

Microbiomes that metabolize lignin fragments obtained by chemical pretreatment of biomass

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Project Goals: Establish the scientific knowledge and new technologies to transform the maximum amount of carbon available in bioenergy crops into biofuels and bioproducts.

Abstract

Overcoming the recalcitrance of lignin and developing conversion strategies for aromatics are key goals to maximize conversion of carbon in plant biomass. While monoaromatics are metabolized by a variety of bacteria, the metabolism of aromatic oligomers remains largely undescribed. Several Sphingomonad are known to depolymerize dimeric lignin model compounds. Beyond the dimeric model compounds, using HMW lignin directly extracted from lignocellulosic biomass would be beneficial to explore diverse lignin-degrading microbes and their enzymes. Here we generated aromatic mixtures of monomers and oligomers from IL pretreatment (referred to BCD liquor) then enriched microbiomes originated from diverse soils on the BCD liquor. Chemical analysis (GPC and 2D HSQC NMR) demonstrated lignin depolymerization by cleavage of β -O-4 linkage during 2-week incubation. Time series metatranscriptomics revealed that predominant microbes produce enzymes involved in lignin depolymerization to aromatic catabolism and ring cleavage.

Reference

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