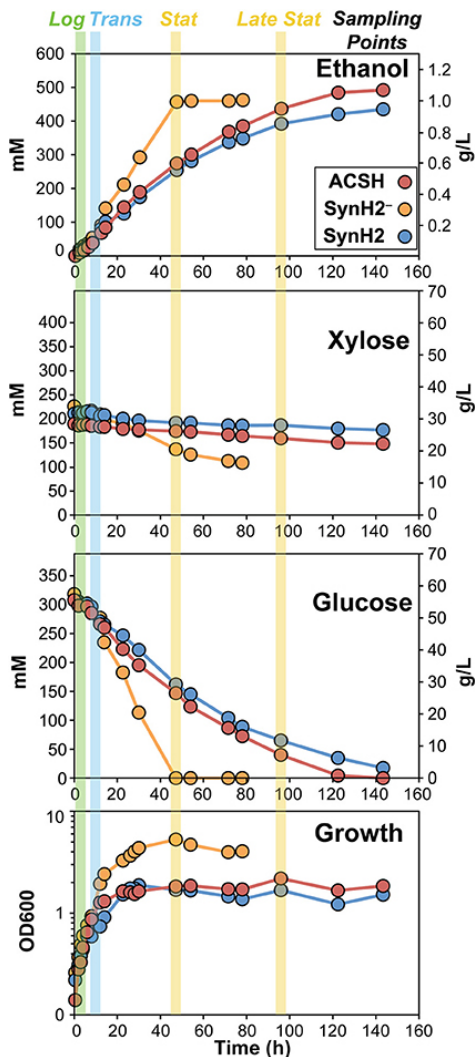
- Stress related
- Efflux pumps
- Anaerobic respiration
- Carbohydrate use (non-glucose)
- Amino acid synthesis
- Flagella, Chemotaxis
- Thiamine synthesis
- Iron metabolism
- Citrate lyase
- Osmotic tolerance

Keating, et al. 2014 *Frontiers Micro.* 5:402  
Schwalbach, et al. 2012 *App. Env. Micro.* 78:3442

# Reiterative microbial design targets sites for<sup>6</sup> strain improvement



# Hydrolysate composition negatively impacts biofuel fermentations

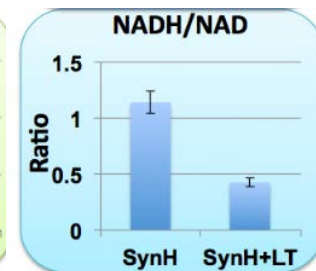
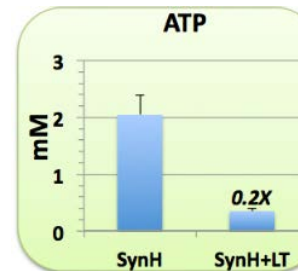
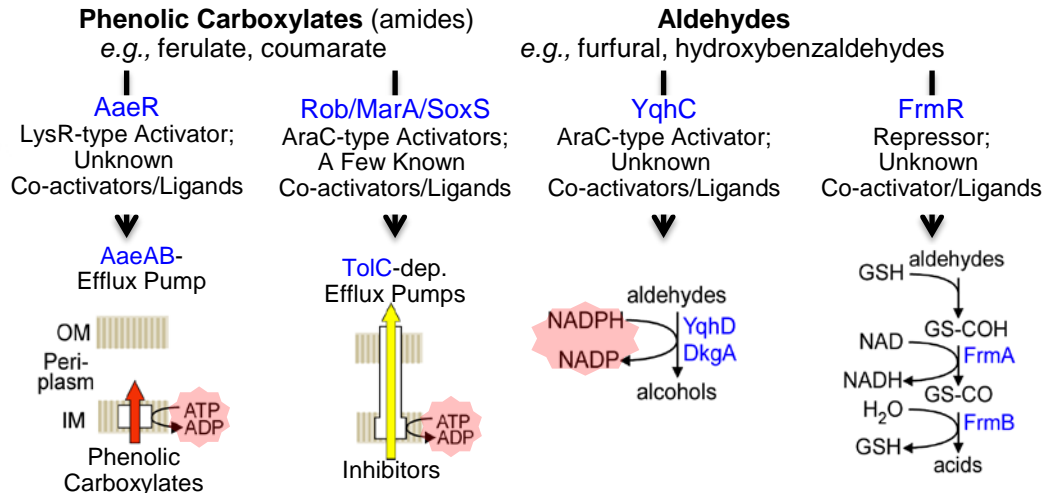
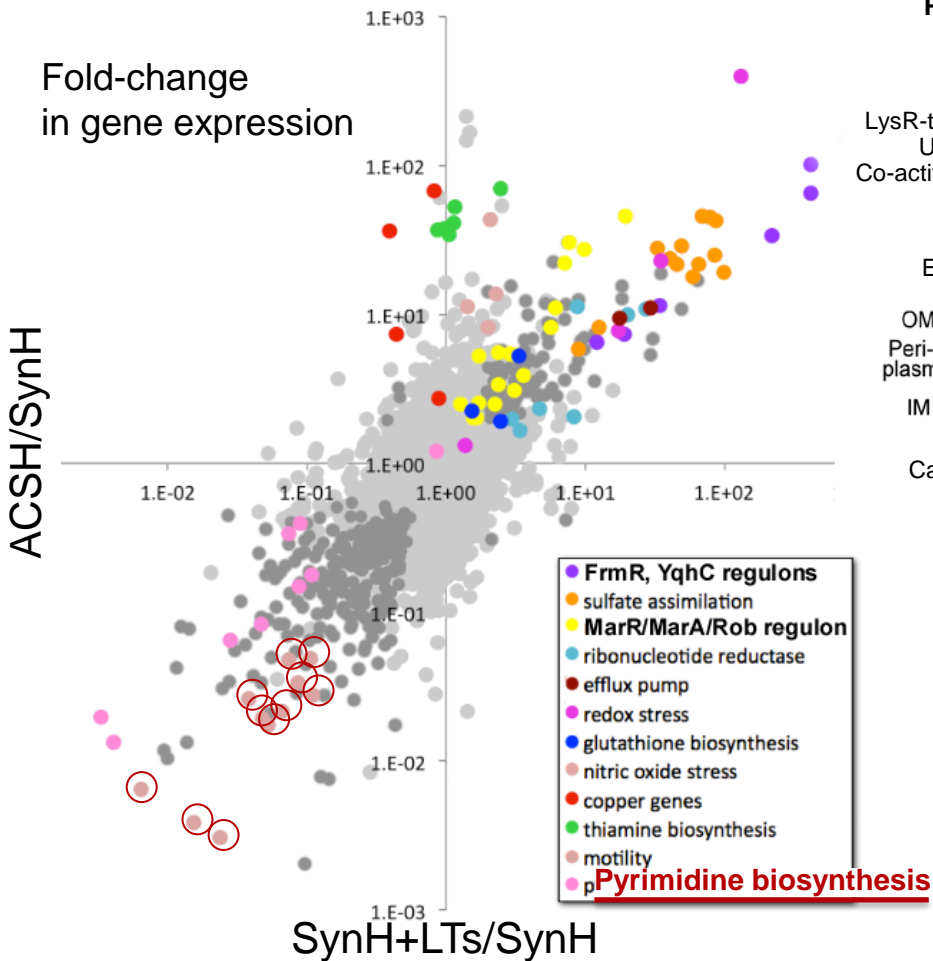


- ✧ Negative impact of ACSH on growth/biomass production, xylose and glucose utilization & ethanol production
- ✧ Recapitulate behavior with synthetic hydrolysate (SynH2)
- ✧ Negative impact overcome in SynH2- (SynH2 minus aromatics from ACSH)

## Lignotoxins (LTs)

- 5.5 mM Feruloyl amide
- 5.5 mM Coumaroyl amide
- 2.1 mM Coumaric acid
- 0.5 mM Benzoic acid
- 1.1 mM Hydroxymethylfurfural
- 0.7 mM Ferulic acid
- 8 minor LTs (<0.5 mM)

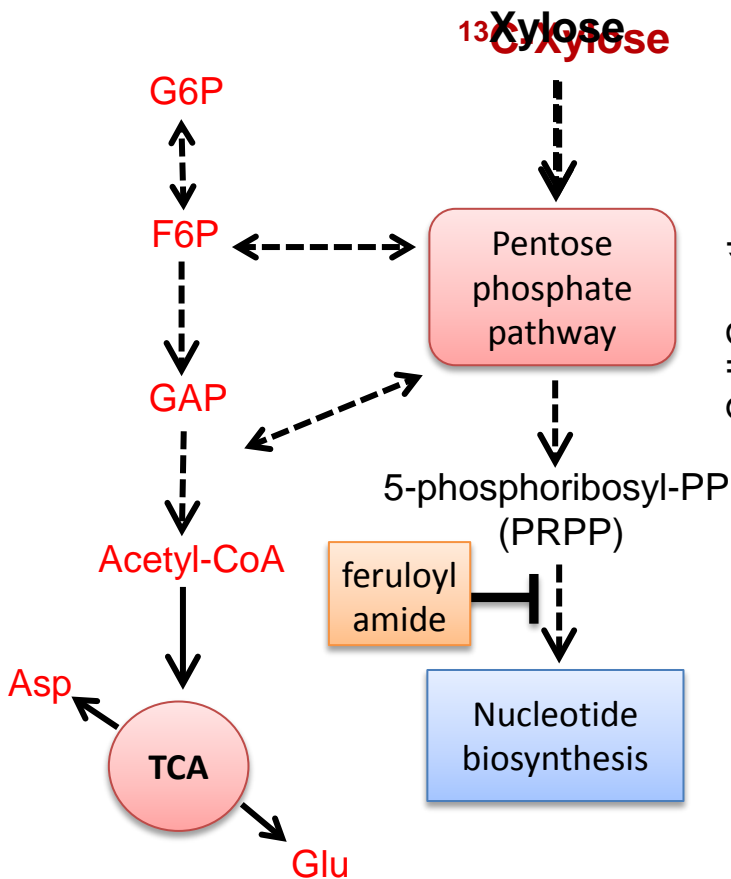
# LTs induce energy-consuming pathways & deplete ATP/NADH



✧ Keating, et al. 2014 *Frontiers Micro.* 5:402

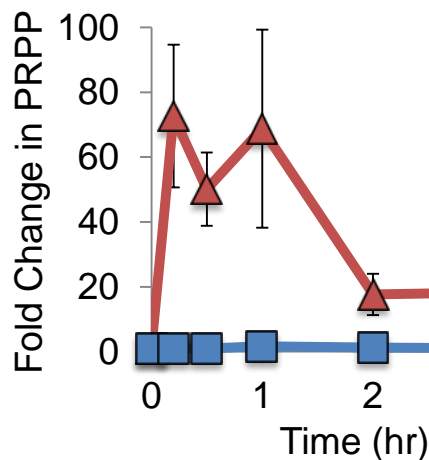
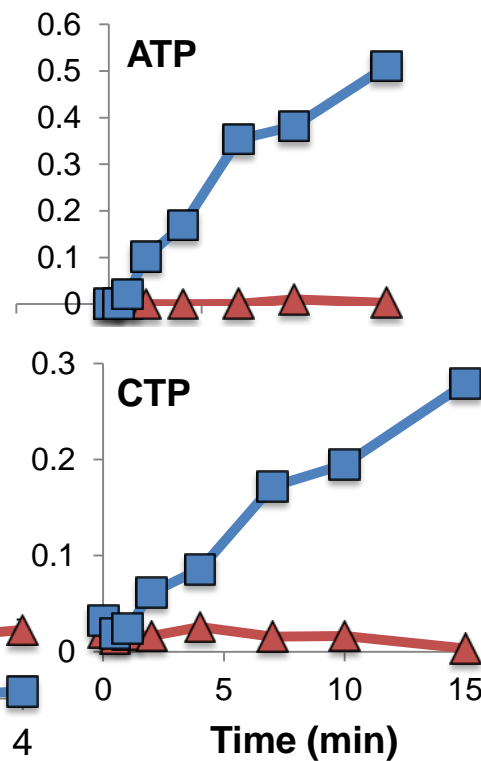
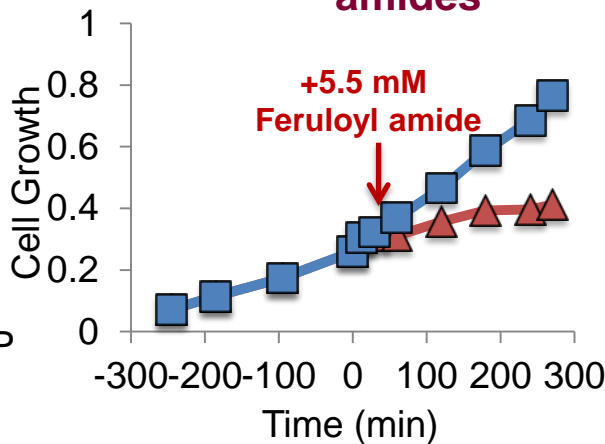
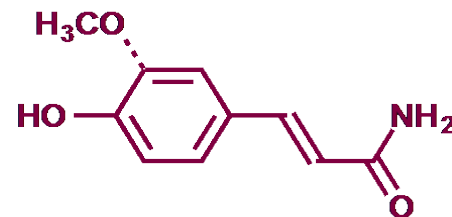


# Feruloyl amide inhibits growth, increases PRPP pools & blocks nucleotide synthesis

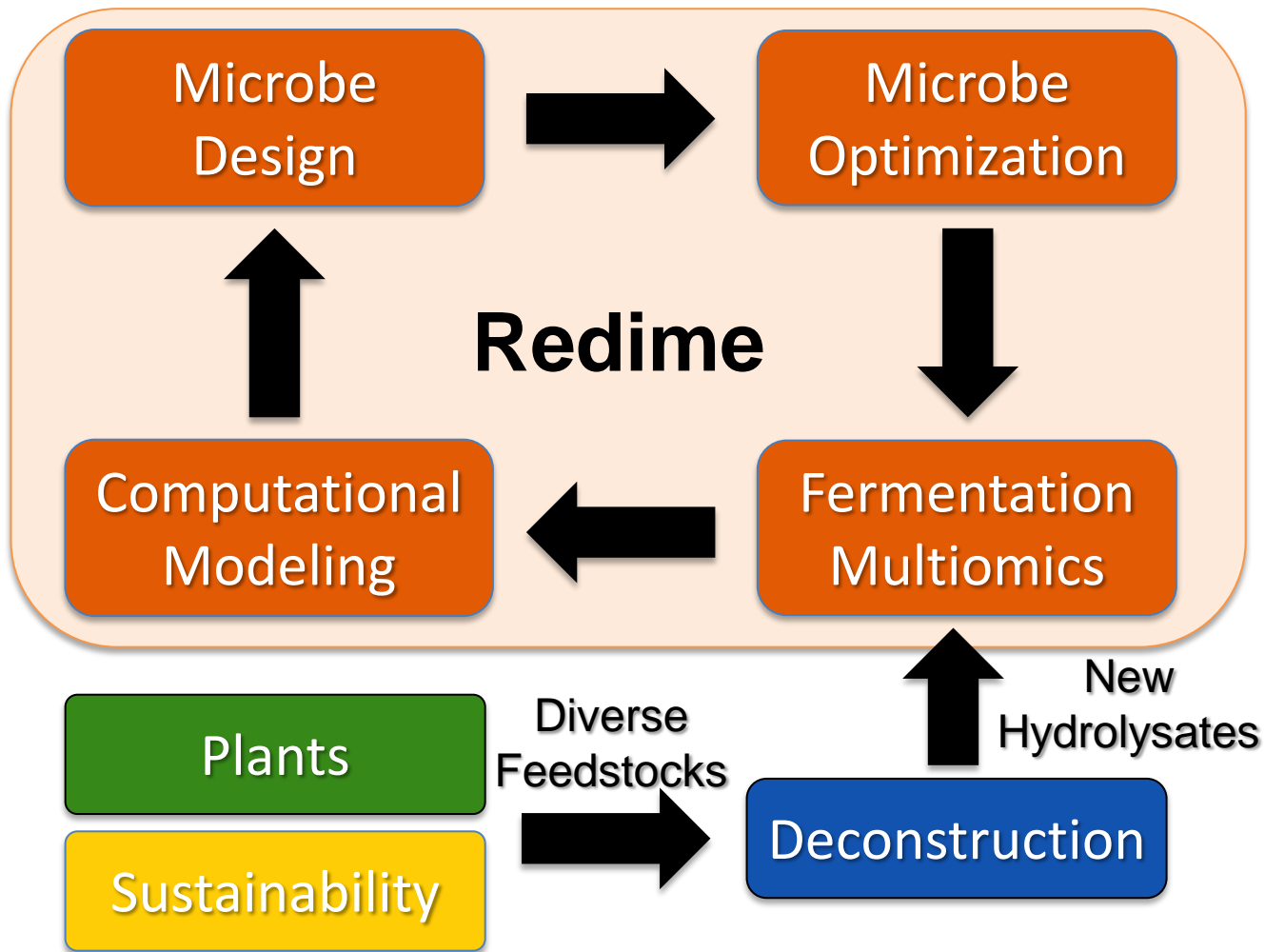


Amador-Noguez, et al. in preparation

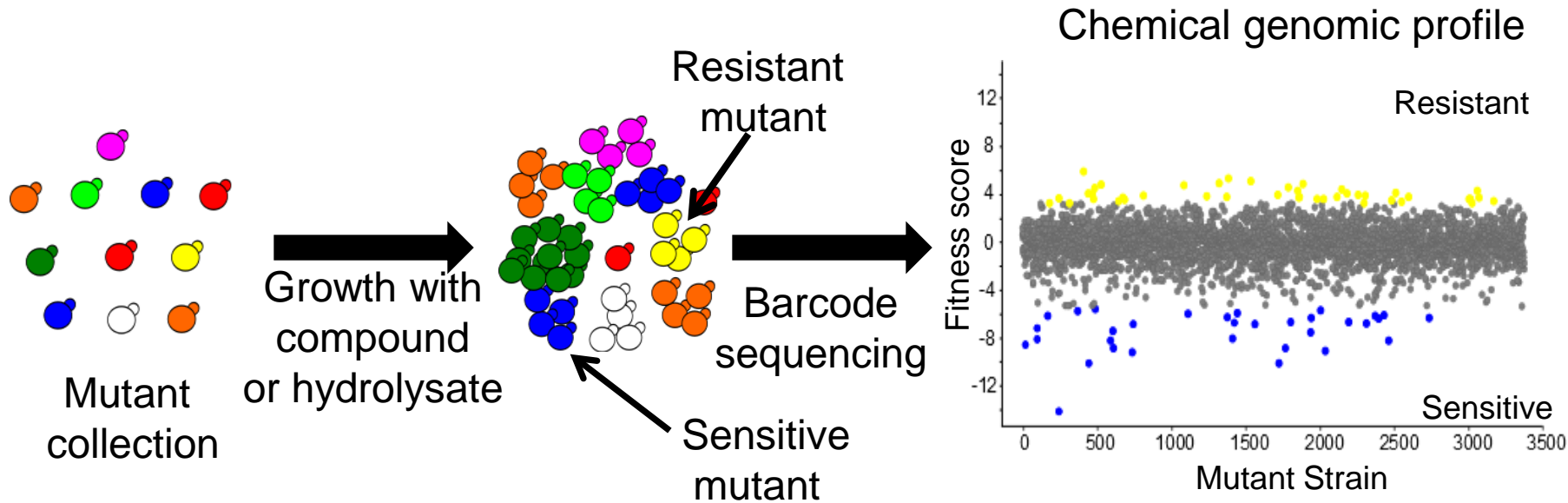
(Feruloyl) Coumaroyl amides



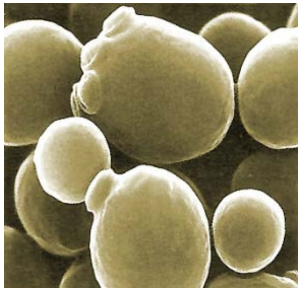
# Reiterative microbial design targets sites for strain improvement<sup>10</sup>



# Genomic fingerprinting of inhibitors



*S. cerevisiae*



Parsons, et al. 2006  
Piotrowski, et al. in press PNAS

*Z. mobilis*



Skerker, et al. 2013

*E. coli*



Otsuka, et al. 2015

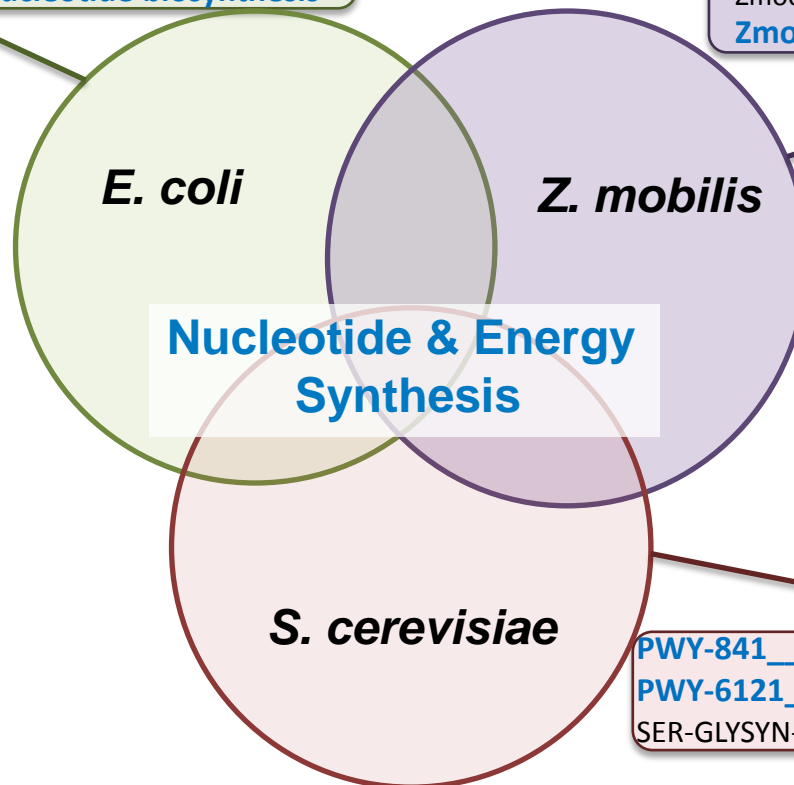
# Conserved mode of action of LT's

Chemical genomic analysis of 29 hydrolysate LT's against 3 "disparate" microbes (*E. coli*, *Z. mobilis*, *S. cerevisiae*)

✧ Nucleotide & energy metabolism responsive across inhibitors & microbes

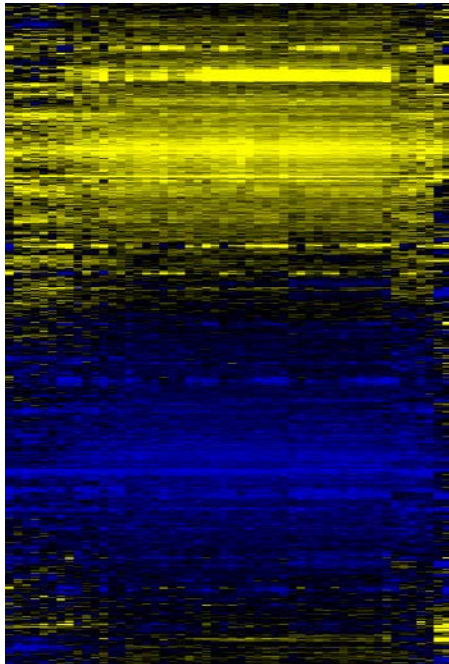
PWY0-1334\_\_electron transfer from NADH to cytochrome *bd* oxidase  
PURINE2-PWY\_\_*de novo* purine nucleotide synthesis  
PWY-7219\_\_*de novo* adenosine nucleotide biosynthesis

Zmo00290 Valine, leucine and isoleucine biosynthesis  
Zmo00340 Histidine metabolism  
Zmo00230 Purine metabolism

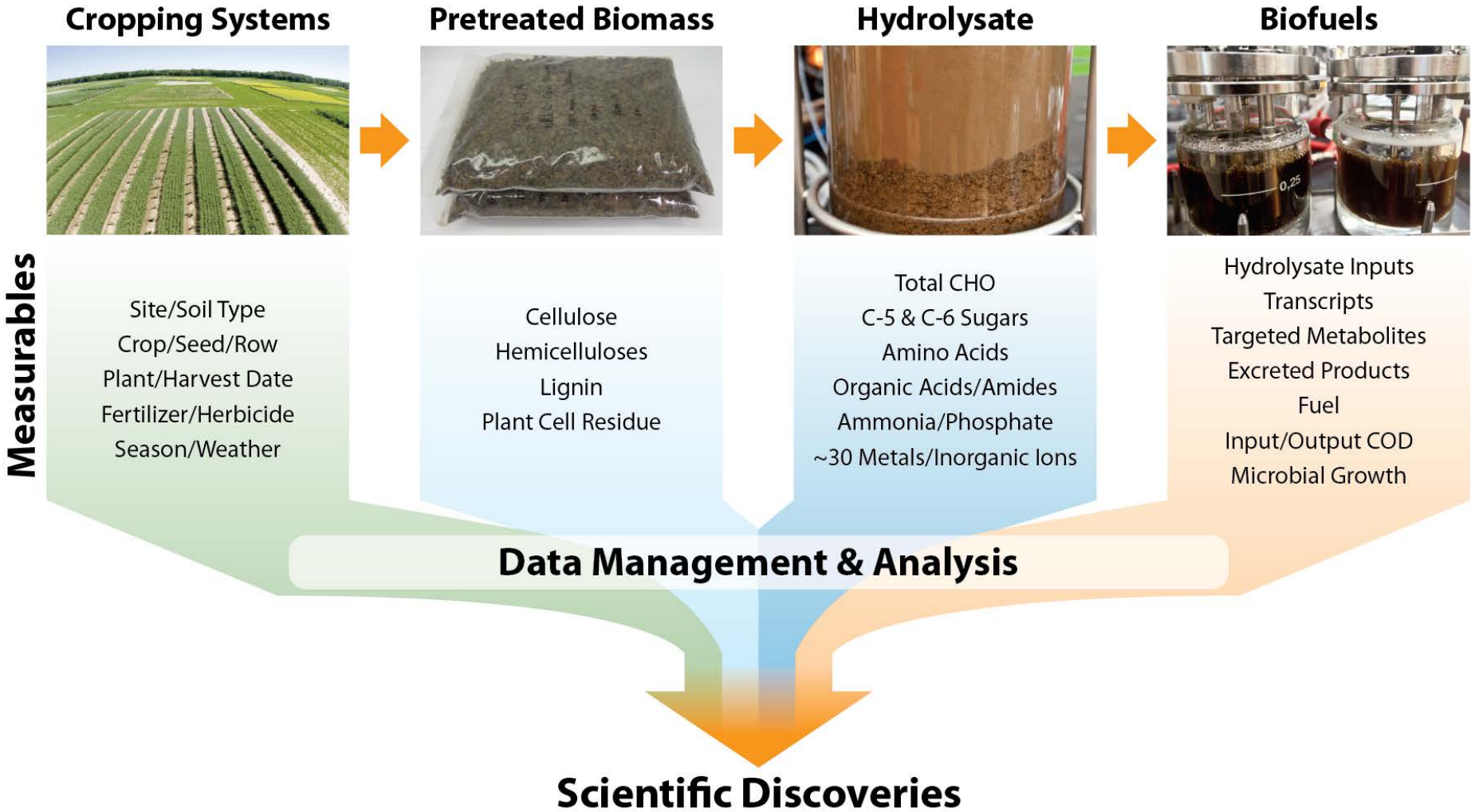


*E. coli*: Otsuka, *et al.* 2015  
*Z. Mobilis*: Skerker, *et al.* 2013

PWY-841\_\_*de novo* purine biosynthesis  
PWY-6121\_5-aminoimidazole ribonucleotide synthesis  
SER-GLYSYN-PWY\_\_Serine & glycine synthesis



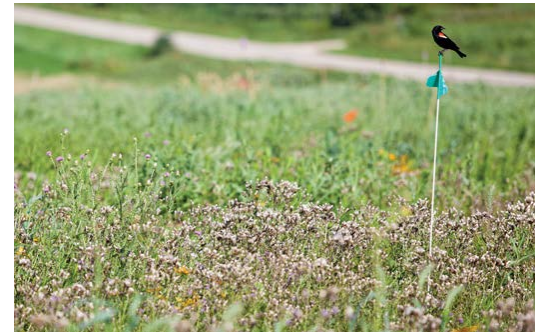
# Systems Analysis of Biomass & Biofuel Production



Measurables



# Systems Analysis of Biomass & Biofuel Production



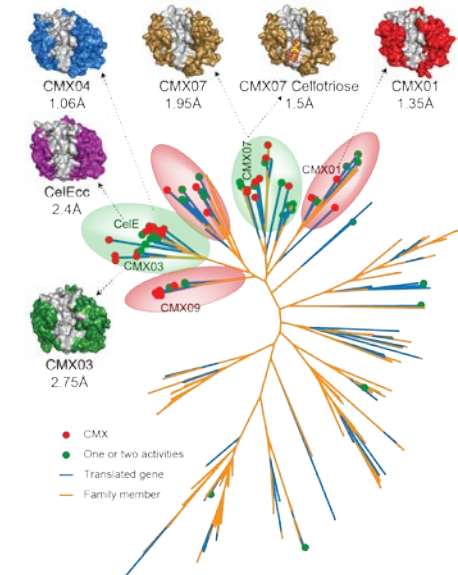
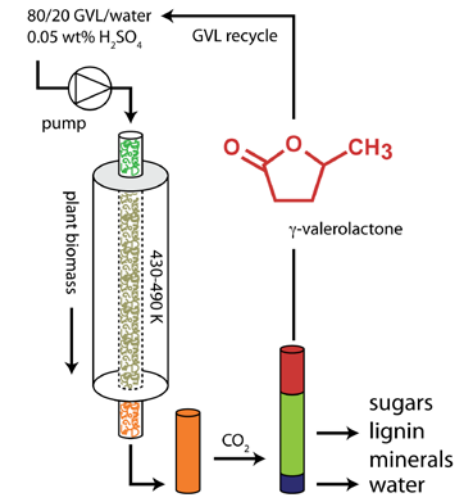
## Analysis of additional biomass crops

- ✘ Poplar
- ✘ Miscanthus
- ✘ Switchgrass
- ✘ Native Prairie
- ✘ Mixed Feedstocks
- ✘ Assess yearly/regional feedstock variations
- ✘ Impact of biomass trait modifications (Zip-Lignin & others)

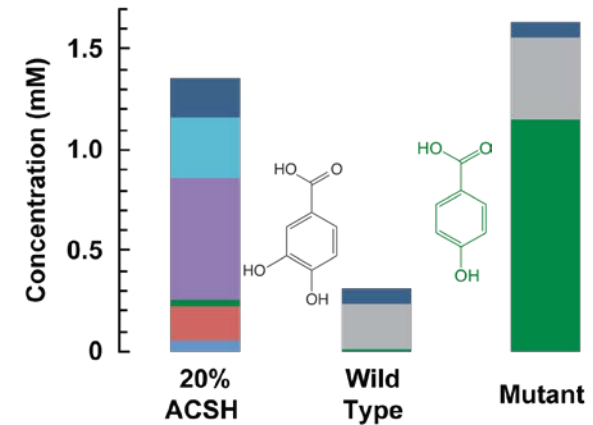
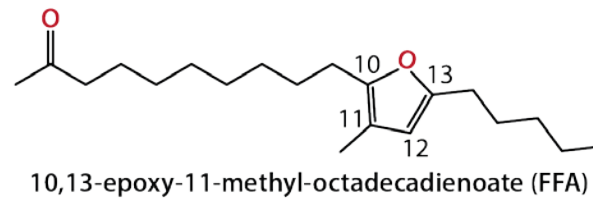
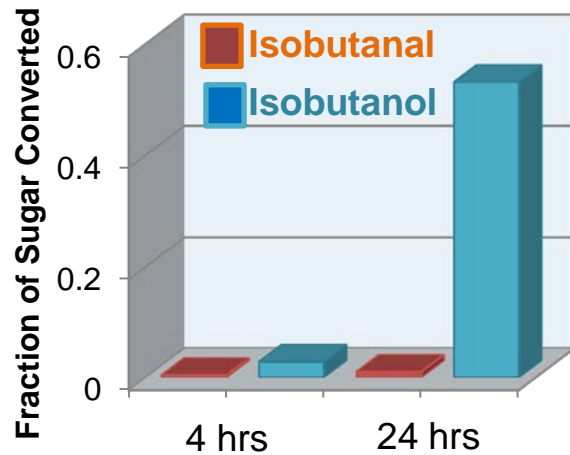
# Systems Analysis of Biomass & Biofuel Production

Analysis of other “polysaccharide hydrolysates” & *lignin streams* from other pretreatments

- ✧  $\gamma$ -Valerolactone (GVL)
- ✧ Alkaline (alkaline hydrogen peroxide, extractive ammonia)
- ✧ Ionic liquids
- ✧ Monitor variations due to changes in enzyme cocktails



# Systems Analysis of Biomass & Biofuel Production



## Analysis of other microbial catalysts

- ✦ Additional fuels & chemicals (long chain alcohols, hydrocarbons, aromatics, etc.)
- ✦ Impact of changes on producing microbes
- ✦ Different single species/consortia



# Questions?