

## A novel xyloglucanase from the white rot fungus *Abortiporus biennis* and its potential role as an accessory biocatalyst in the enzymatic degradation of xyloglucan-containing substrates

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### INTRODUCTION & AIM

Lignocellulosic biomass is a composite material consisting of cellulose, hemicellulose, and lignin. Usually, cellulose fibrils are covered with **xyloglucan**, a complex, highly substituted plant biomass hemicellulose. Xyloglucan is present in many plant species as a seed storage polysaccharide or as a component of the primary cell wall.

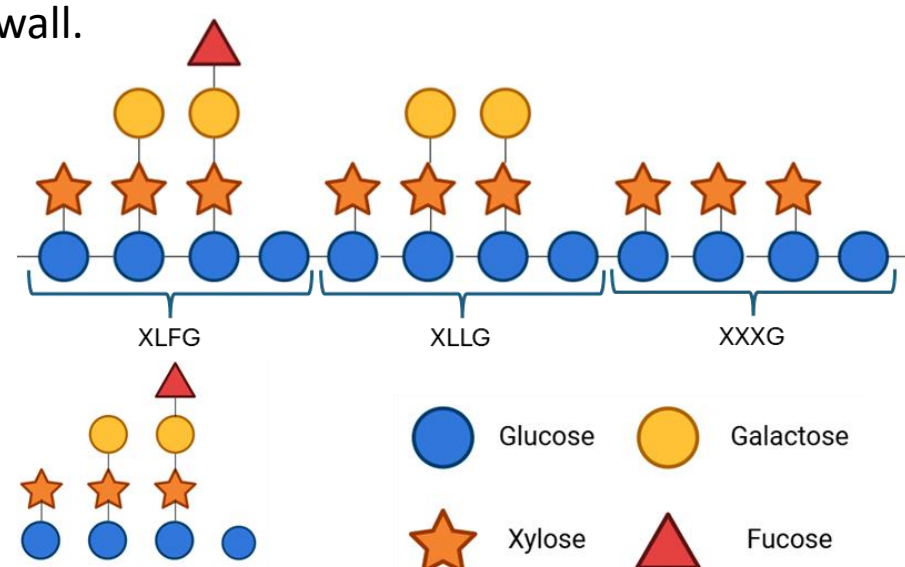


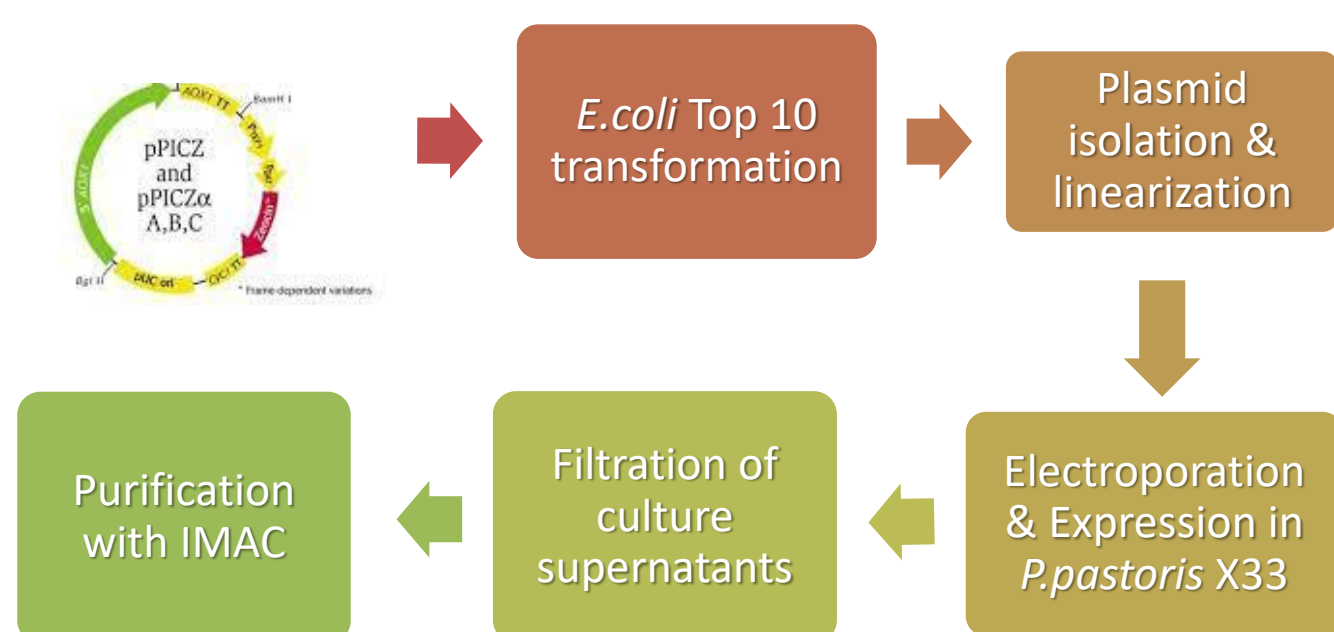
Figure 1:  
Primary  
structure of  
xyloglucan

Xyloglucan requires many enzymes with complementary specificities for its complete breakdown called **xyloglucanases** (XEGs, EC 3.2.1.151). They hydrolyze backbone  $\beta$ -1,4-glucosidic bonds, releasing shorter oligomers. In this study, we characterized **AbiXega**, a novel XEG from the white-rot fungus *Abortiporus biennis*, known for its lignocellulose-degrading capabilities.

The need for xyloglucan removal is evident in biorefinery applications, where platform sugars must be obtained for the valorization of cellulosic biomass, but also in the improvement of animal feed, such as corn bran and apple pulp.

### METHOD

#### Expression and purification of *AbiXega*



- Bioinformatics analysis
- Biochemical characterization
- Mode of action studies using DNS method to detect reducing sugars
- Synergism with commercial cellulases

### RESULTS & DISCUSSION

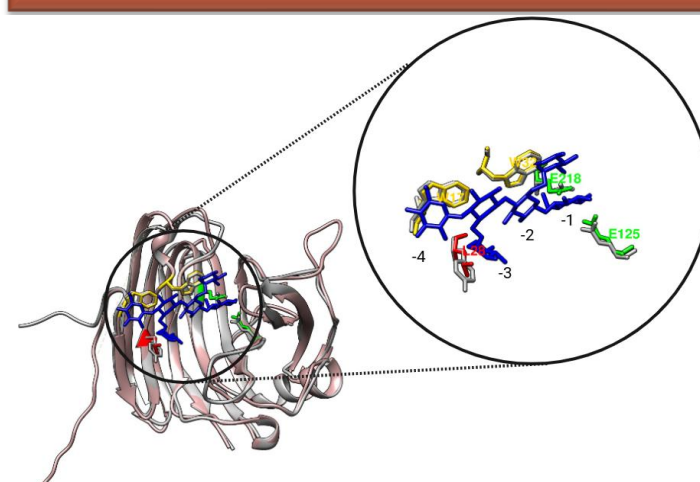


Figure 2: Superimposition of the AlphaFold model structure of *AbiXega* (pink) with *Aspergillus aculeatus* XEG (grey) and close-up view of the ligand binding site. (Ligand GXXG: blue, catalytic glutamates of *AbiXega* green, Leu28: red, Trp17 and Trp32: yellow)

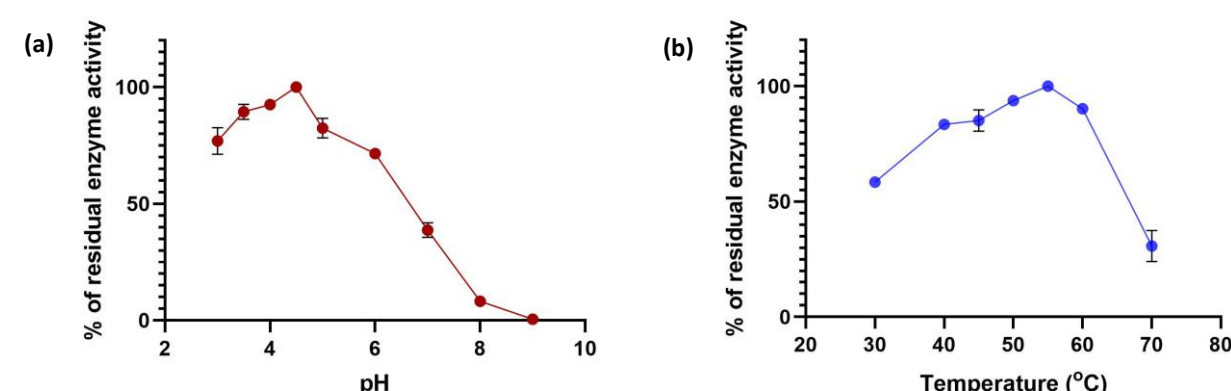


Figure 3: Optimum activity conditions of *AbiXega*. Effect of (a) pH and (b) temperature on the activity of *AbiXega* during xyloglucan hydrolysis.

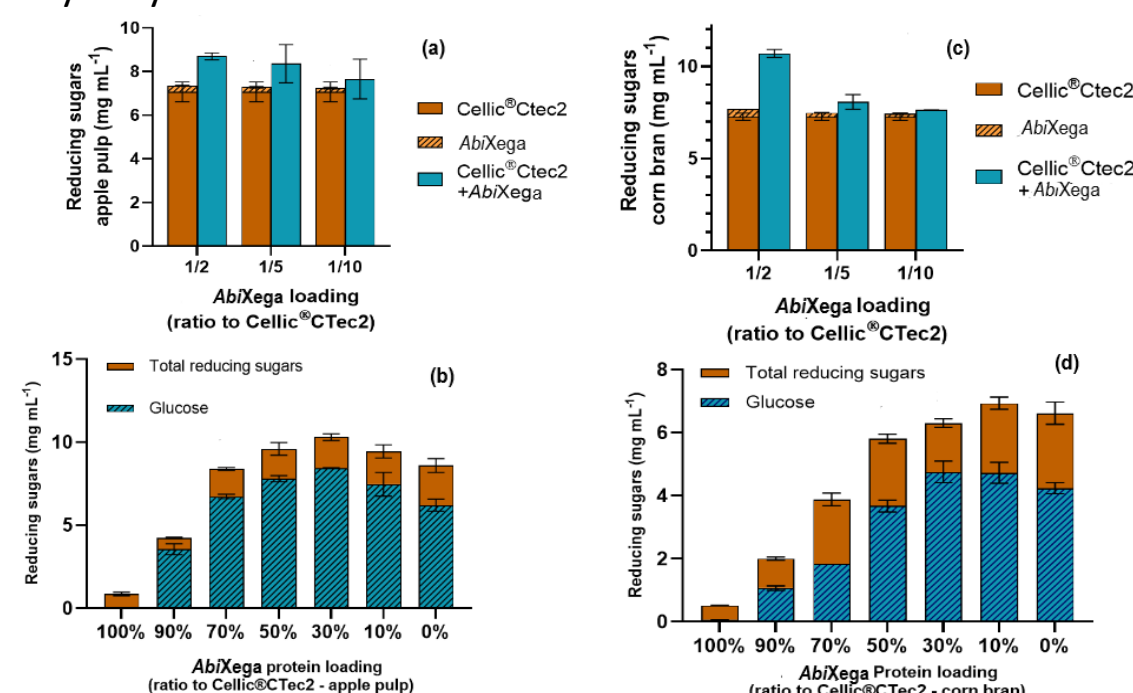


Figure 4: Synergism of *AbiXega* and cellulases in apple pulp (a, b) and corn bran (c, d) hydrolysis. Effect of *AbiXega* protein loading on the reducing sugars release from (a) apple pulp and (c) corn bran. (b), (d) Effect of *AbiXega* dosage on a total protein content of fixed protein loading on the production of reducing sugars and glucose by cellulases.

### CONCLUSION

- Detectable activity in xyloglucan and  $\beta$ -glucan
- Enhances the action of cellulases in corn bran and apple pulp
- Reaction's total protein loading minimized, without loss in reducing sugars release → lower enzyme costs for substrates with a high xyloglucan content

### REFERENCES

- Matsuzawa, T., Watanabe, A., Shintani, T. et al. Enzymatic degradation of xyloglucans by *Aspergillus* species: a comparative view of this genus. *Appl Microbiol Biotechnol* 105, 2701–2711 (2021). <https://doi.org/10.1007/s00253-021-11236-8>
- Sun, P., Li, X., Dilokpimol, A., Henrissat, B., de Vries, R. P., Kabel, M. A., & Mäkelä, M. R. (2022). Fungal glycoside hydrolase family 44 xyloglucanases are restricted to the phylum Basidiomycota and show a distinct xyloglucan cleavage pattern. *IScience*, 25(1), 103666. <https://doi.org/10.1016/j.isci.2021.103666>