

New Separation Method Reveals Thousands of Hidden Compounds in Renewable Bio-Oils

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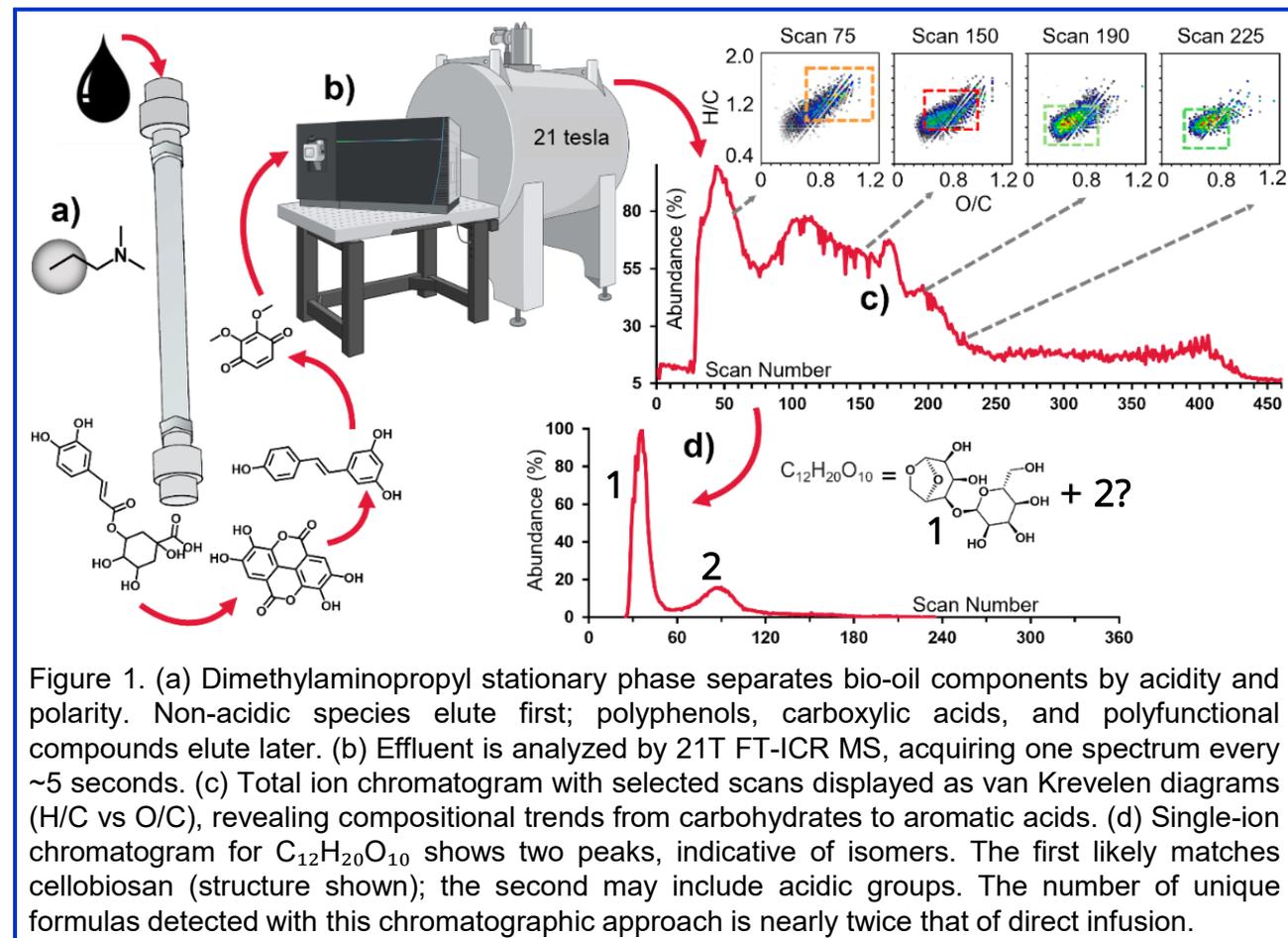
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Fast pyrolysis of lignocellulosic biomass yields oxygen-rich bio-oils, which serve as promising feedstocks for sustainable fuel and chemical production. However, their molecular complexity and abundance of acidic, polyfunctional species complicate analysis by direct infusion FT-ICR mass spectrometry (MS), which suffers from ion suppression and limited structural resolution. To address this, MagLab users developed a partition-based chromatography method using a polymeric stationary phase with amino functionalities (Figure 1a). This phase retains acidic compounds through reversible acid–base interactions. A solvent gradient from methanol to water with diethylamine creates a high-pH, hydrogen-bonding environment that separates compounds based on their acidity and polarity. As more water is added, different types of acidic compounds—like phenols, carboxylic acids, and lignin-based oligomers—elute sequentially for MS analysis.

Coupling the separation to 21T FT-ICR MS (Figure 1b) enables assignment of tens of thousands of molecular formulas per experiment. This time-resolved approach captures compositional trends, as shown with van Krevelen diagrams (Figure 1c) plotting formulas by H/C and O/C ratios for selected scans. Early spectra (e.g., scan 75) contain methanol-soluble carbohydrates (high H/C, O/C); intermediate scans (e.g., 150) show biomass-like polyphenols; and later scans (e.g., 190/225) highlight aromatic, acidic species (low H/C). The method also resolves structural isomers: compounds with the same molecular formula but different atomic connectivity. For example, the single-ion chromatogram for $C_{12}H_{20}O_{10}$ reveals two distinct peaks (Figure 1d), which indicates the presence of isomers. One likely corresponds to cellobiosan, a known pyrolysis product; the second may represent a more acidic variant. Such detail is not accessible by direct infusion analysis and is essential for improving bio-oil upgrading processes.



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