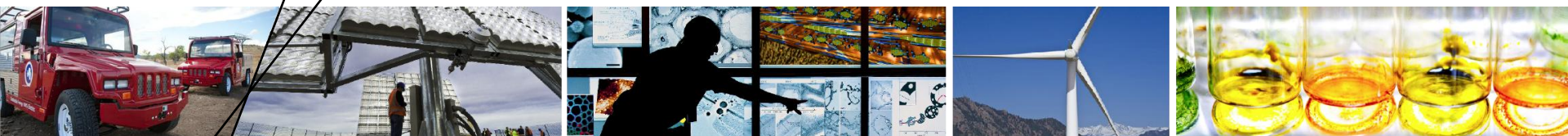


# Enzyme Synergy and Cooperation in Biomass Conversion



***Pacific Rim Summit on Industrial  
Biotechnology & Bioenergy***

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# Problems in Biomass Enzyme Synergy

- **“Real” substrates are highly variable**
  - Different feedstocks, pretreatment chemistries and severities
    - Highly heterogenous
    - Variable particle size, surface area:volume ratios
- **Model substrates are also highly variable**
  - Cellulose
    - different levels of crystalline, paracrystalline, and amorphous content
      - Different crystalline isoforms
    - lignin and hemicellulose residuals
    - degree of polymerization
    - processing changes, etc.
  - “native” hemicelluloses are likely soluble
    - Most available xylans have been de-esterified
    - Hemicellulose varies greatly in structure and chemical composition
- **Complex enzyme systems**
  - Endo/exo/b-glucosidase/PMOs for cellulases
    - Multiple variants of each
  - More than a dozen activities involved in xylan hydrolysis
    - Endos, exos, debranching, accessory enzymes
- **Conflicts in literature**
  - No standardized assay methods, variable substrate composition determination, wide range of enzyme purities and sources
  - ***Synergy in biomass conversion has few hard and fast rules; studies should be evaluated primarily as stand-alone results***





# Classical Cellulase Synergy

## • Endo-Exocellulase synergy

- Hinted at by Elwyn Reese et al. 60+ years ago

- [Reese E. T., Siu R. G., Levinson H. S. \(1950\) The biological degradation of soluble cellulose derivatives and its relationship to the mechanism of cellulose hydrolysis. J. Bacteriol. 59, 485-497](#)

- $C_1/C_x$  model:  $C_1$  releases substrates for  $C_x$

- Classical synergy model- Wood and McCrae

- [T. M. Wood and S. I McCrae. 1979. Synergism between enzymes involved in the solubilization of native cellulose. Advances in chemistry. 181:181-209.](#)

- Endos ( $C_x$ ) create new chain ends for exos
    - » # chain ends is rate-limiting

- Exos ( $C_1$ ) disrupt crystalline regions, exposing more sites for endos

- Recently revisited by Jürgen Jalak et al.

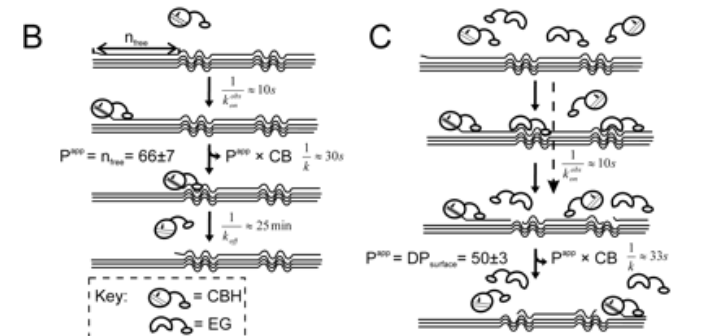
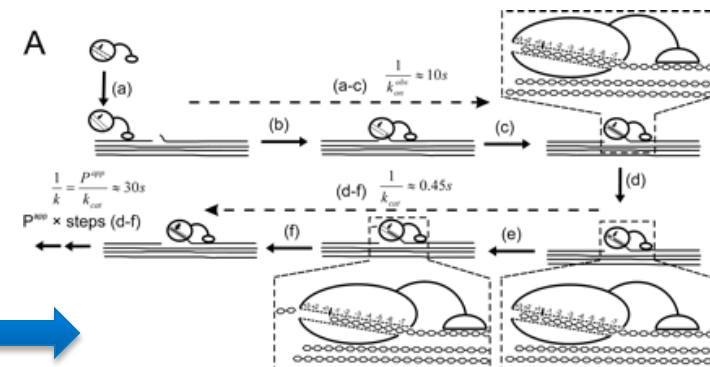
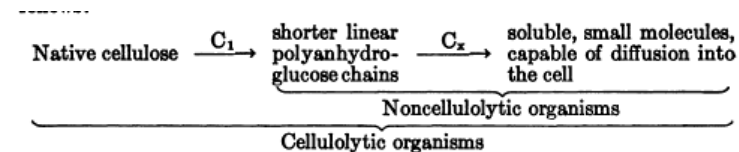
- [J. Jalak, M. Kurasin, H. Teugjas, P. Välijamäe. 2012. Endo-exo Synergism in Cellulose Hydrolysis Revisited. J Biol Chem. 17:287:28802-15](#)

- Endocellulases increase exocellulase average processivity

- $k_{off}$  for cel7A is rate-limiting

- Removes amorphous regions which stall exos

- Remove single chains from surface



# Exo-Exo Cellulase Synergy

- **Different exocellulases attack different ends**

- Proposed by Fagerstam and Pettersson in 1980

- [Fagerstam, L. and L. Pettersson. 1980. The 1,4-b-glucan cellobiohydrolases of \*Trichoderma reesei\* QM9414. An new type of cellulolytic synergism. FEBS Lett. 119\(1\), 97-100.](#)

- Demonstrated by Barr *et al* in 1996

- [Barr, B. K., Y. L. Hsieh, B. Ganem, D. B. Wilson. 1996. Identification of two functionally different classes of exocellulases. Biochem. 35\(2\), 586-592](#)

- Reducing vs. non-reducing ends

- GH7s are reducing end specific, GH6s are non-reducing end specific

- Insight from ultra-high speed AFM

- [Igarashi, K., T. Uchihashi, A. Koivula, M. Wada, S. Kimura, T. Okamoto, M. Penttilä, T. Ando, and M. Samejima. 2011. Traffic Jams Reduce Hydrolytic Efficiency of Cellulase on Cellulose Surface. Science 333 \(6047\) pp. 1279-1282](#)

- Demonstrated cel7A is processive and peels layers

- Gets stuck and stalls on surface

- Cel6A frees stuck cel7A

- Supports Jalak *et al* model

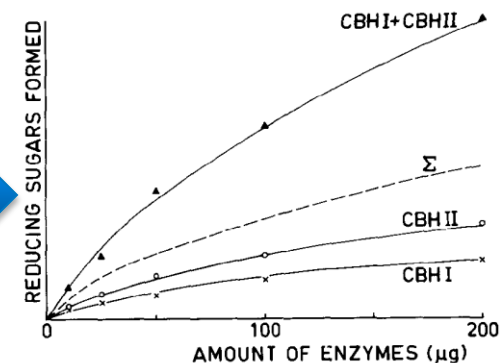
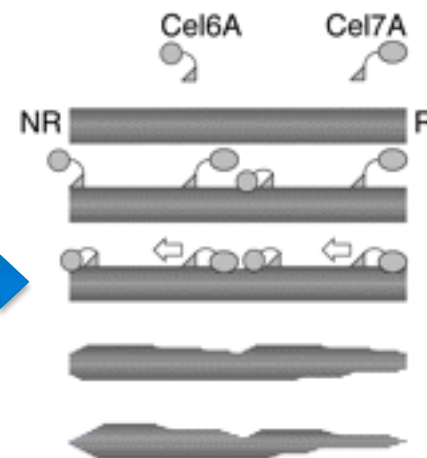
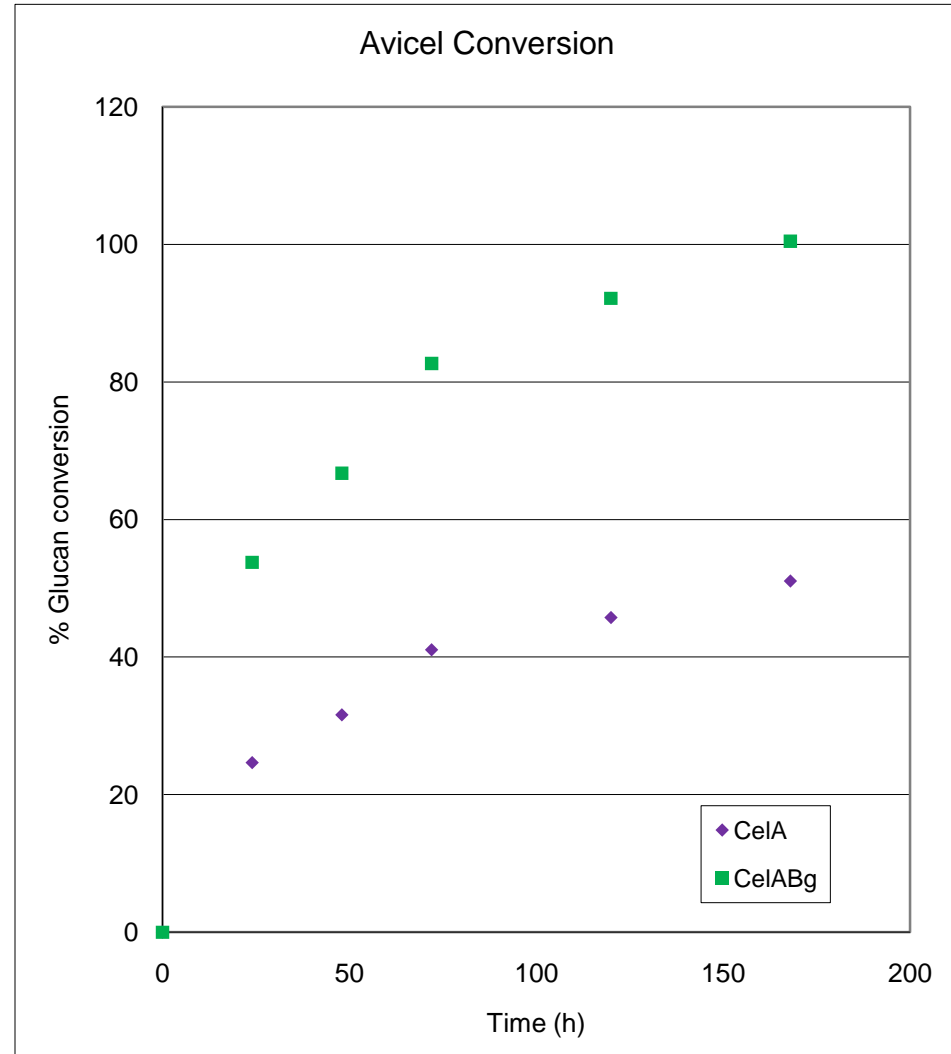


Fig.2. Synergism towards crystalline cellulose. CBH I and CBH II were incubated alone and in combination with 1% Avicel as substrate. The samples were incubated for 1 h at 40° C in 50 mM sodium acetate buffer (pH 5.0). The non-synergistic sum of the activities of CBH I and CBH II is indicated by  $\Sigma$ .



# Cellulase- $\beta$ -glucosidase Synergy

- **Generally thought to be due to end-product (cellobiose) inhibition of cellulase**
  - Usually cel7A or bacterial analog
- **Typical experiment shows increased digestion**
  - CelA from *Caldicellulosiruptor bescii*
    - 15 mg/g cellulose
  - $\beta$ -G from *Thermotoga maritima*
    - 1 mg/g cellulose
  - Similar to many other cellulase/ $\beta$ -G combinations
  - 1.5% avicel



# A New Cellulose Hydrolysis Mechanism

Biochemistry 2010, 49, 3305–3316

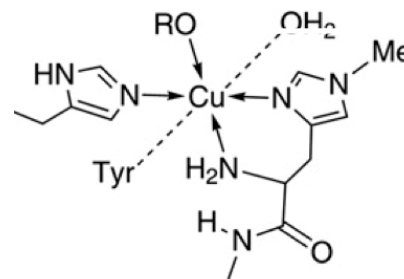
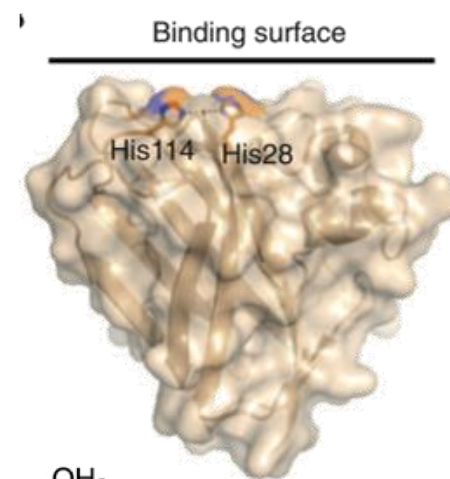
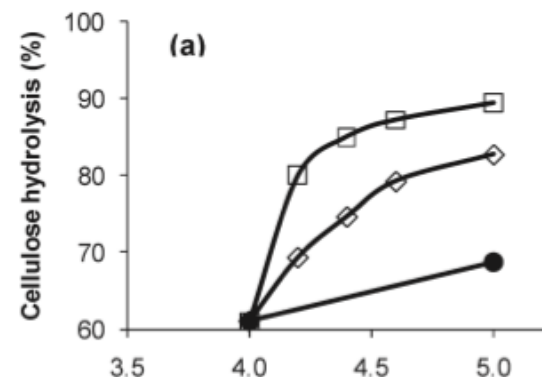
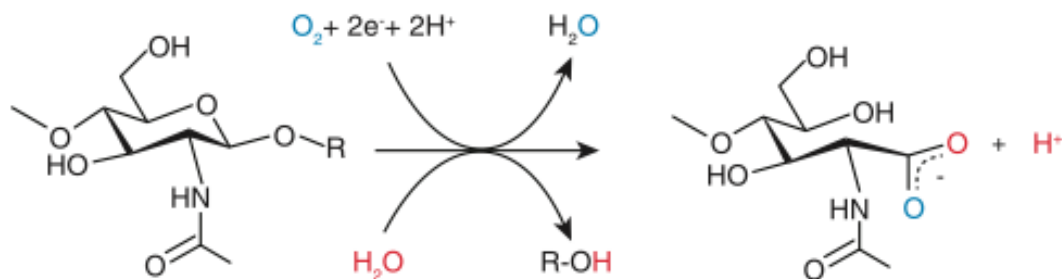
Stimulation of Lignocellulosic Biomass Hydrolysis by Proteins of Glycoside Hydrolase Family 61: Structure and Function of a Large, Enigmatic Family<sup>†</sup>

Paul V. Harris,<sup>\*,†,‡,§</sup> Ditte Welner,<sup>§,||</sup> K. C. McFarland,<sup>‡</sup> Edward Re,<sup>‡</sup> Jens-Christian Navarro Poulsen,<sup>§</sup> Kimberly Brown,<sup>‡</sup> Rune Salbo,<sup>§</sup> Hanshu Ding,<sup>‡</sup> Elena Vlasenko,<sup>‡</sup> Sandy Merino,<sup>‡</sup> Feng Xu,<sup>‡</sup> Joel Cherry,<sup>‡</sup> Sine Larsen,<sup>§</sup> and Leila Lo Leggio<sup>§</sup>

7 months later:

## An Oxidative Enzyme Boosting the Enzymatic Conversion of Recalcitrant Polysaccharides

Gustav Vaaje-Kolstad, Bjørge Westereng, Svein J. Horn, Zhanliang Liu, Hong Zhai, Morten Sørlie, Vincent G. H. Eijsink<sup>\*</sup>



10 months later:

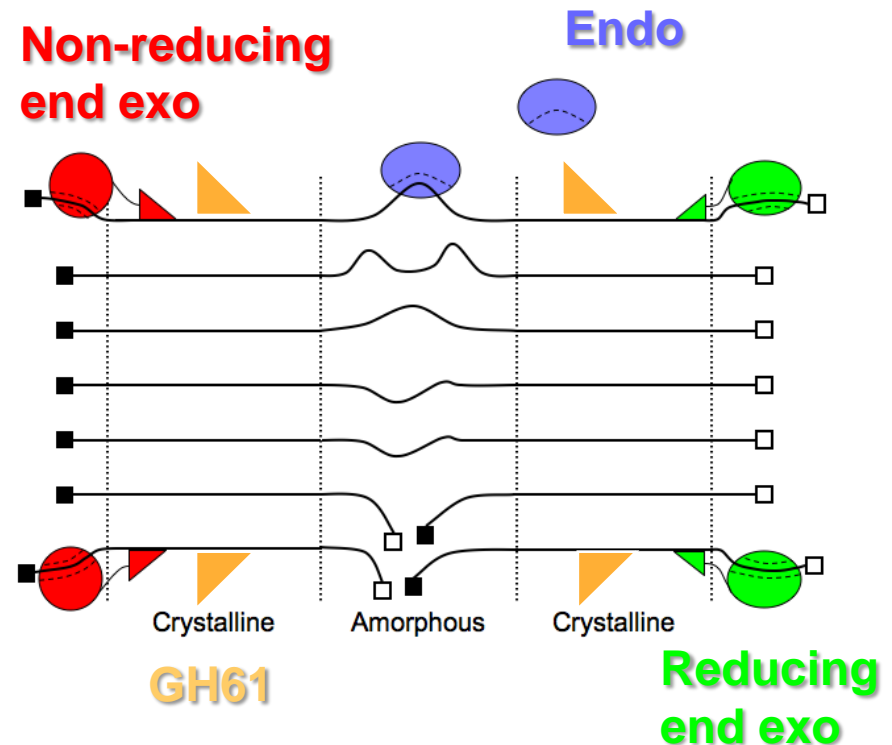
## Insights into the oxidative degradation of cellulose by a **copper** metalloenzyme that exploits biomass components

R. Jason Quinlan<sup>a,1</sup>, Matt D. Sweeney<sup>a,1</sup>, Leila Lo Leggio<sup>b</sup>, Harm Otten<sup>b</sup>, Jens-Christian N. Poulsen<sup>b</sup>, Katja Salomon Johansen<sup>c,2</sup>, Kristian B. R. M. Krogh<sup>c</sup>, Christian Isak Jørgensen<sup>c</sup>, Morten Tovborg<sup>c</sup>, Annika AnthonSEN<sup>c</sup>, Theodora Tryfona<sup>d</sup>, Clive P. Walter<sup>c</sup>, Paul Dupree<sup>d</sup>, Feng Xu<sup>a</sup>, Gideon J. Davies<sup>e</sup>, and Paul H. Walton<sup>e</sup>



# Where does GH61 fit in Cellulose Hydrolysis?

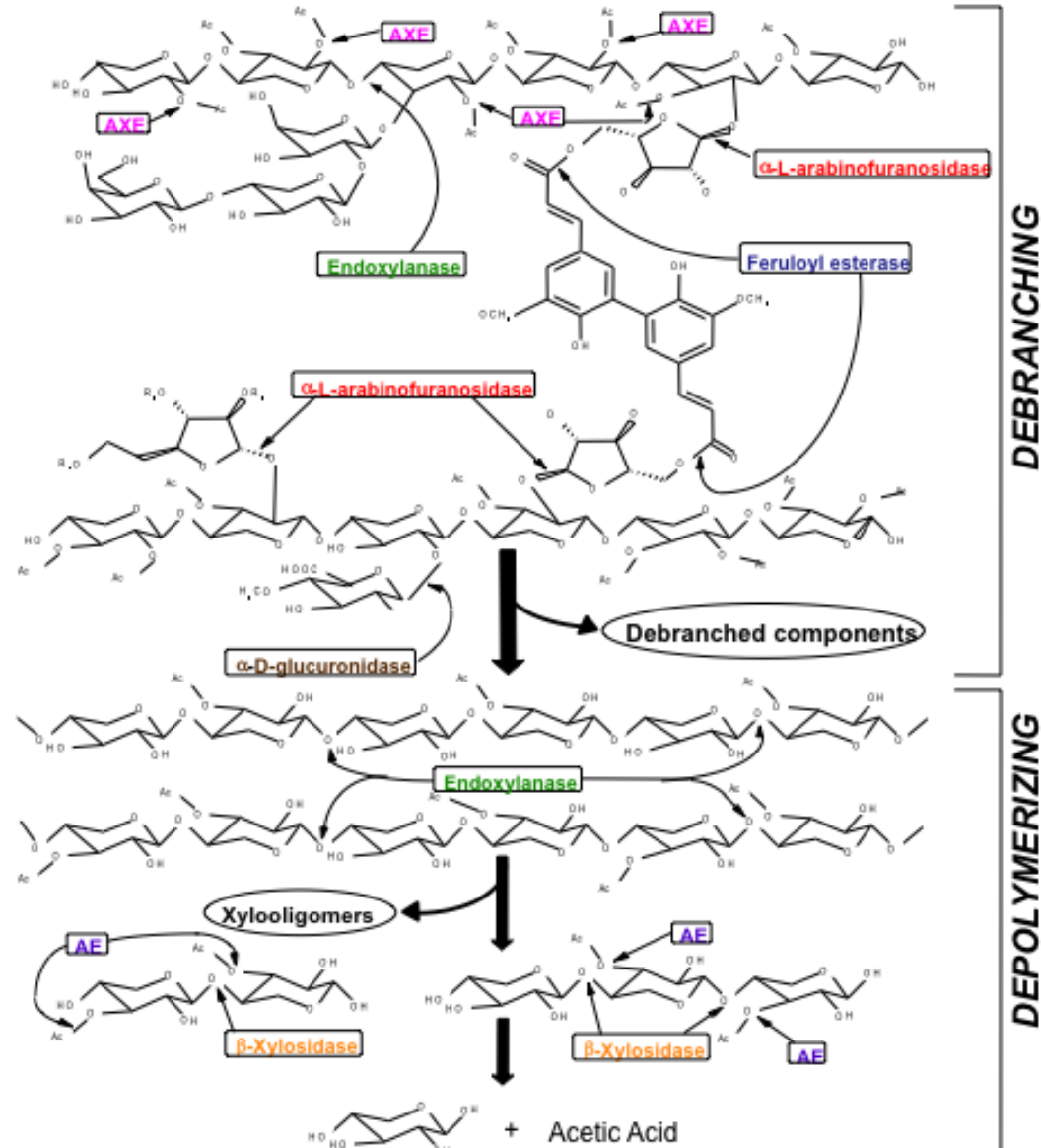
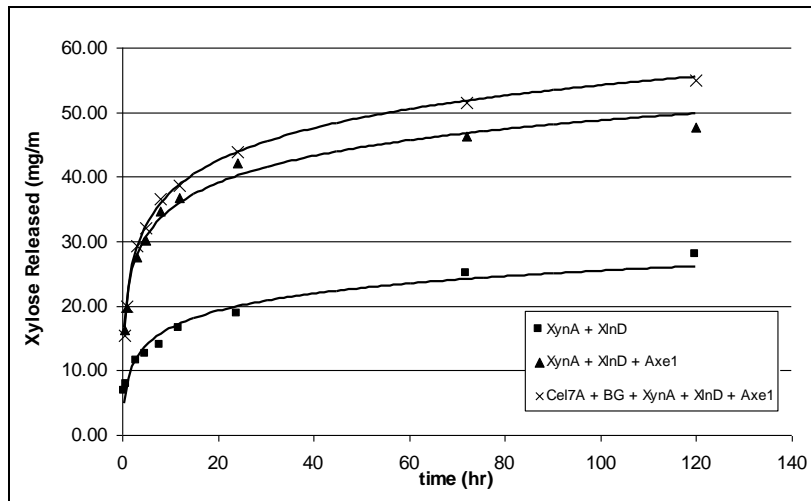
- **GH61 likely acts to remove cellulose surface aberrations**
  - Analogues to endos or cel6A
  - May initiate new binding sites for exos
- **Many questions remain including:**
  - What is the mechanism of GH61 action?
  - What residues are important for specificity?
  - What other substrates can GH61s act on?
  - How do GH61s behave on cellulose?
  - What is the structural diversity of GH61s?





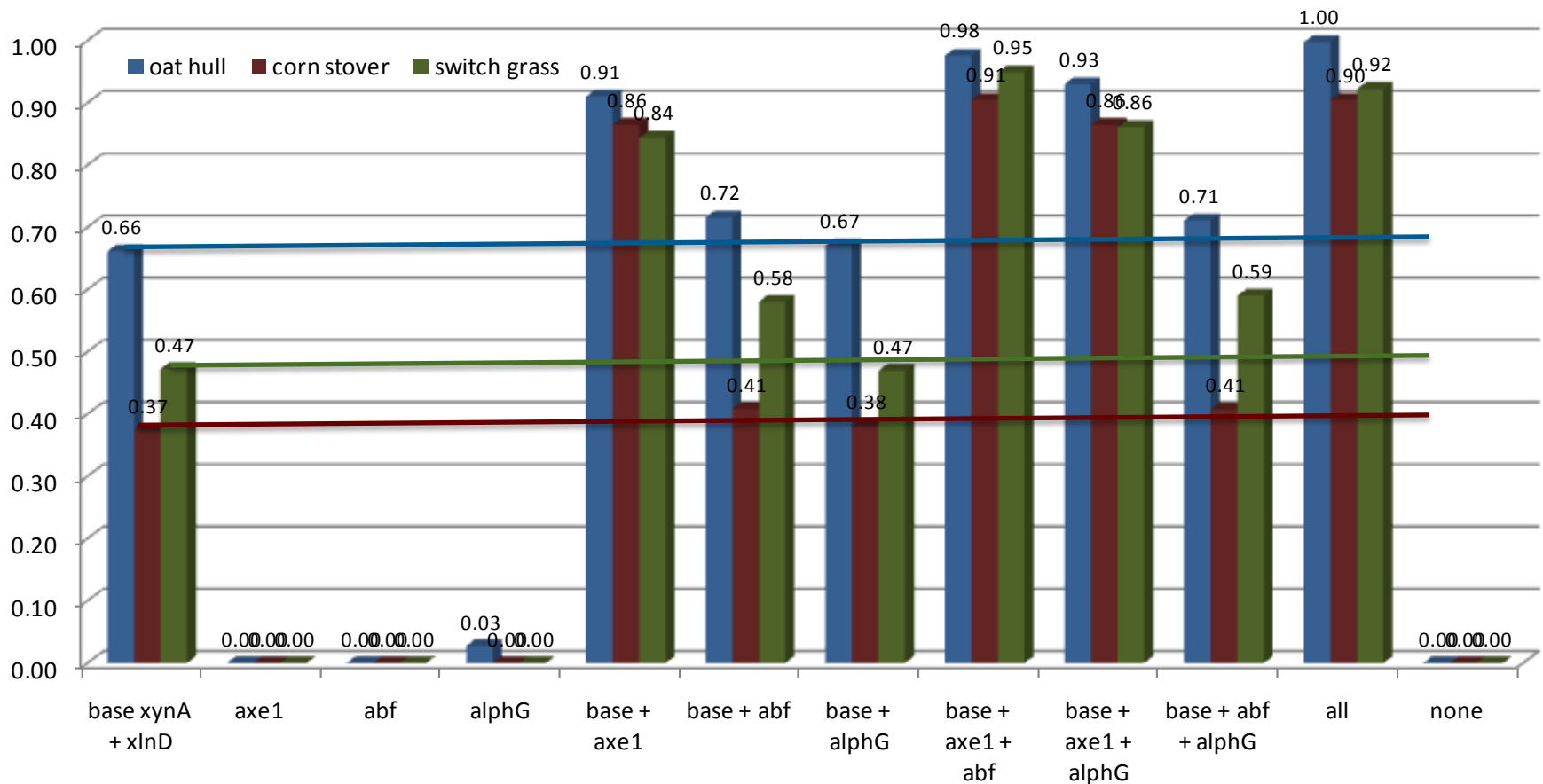
# Synergy in Hemicellulose Hydrolysis

- **Hemicellulose complexity requires a variety of enzyme classes**
  - Hardwoods vs. softwoods vs. herbaceous
- **Native hemicellulose is thought to be “soluble”**
  - Extracted with ester-links intact, it is
  - Side chains prevent intermolecular association and retain water
  - Debranching leads to insoluble backbone
    - Slower enzyme binding kinetics
- **Not really synergy, as enzymes release different products**
  - Cooperativity is closer



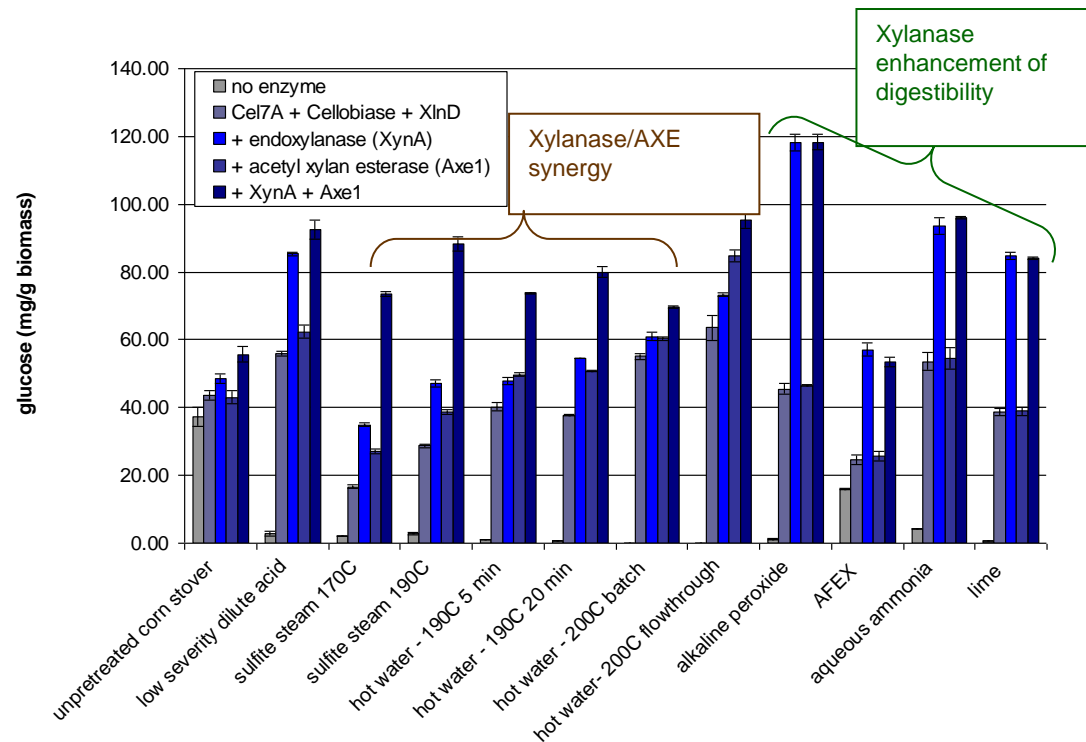
# Digestion of Natively Extracted (DMSO) Xylans

- **Cooperativity of accessory enzymes mixes on purified xylans**
  - xynA/xInD + abfB, glrA, axe1



# Enzyme Synergy in Pretreatment Solids

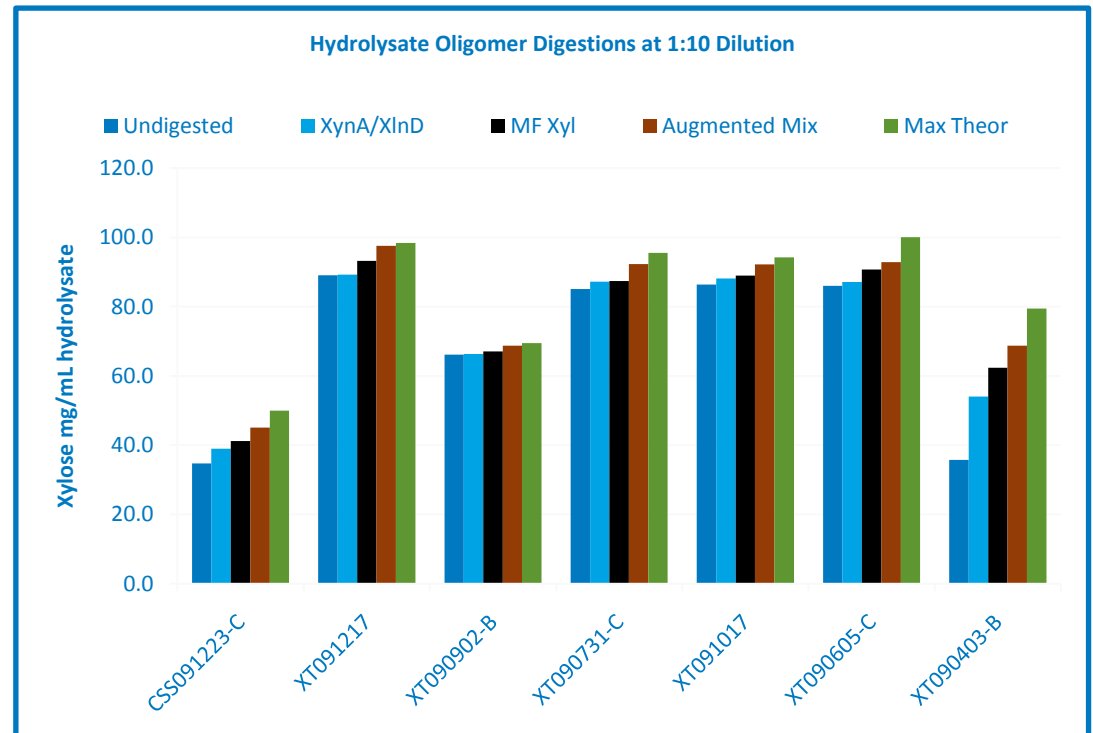
- **Xylanase enhances glucose release**
  - Xylanase/AXE synergy on SSE/low severity HW
  - No AXE synergy- alkaline de-esterification
- **Cellulase enhances xylose release**
  - Sulfite steam exploded corn stover-low severity
- **Improvements in xylan and glucan conversion continue over long digestions**
  - Synergism improves rates
  - Synergism improves extent of conversion
  - Improved accessibility to target structures
- **Can lower enzyme loading for given severity**
  - DA-180°C, 40 → 15 mg prt/g cellulose



Selig, M. J., W. S. Adney, M. E. Himmel and S. R. Decker. (2009). [The Impact of Cell Wall Acetylation on Corn Stover Hydrolysis by Cellulolytic and Xylanolytic Enzymes](#). Cellulose. 16:711-722

# Enzyme Synergy on Hydrolysate Oligomers

- **End product inhibition is significant**
  - enzymes are not capable of digesting all of the oligomers and solids to completion in SSF.
- **Commercial xylanases lack critical activity(s)**
  - $\beta$ -xylosidase, AXE,  $\beta$ -glucuronidase, and/or  $\beta$ -L-arabinofuranosidase.
  - Even high-xylose yield pretreatments can benefit from additional enzyme activities
- **Endoxylanase and  $\beta$ -xylosidase can be as effective as a commercial enzyme prep.**
  - Some xylanases are deficient in debranching/accessory enzymes

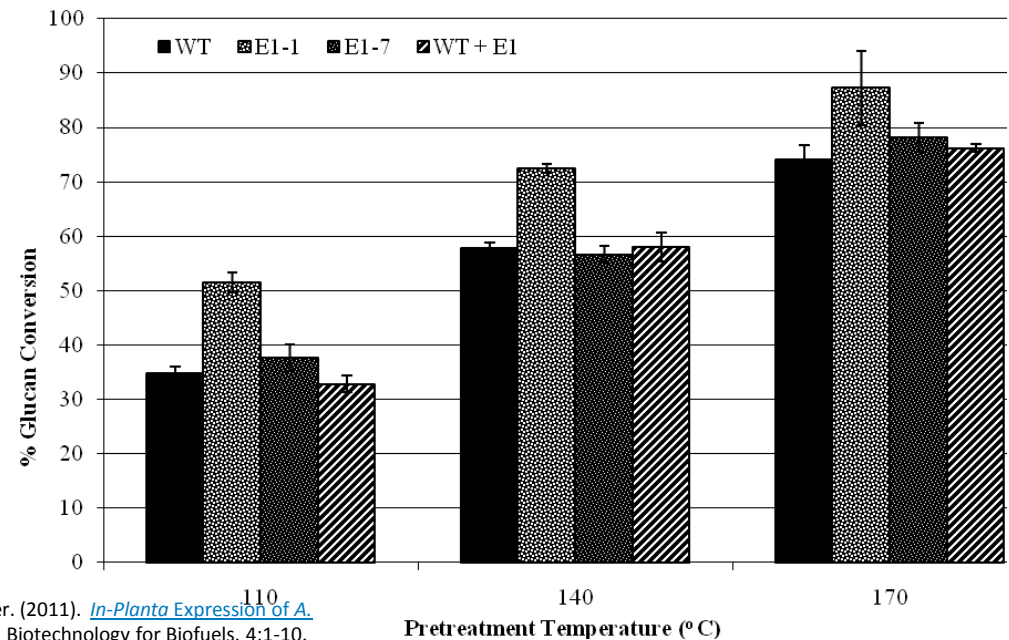
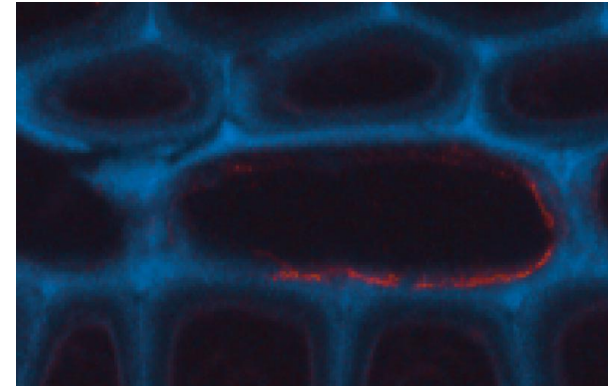


Digestions of hydrolysate oligomers at 1:10 dilution with commercial, component, and augmented commercial xylanases. XynA/XInD was a mix of XynA (10 mg per g oligomer) and XylD (1 mg per g oligomer). MFXyl was Multifect Xylanase loaded at 10 mg per g oligomer. The Augmented Mix The augmented mix consisted of MF Xylanase (20 mg/g oligomer), AXE (5 mg/g), abf (5 mg/g),  $\alpha$ -glucuronidase (2.5 mg/g), and  $\beta$ -xylosidase (1 mg/g). Conversion is corrected to undiluted sample.

# “Other” Enzyme Enhancements

- **Plant expression of cellulase**

- Old idea
  - enzyme production
  - Post-harvest digestion
- New idea
  - Specifically weaken cell wall to pretreatment
  - Lower pretreatment severity and/or decrease enzyme loading
- Expressed cel5A in corn
  - Pretreated with dilute acid at several temperatures
  - E1-stover was consistently more digestible than WT



Brunecky, R., M. J. Selig, T. B. Vinzant, M. E. Himmel, D. Lee, M. J. Blaylock and S. R. Decker. (2011). [In-Planta Expression of \*A. cellulolyticus\* Cel5A Endocellulase Reduces Cell Wall Recalcitrance in Tobacco and Maize](#). *Biotechnology for Biofuels*. 4:1-10.

# Summary

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- **Endo-exo and exo-exo cellulase synergies are likely due to prevention of cel7A stalling**
  - GH61 may act similarly
- **Cellulase -  $\beta$ -glucosidase synergy is likely due to end-product inhibition relief**
  - Similar effect in xylanase -  $\beta$ -xylosidase
- **Hemicellulase – accessory enzyme synergy is in part a function of maintaining backbone chain solubility**
  - Debranching can lead to insoluble chains
  - Synergy in oligomer conversion can enhance yield
- **Free cellulases and cellulosomes are synergistic**
  - Only on cellulose, not biomass
  - Mechanism is not understood
- ***Enzyme synergy in biomass conversion is complicated***

# Acknowledgements

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- **Mike Himmel- our fearless leader**
- **John Baker- enzyme kinetics**
- **Mike Selig- hemicellulase assayer, temporary Canadian**
- **Mike Resch- cellulosome and GH61 biochemistry**
- **Bryon Donohoe- TEM imaging**
- **Todd Vinzant- pretreatment**
- **Gregg Beckham- GH61 research coordinator**
- **Roman Brunecky- bacterial enzyme kinetics**