Supplementary data for

Functional Study of Different Lignocellulases from *Trichoderma guizhouence* NJAU4742 in the Synergistic Degradation of Natural Straw

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Number of pages: 7

Number of figures: 6

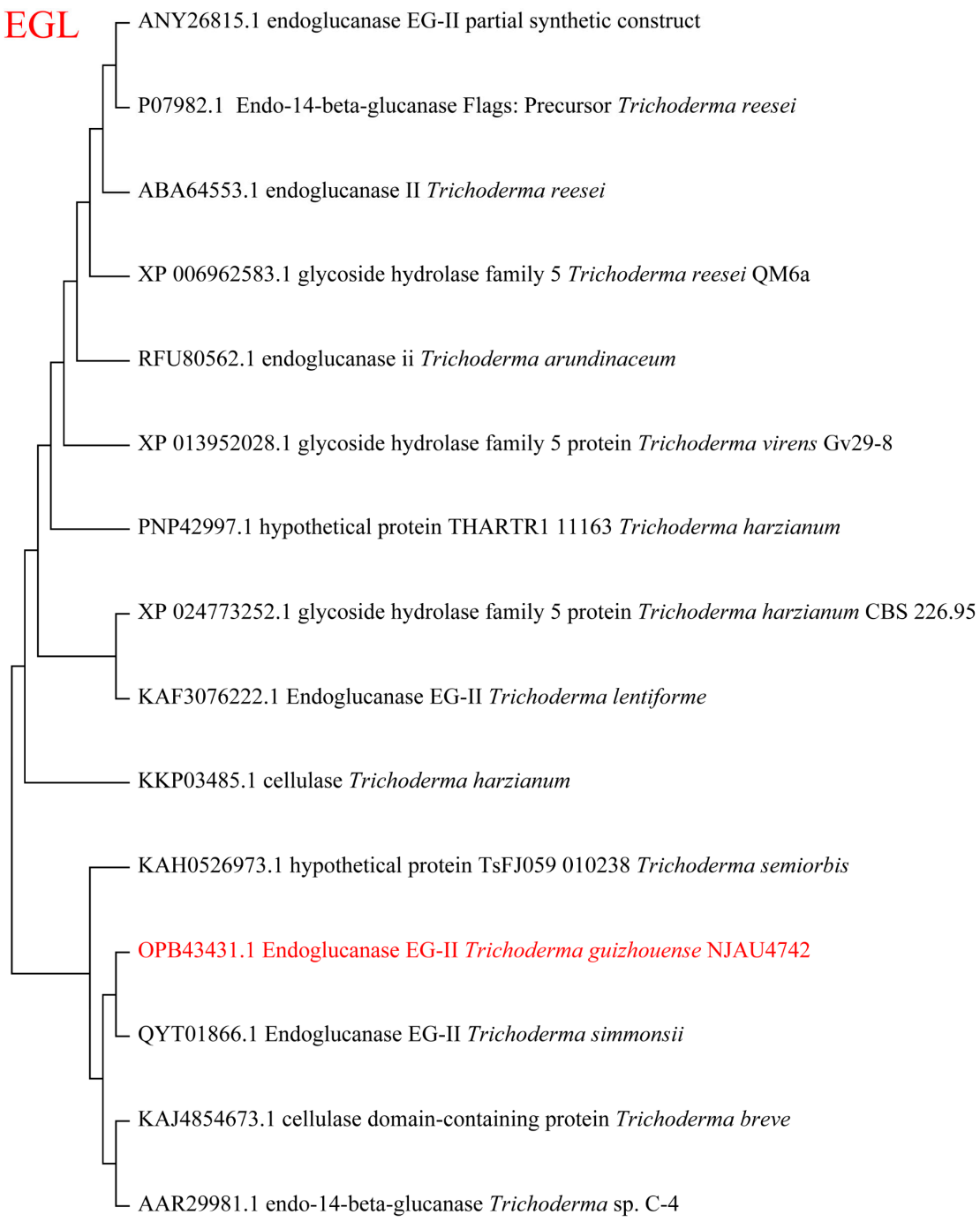

Figure S1. The phylogenetic trees of EGL by performing multiple sequence alignments using MEGAX software.

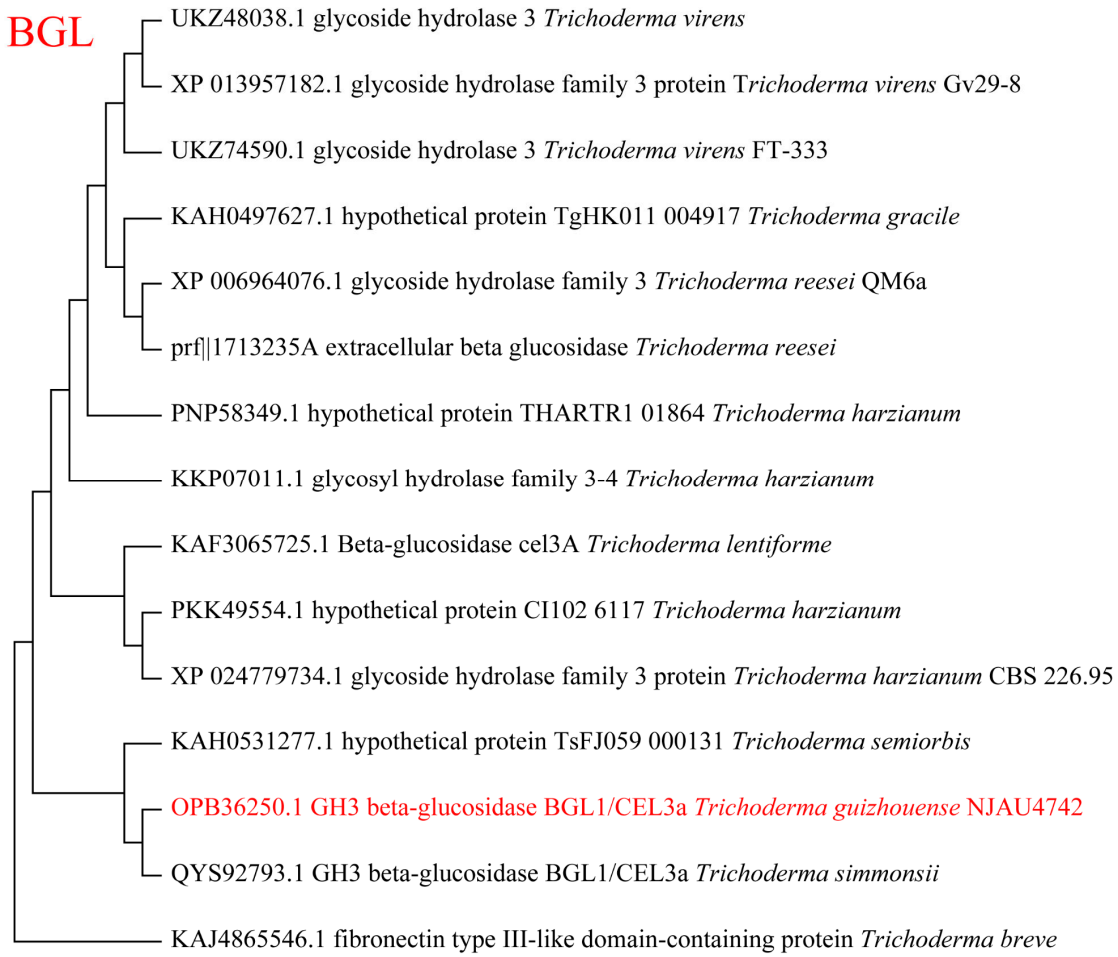

Figure S2. The phylogenetic trees of BGL by performing multiple sequence alignments using MEGAX software.

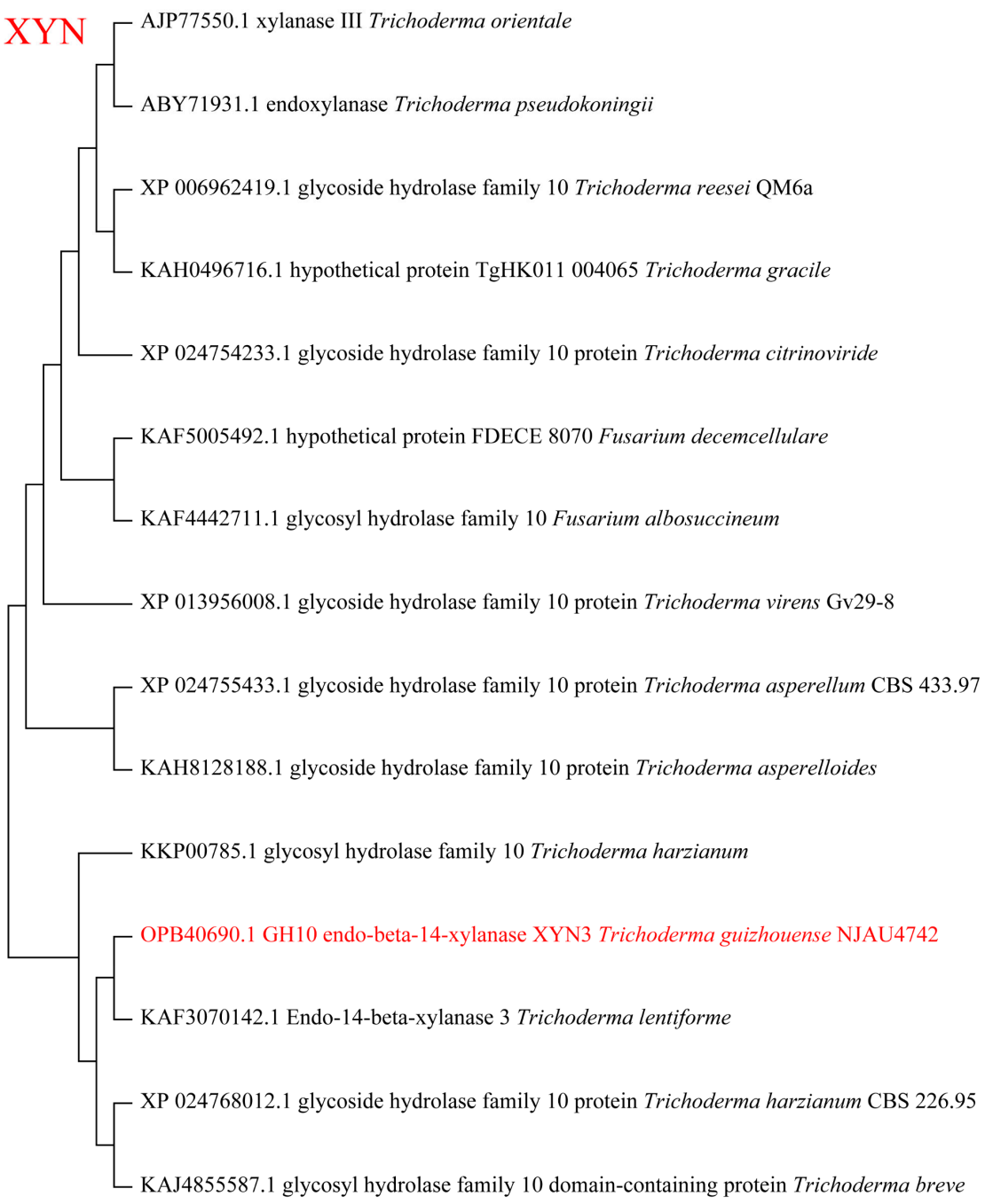

Figure S3. The phylogenetic trees of XYN by performing multiple sequence alignments using MEGAX software.

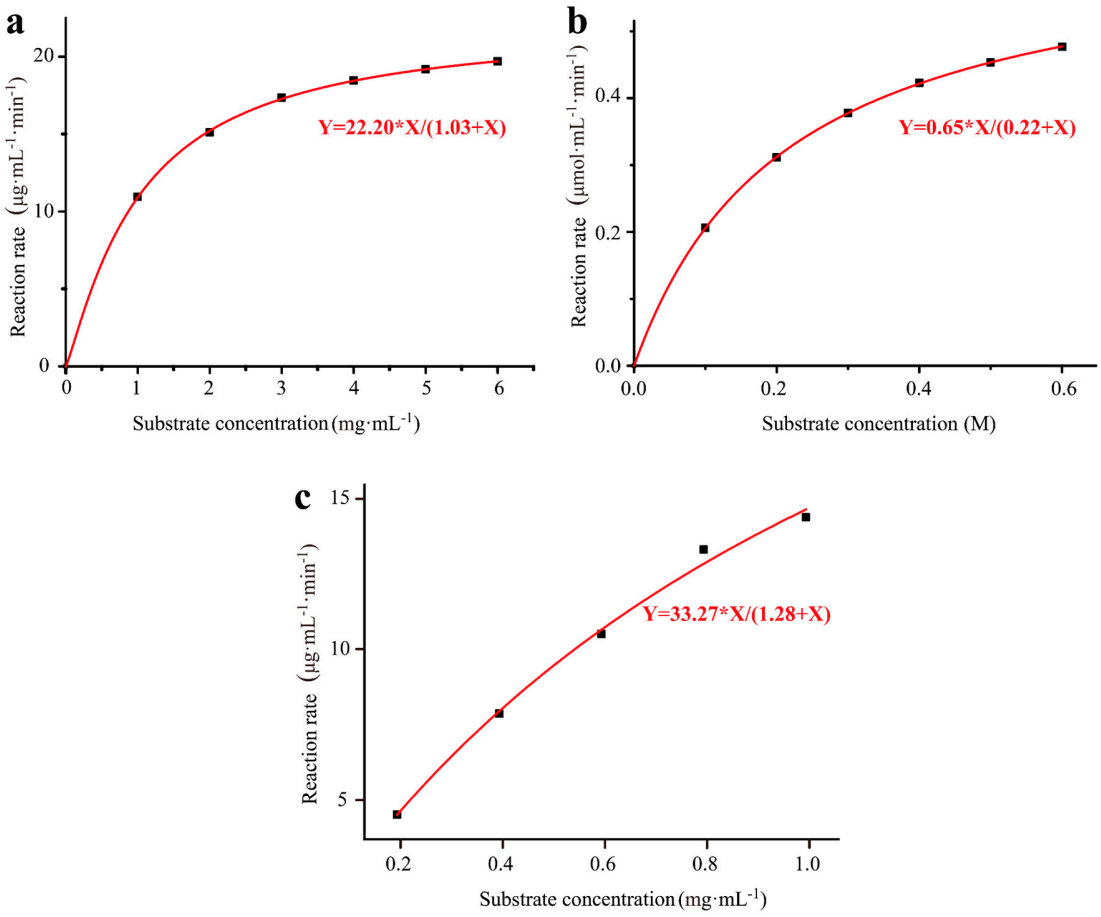

Figure S4. Kinetic analysis of EGL (a), BLG (b), and XYN (c), respectively.

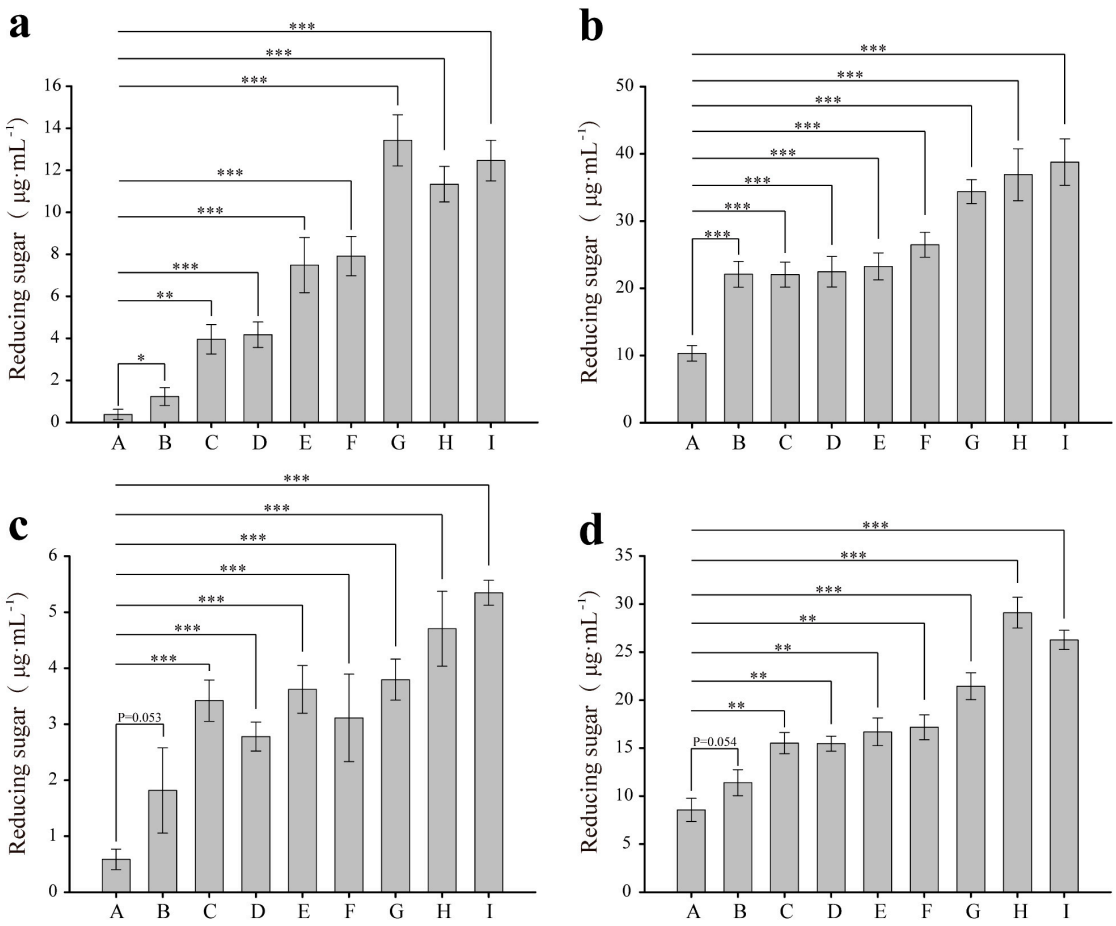

Figure S5. Synergistic degradation of different natural substrates by EGL and BGL. a: wheat straw; b: soybean straw; c: rice straw; d: corn straw. Different letters on the x-axis represent different proportions of the three hydrolytic enzymes added. A: 0.05 M EGL; B: 0.05 M EGL + 0.01 M BGL; C: 0.05 M EGL + 0.02 M BGL; D: 0.05 M EGL + 0.03 M BGL; E: 0.05 M EGL + 0.04 M BGL; F: 0.05 M EGL + 0.05 M BGL; G: 0.05 M EGL + 0.06 M BGL; H: 0.05 M EGL + 0.07 M BGL; I: 0.05 M EGL + 0.08 M BGL. *** $P < 0.001$, ** $P < 0.01$, * $P < 0.05$.

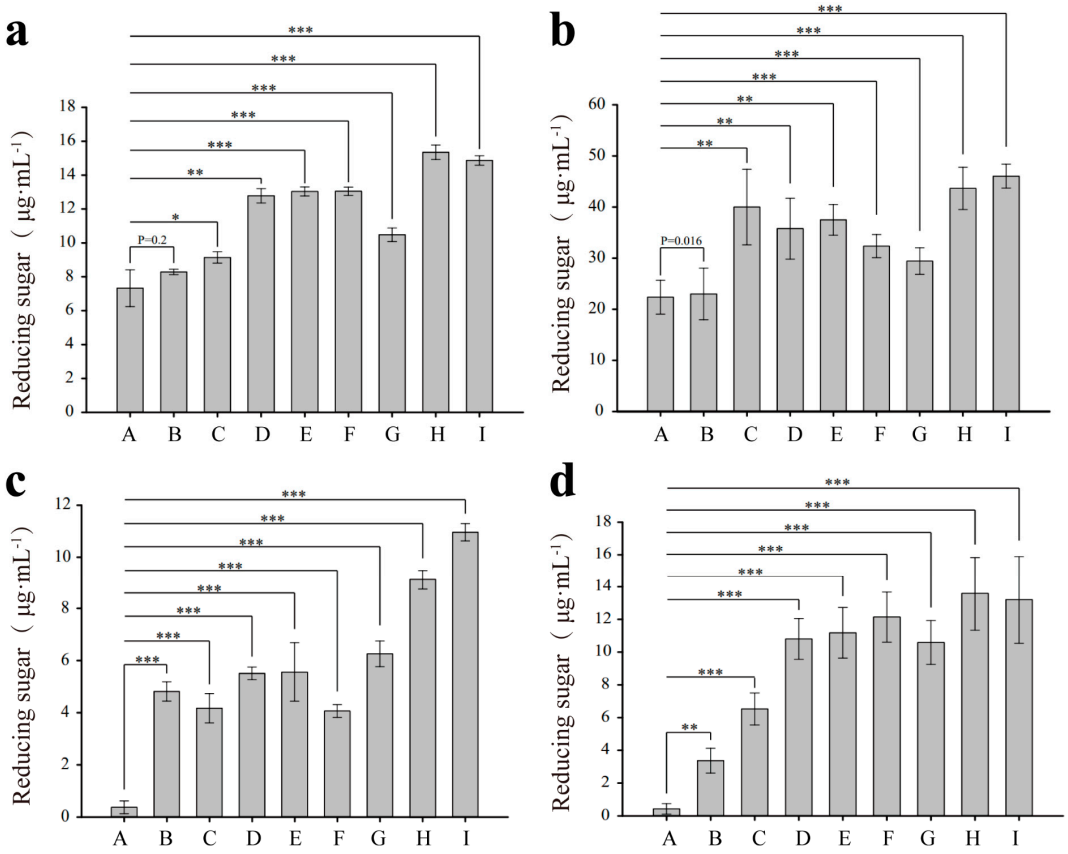

Figure S6. Synergistic degradation of different natural substrates by EGL and XYN. a: wheat straw; b: soybean straw; c: rice straw; d: corn straw. Different letters on the x-axis represent different proportions of the three hydrolytic enzymes added. A: 0.1 M XYN; B: 0.1 M XYN + 0.01 M EGL; C: 0.1 M XYN + 0.02 M EGL; D: 0.1 M XYN + 0.03 M EGL; E: 0.1 M XYN + 0.04 M EGL; F: 0.1 M XYN + 0.05 M EGL; G: 0.1 M XYN + 0.06 M EGL; H: 0.1 M XYN + 0.07 M EGL; I: 0.1 M XYN + 0.08 M EGL. *** $P < 0.001$, ** $P < 0.01$, * $P < 0.05$.