

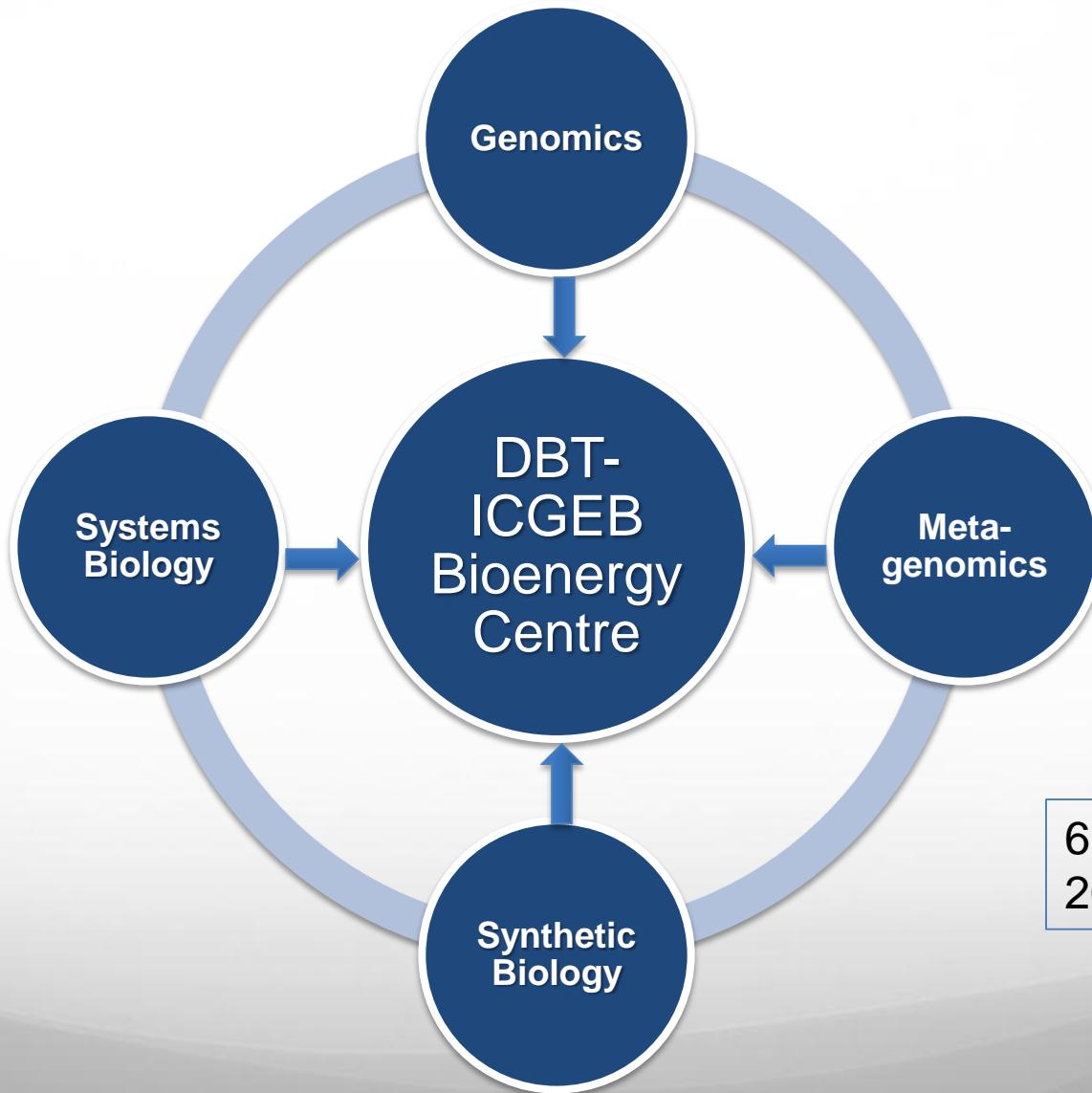


Molecular interventions towards bioenergy

Dr. Syed Shams Yazdani

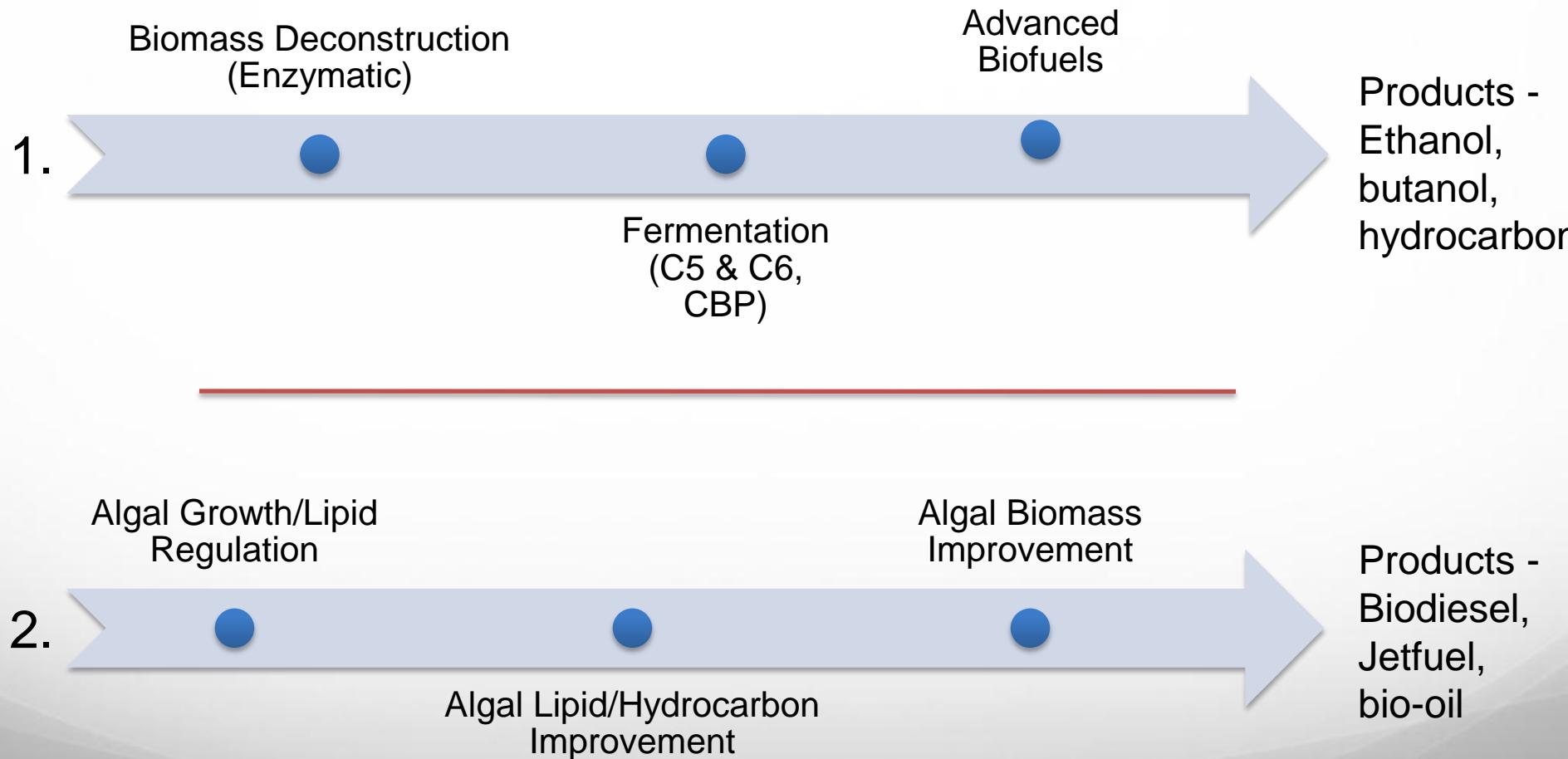
**Synthetic biology and Biofuels Group
DBT-ICGEB Centre for Advanced Bioenergy Research
International Centre for Genetic Engineering and Biotechnology
New Delhi**

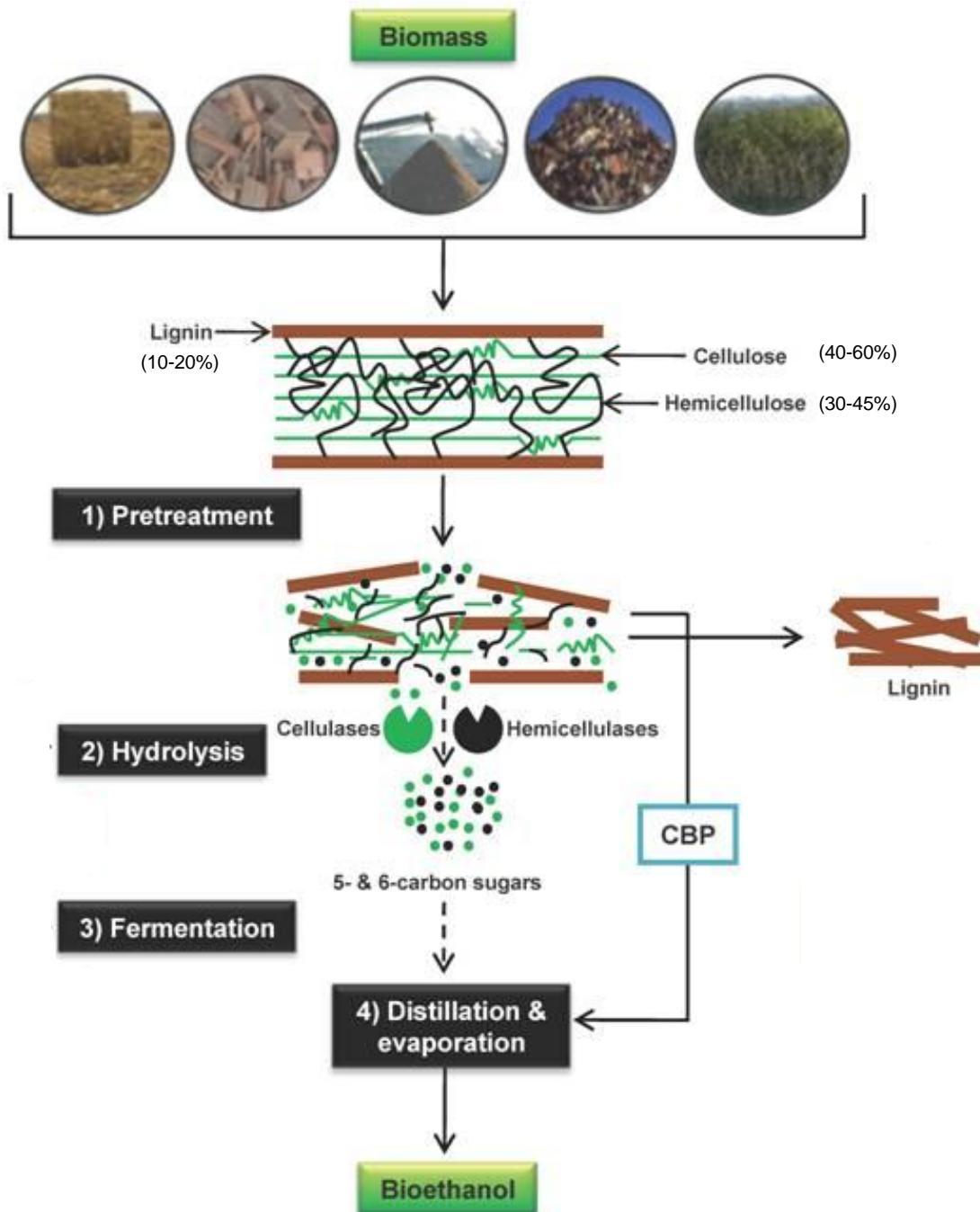
Four Pillars of Our Research

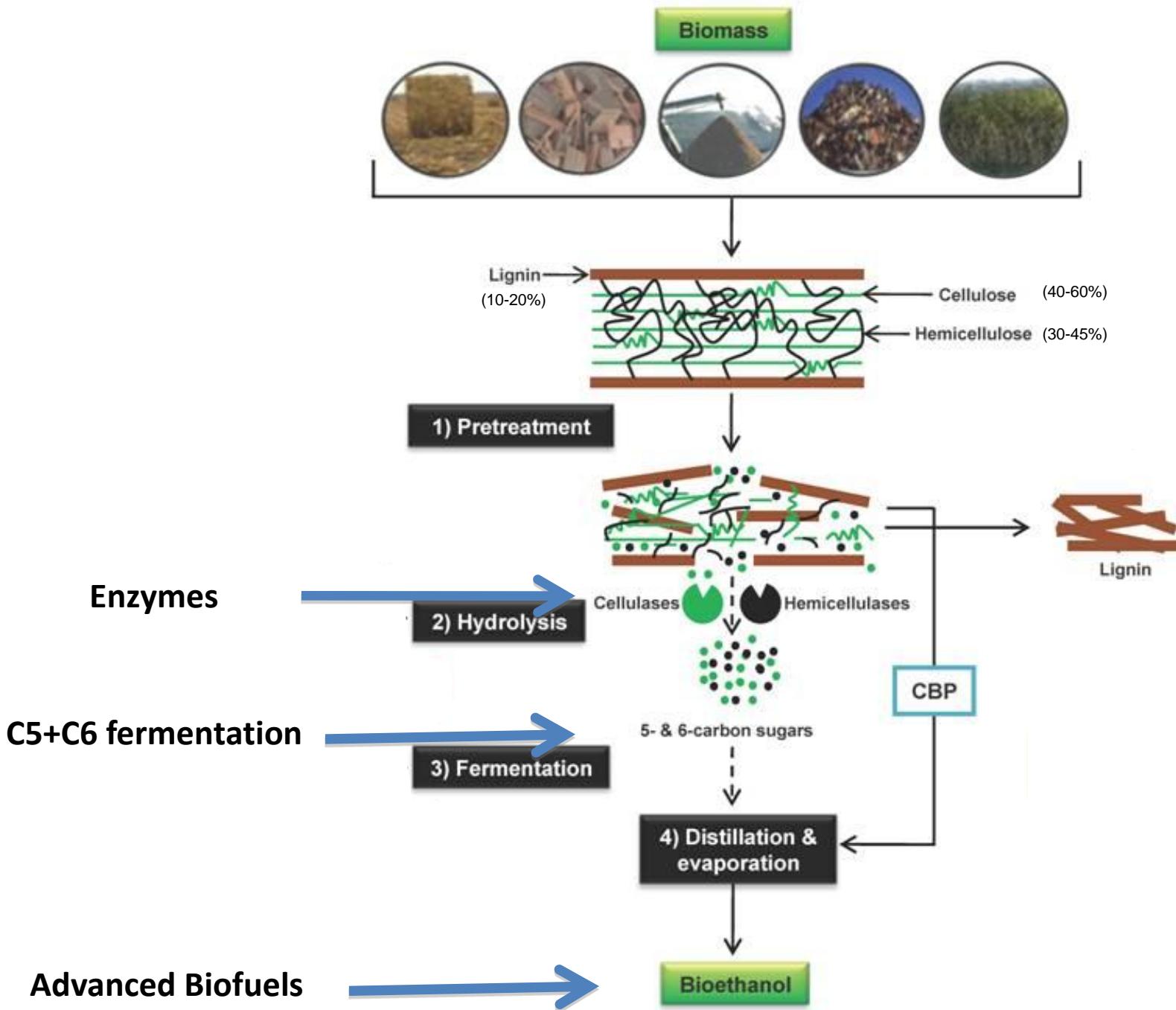


6 investigators
20 students

Research Theme at the Bioenergy Centre

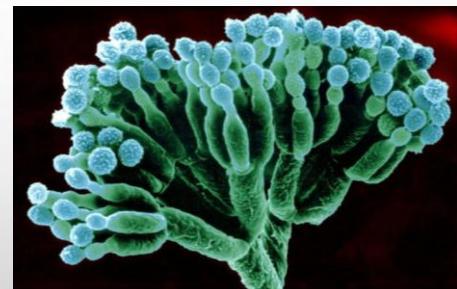
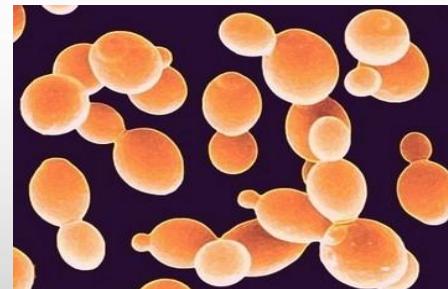
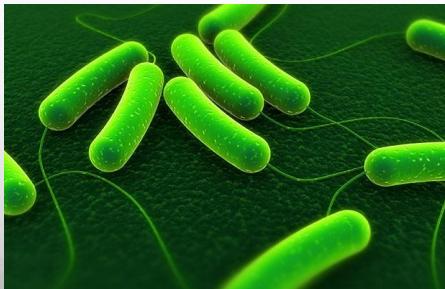




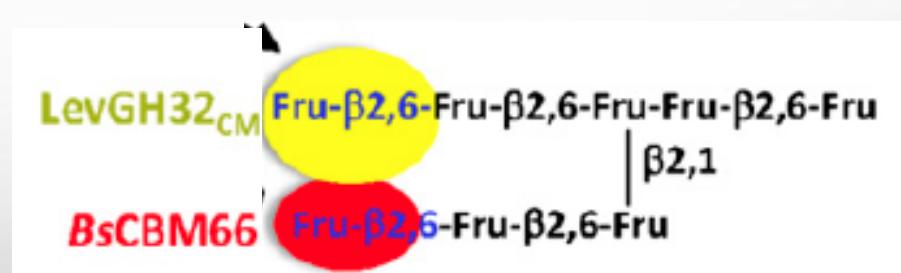
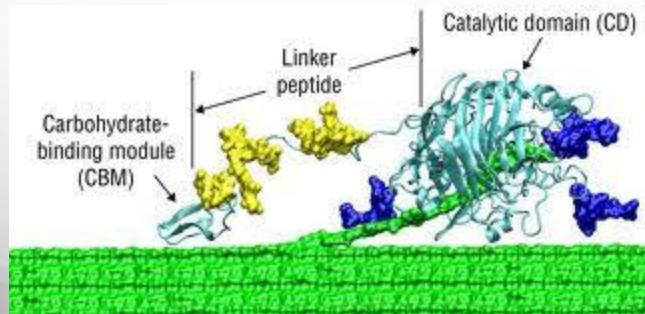
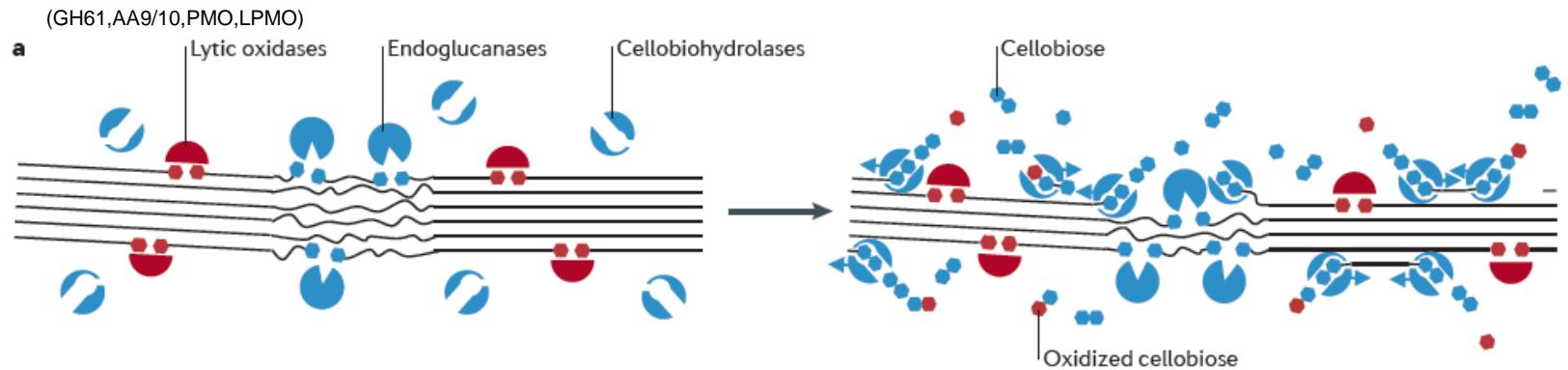


Enzyme Research

- Mainly to identify and design enzymes for efficient degradation of lignocellulosic biomass
- Source of enzymes are either free living fungi or microbes residing in insect guts that destroy plants
- Expression platforms are either native hosts or recombinant hosts such as *E. coli*, yeast or fungi



Cellulolytic enzymes and CBMs



From Literatures

Identifying novel cellulolytic enzymes



Cotton Bollworm



Rice Stem Borer



Termite



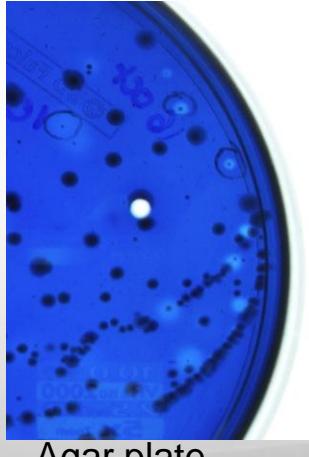
Pillbug



Gut screened
for cellulase producer



Strain I

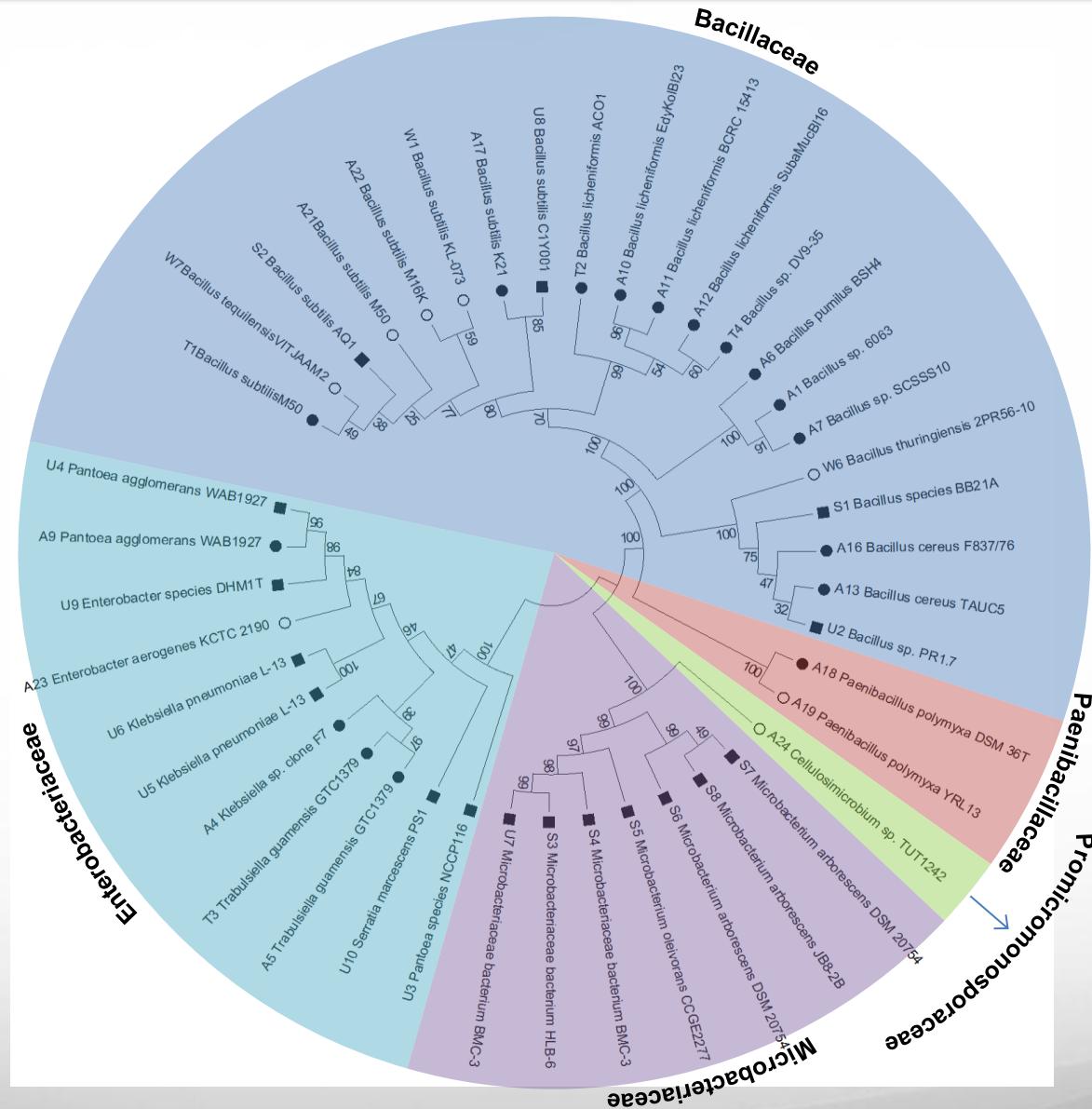


Agar plate

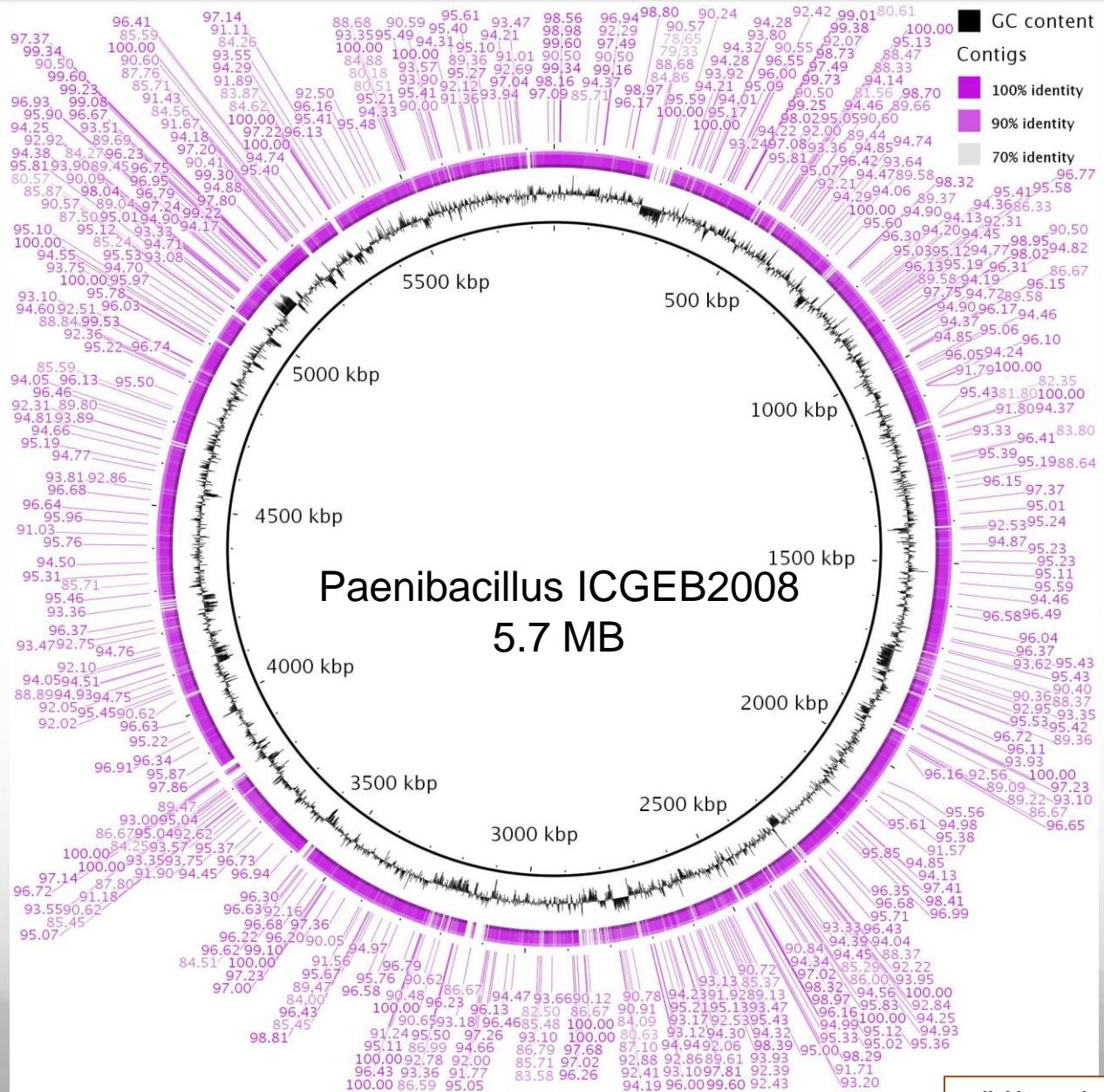


Strain II

Phylogeny of Gut Cellulolytic Microbes

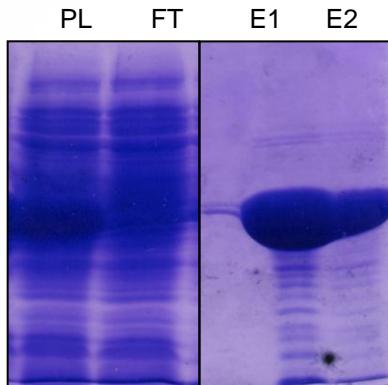


Genome Sequencing of *Paenibacillus ICGEB2008*



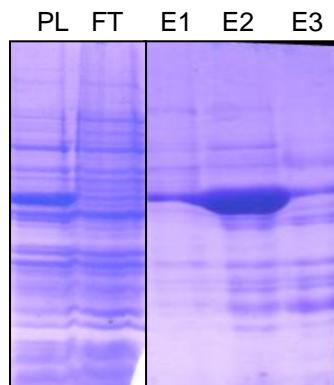
Novel cellulolytic enzymes overexpressed with high specific activity

Endocellulase – 41 kDa



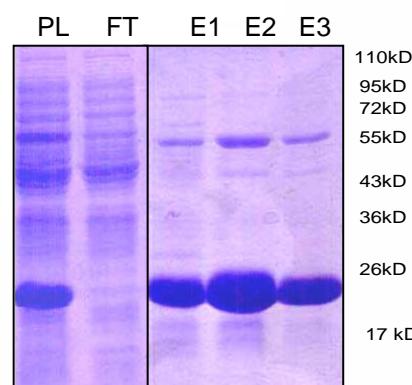
pH optima ~ 6-7
Temp optima ~50 °C

β -glucosidase – 51 kDa



pH optima ~ 6
Temp optima ~50 °C

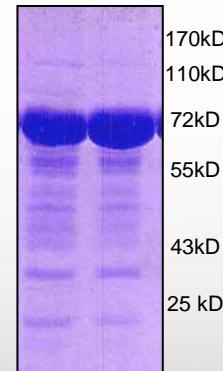
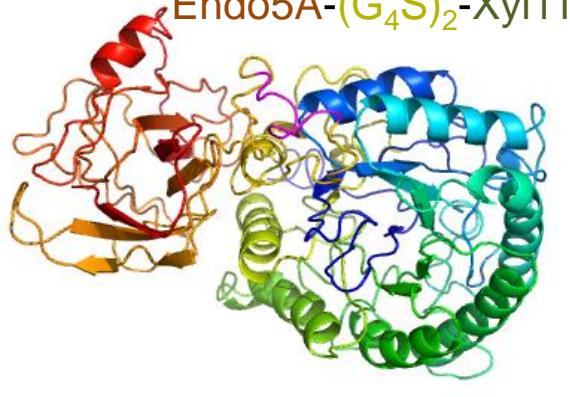
Xylanase – 21 kDa



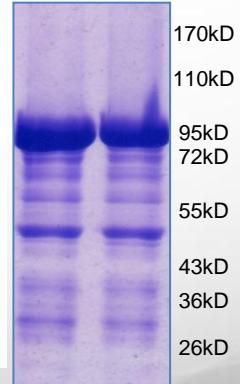
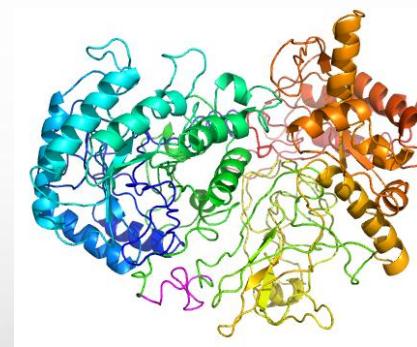
pH optima ~ 6-7
Temp optima ~50 °C

110kD
95kD
72kD
55kD
43kD
36kD
26kD
17 kD

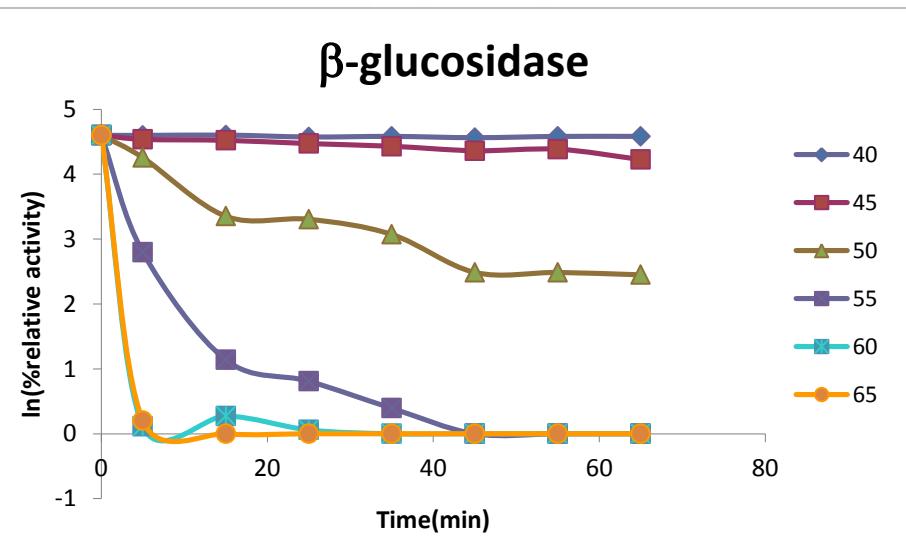
Endo5A-(G₄S)₂-Xyl11D



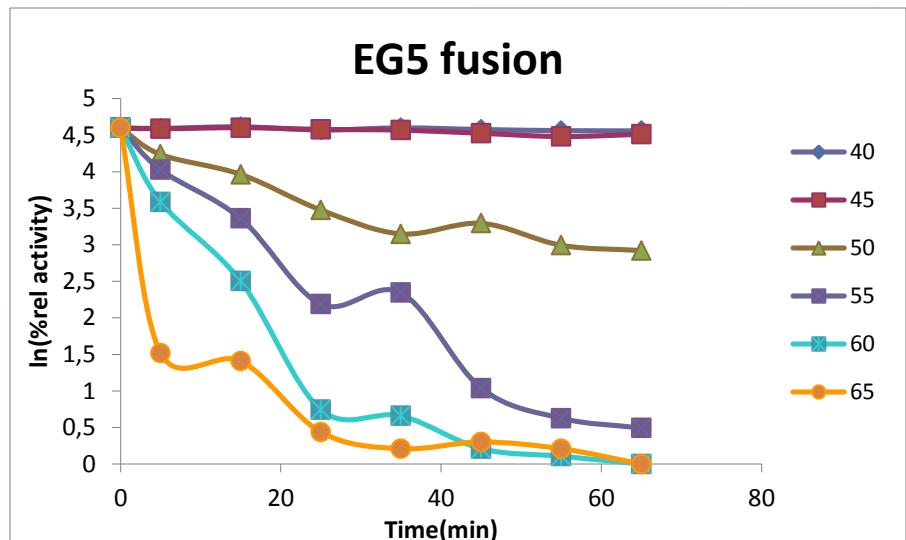
Endo5A-(G₄S)₃-Gluc1C



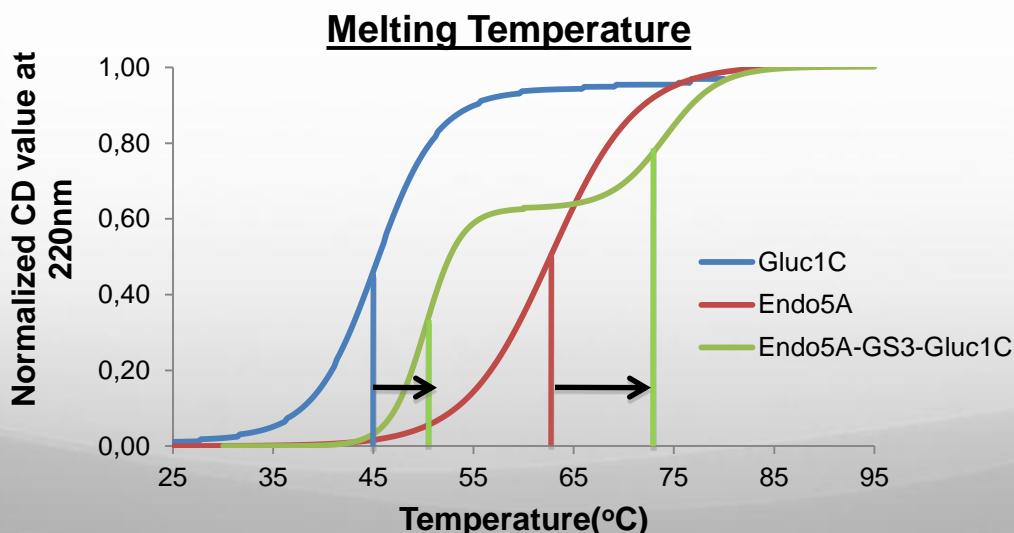
Fusion enhances thermostability



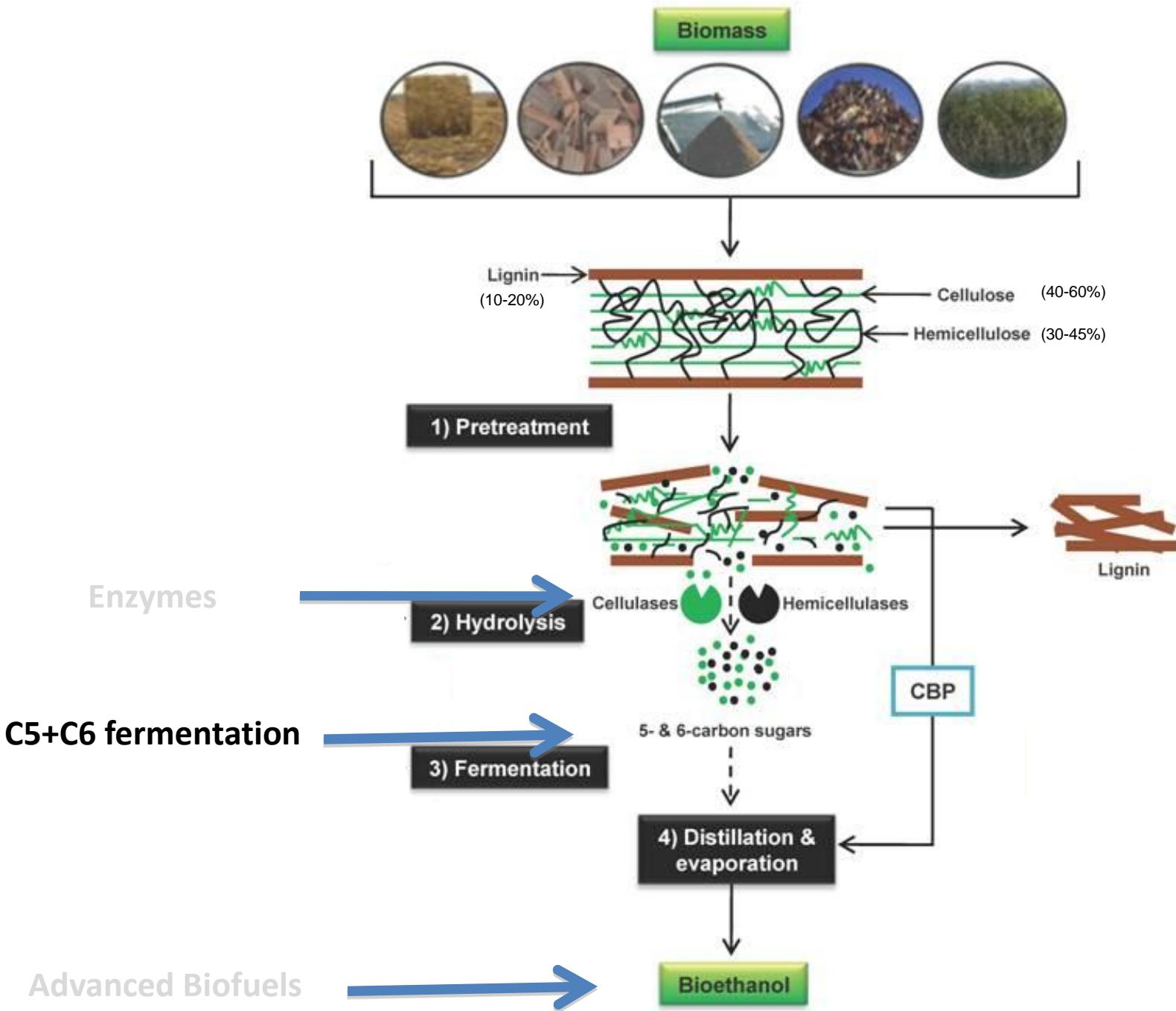
Half life of activity at 55 °C – 1.9 min



Half life of activity at 55 °C – 7.5 min



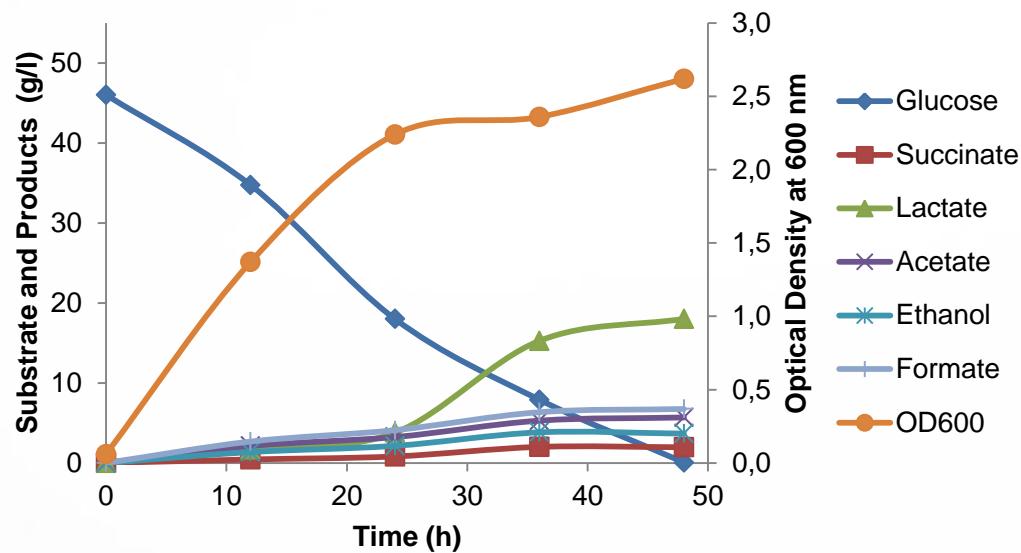
Secondary
structure more
stable in fusion



Engineering microbe for converting C5/C6 sugars to ethanol

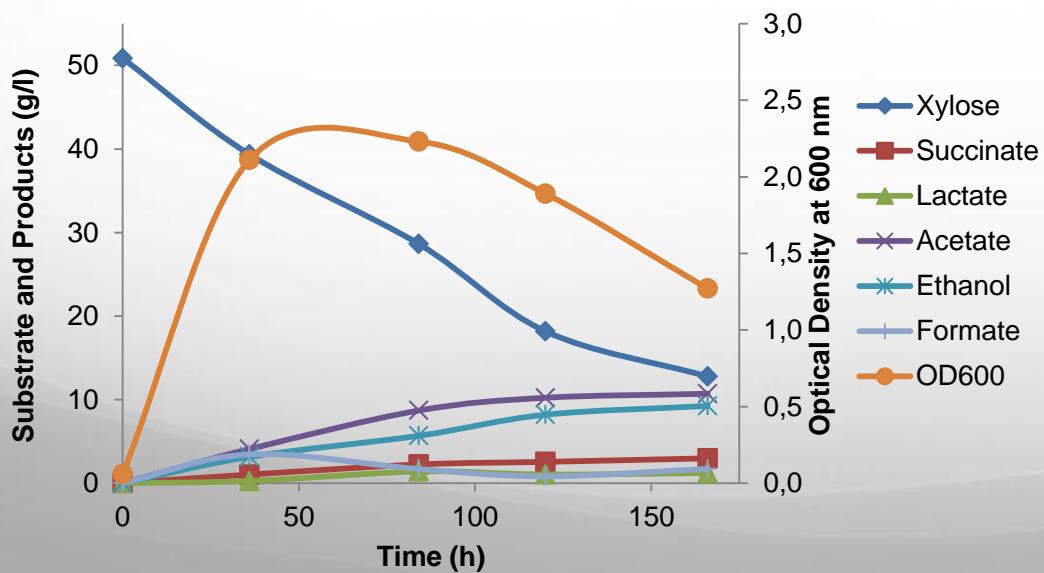
- Up to 30% pentose sugars in agricultural residues
- Traditional yeast, *Saccharomyces cerevisiae*, cannot utilize pentose sugars
- *E. coli* could ferment all pentose and hexose sugars present in the lignocellulosic biomass
- It, however, produces various competing products under anaerobic condition

Competing co-products of *E. coli* under fermentative condition



Max Ethanol Productivity = 0.14 g/l/h

Ethanol Yield = 0.08 g/g glucose

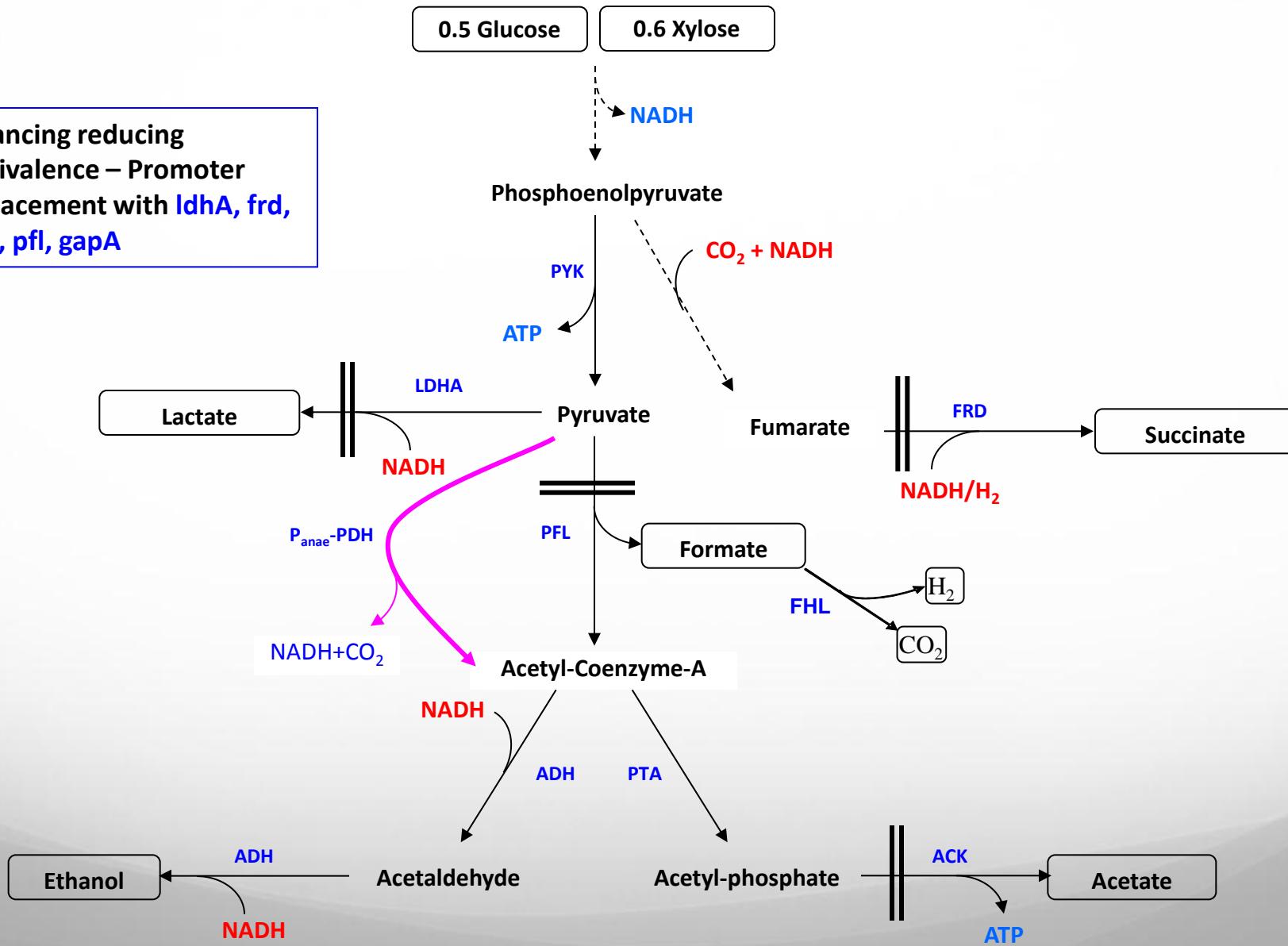


Max Ethanol Productivity = 0.09 g/l/h

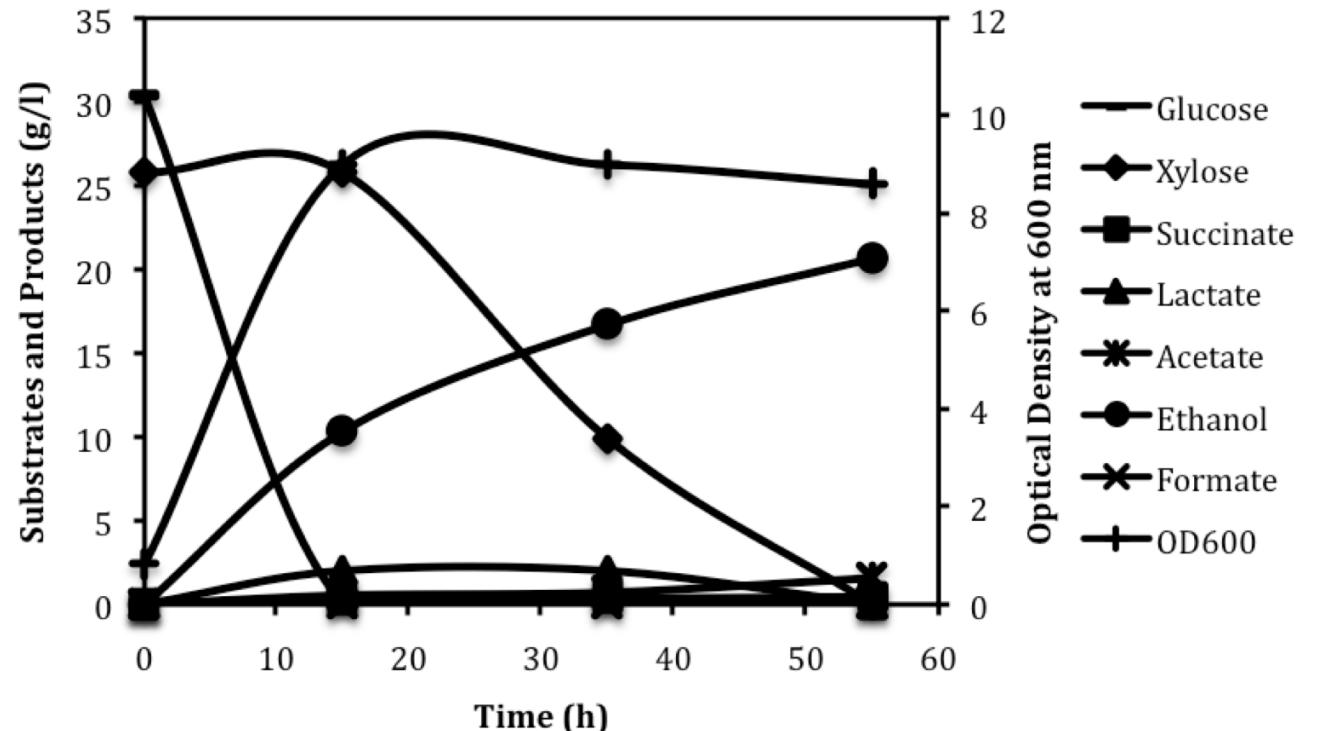
Ethanol Yield = 0.24 g/g xylose

Engineering *E. coli* for ethanol production

Balancing reducing equivalence – Promoter replacement with **ldhA, frd, adh, pfl, gapA**



Co-Fermentation of Glucose and Xylose by SSY10

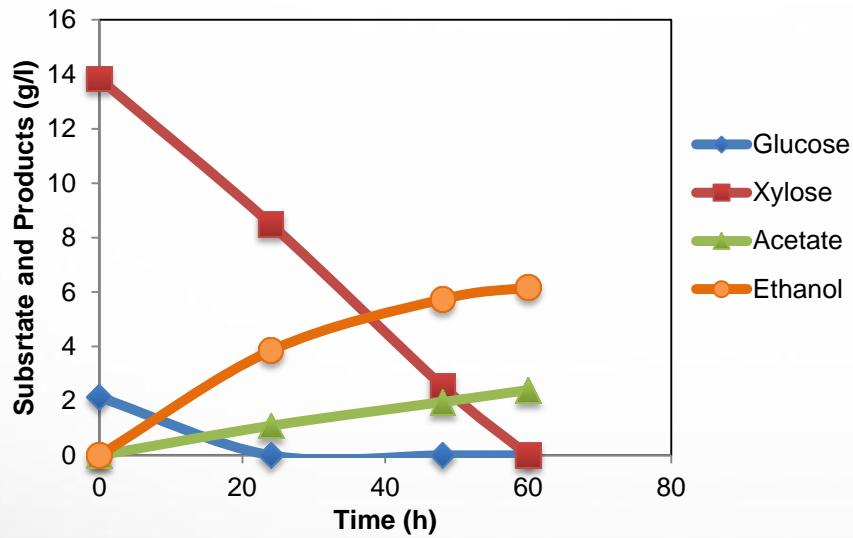


Ethanol Productivity = 0.7 g/l/h

Ethanol Yield = 0.43 g/g glucose

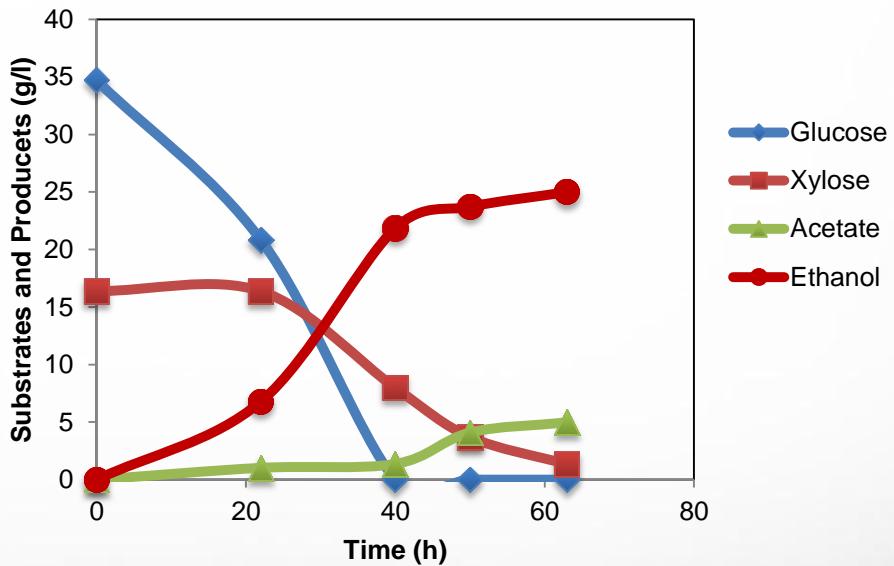
Second generation lignocellulosic ethanol

Acid Treated Rice Straw



Ethanol Yield = 0.4 g/g

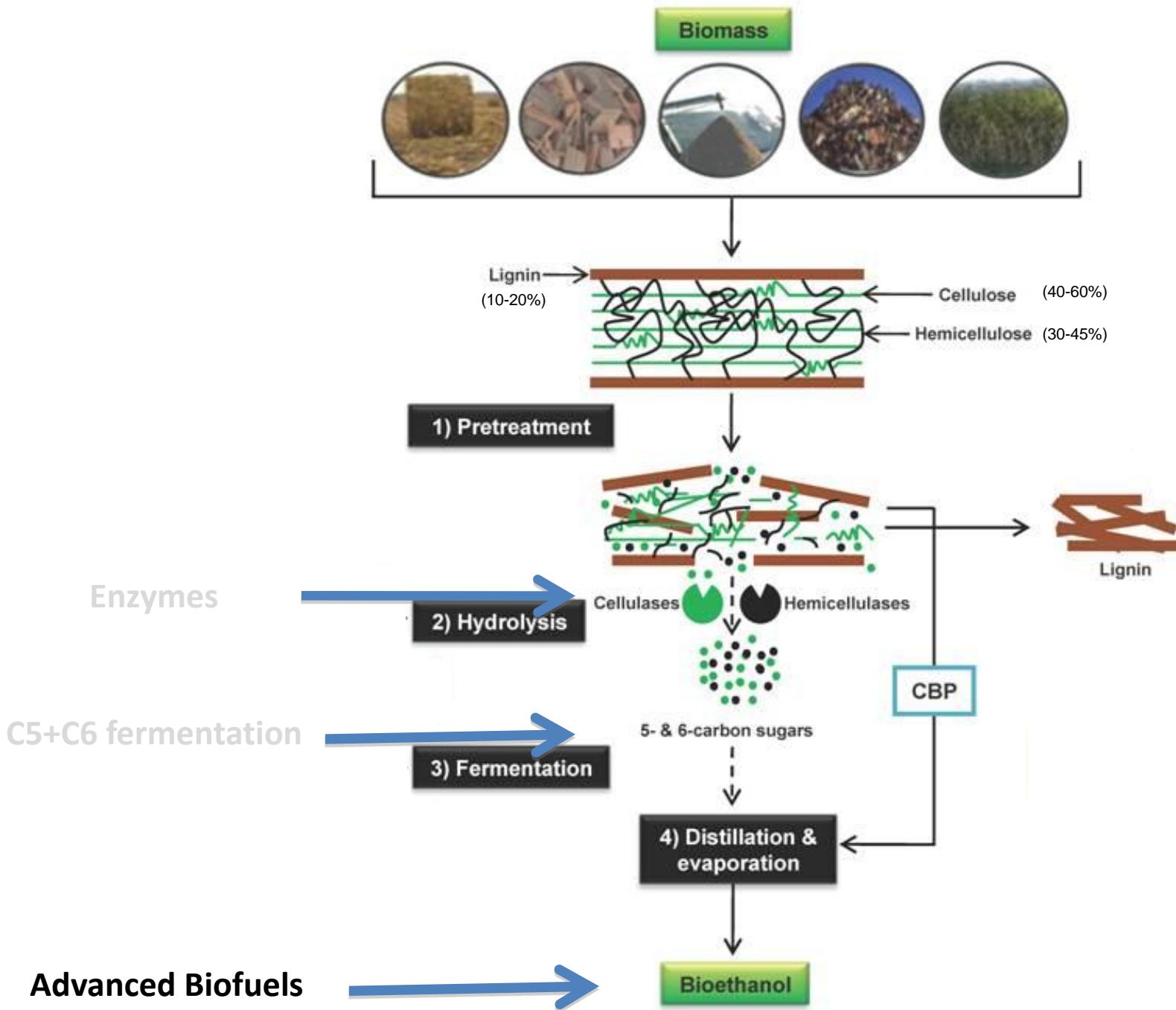
Ammonia Treated Rice Straw



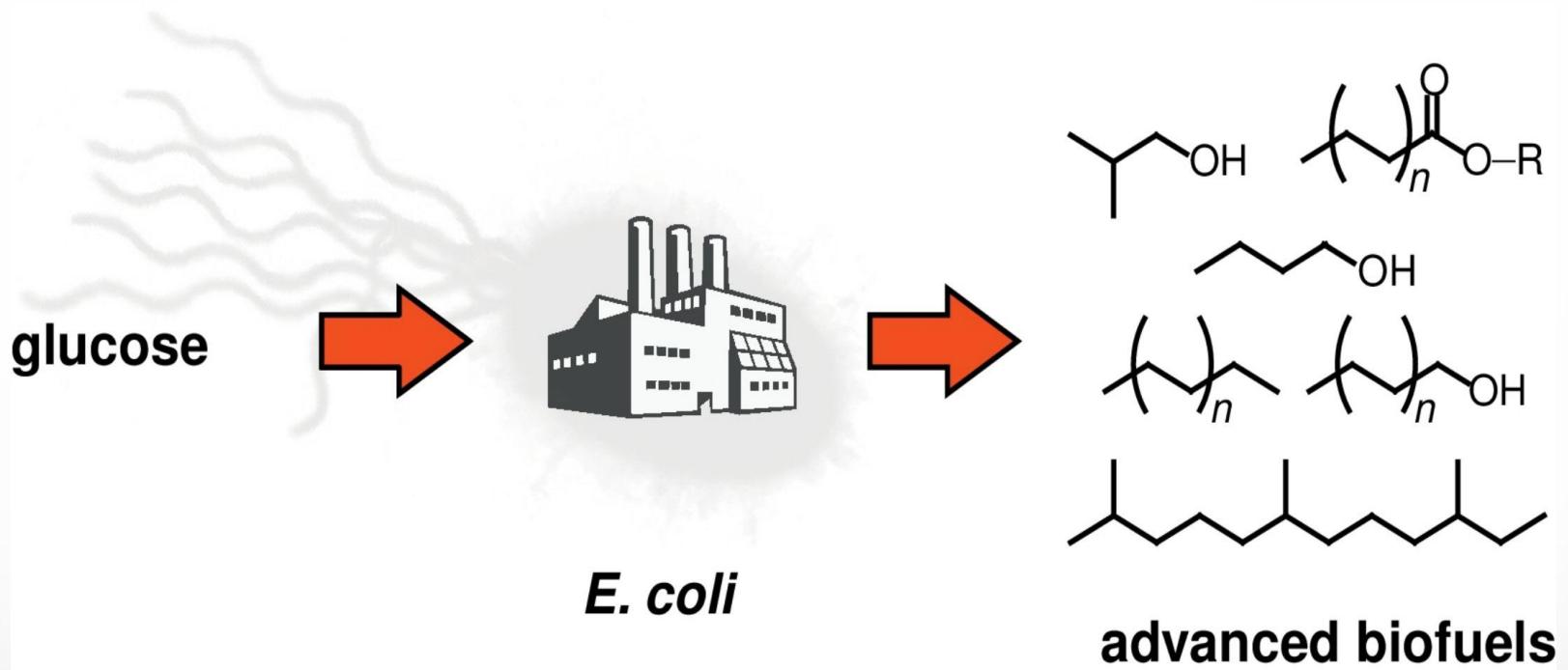
Ethanol Yield = 0.5 g/g

Pre-treated Biomass from IOCL

Pre-treated Biomass from IGL/ICT



Advanced Biofuels



LETTERS

Non-fermentative pathways for synthesis of branched-chain higher alcohols as biofuels

Shota Atsumi¹, Taizo Hanai¹ & James C. Liao^{1,2}



Microbial Biosynthesis of Alkanes

Andreas Schirmer, *et al.*
Science **329**, 559 (2010);
DOI: 10.1126/science.1187936

Synthesis of customized petroleum-replica fuel molecules by targeted modification of free fatty acid pools in *Escherichia coli*

Thomas P. Howard^a, Sabine Middelhaufe^a, Karen Moore^a, Christoph Edner^a, Dagmara M. Kolak^a, George N. Taylor^a, David A. Parker^{a,b}, Rob Lee^{a,b}, Nicholas Smirnoff^a, Stephen J. Aves^a, and John Love^{a,1}

^aBiosciences, College of Life and Environmental Sciences, University of Exeter, Exeter EX4 4QD, United Kingdom; and ^bBiodomain, Shell Technology Centre Thornton, Chester CH1 3SH, United Kingdom

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RESEARCH

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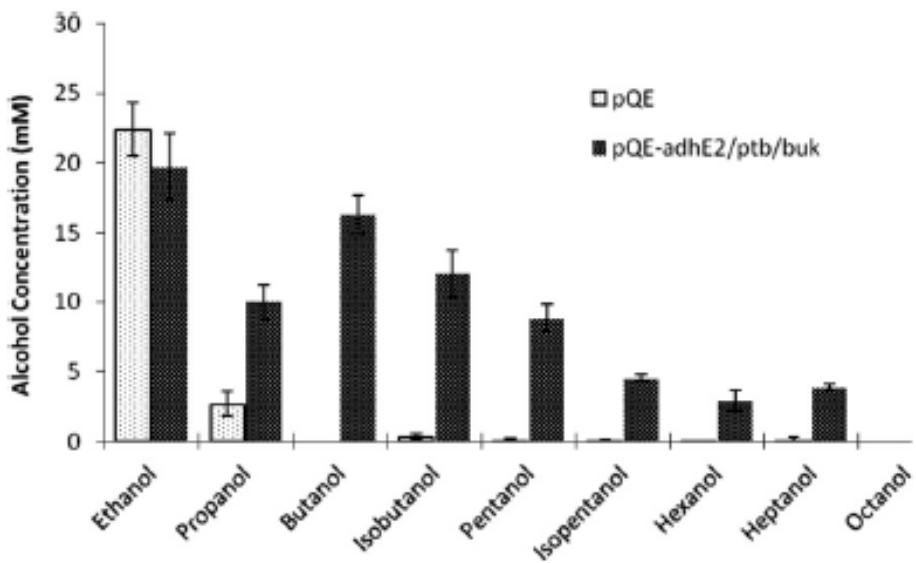
Engineering *E. coli* strain for conversion of short chain fatty acids to bioalcohols

Anu Jose Mattam and Syed Shams Yazdani*

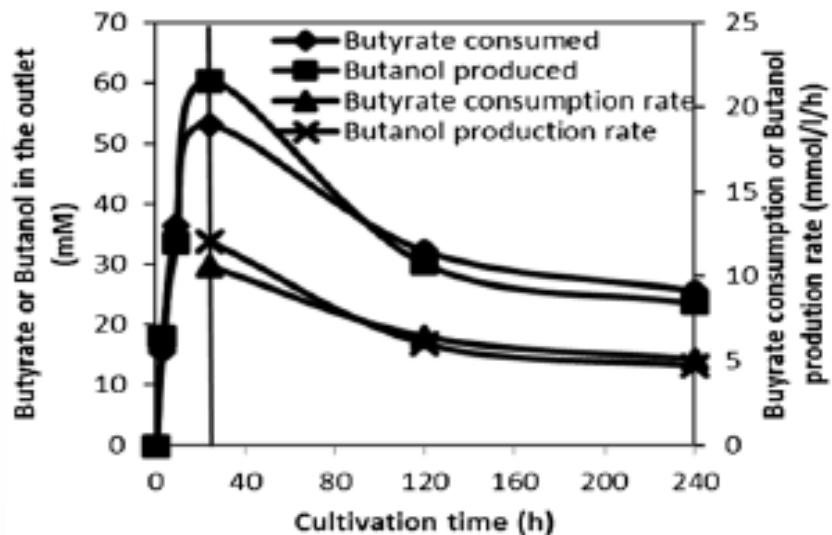


Figure 1 Metabolic pathway of *Clostridium acetobutylicum* engineered in *E. coli*. Abbreviations: Buk – butyrate kinase, PtB – phosphotransbutyrylase, AdhE2 – aldehyde-alcohol dehydrogenase.

Engineering *E. coli* for short chain alcohols



Production of C3-C7 alcohol



Continuous Production of Butanol

Acknowledgment

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 - Nidhi, Research Associate
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 - Anu, Graduate Student
 - Zia, Graduate Student
 - Zeenat, Junior Res Fellow
 - Vamsi, Junior Res Fellow
 - Tabinda, Junior Res Fellow
 - Kamran, Junior Res Fellow
- Collaborators
 - Insect Resistance Group, ICGEB
 - DBT-ICT Centre for Energy Bioscience, Mumbai
 - DBT-IOC Centre for Advanced Bioenergy Research, Faridabad
- Funding
 - DBT, Govt of India
 - DST, Govt of India



Thank You