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Detection and Description of Auxotrophic Bacteria in Maize Rhizosphere and Bulk Soil: Insights into Microbial Dependencies

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By releasing amino acids and other compounds, plants shape bacterial communities in the rhizosphere, potentially favoring the enrichment of auxotrophic bacteria organisms that cannot synthesize specific amino acids and depend on external sources. We hypothesize that these auxotrophs are more abundant in the maize rhizosphere than in bulk soil and may enhance plant resilience under nutritional stress. To test this, we employed a dependency-based cultivation strategy to assess auxotrophy profiles in rhizospheric and bulk soil microbial communities associated with maize. The analysis was conducted using two contrasting soil textures (clay and sandy soils) collected from a greenhouse mesocosm experiment carried out over 50 days. Enrichment cultures were grown in minimal medium containing only minerals and carbon sources (pyruvate and glucose). Amino acids and vitamins were classified into six chemical categories and individually supplemented into the growth medium, with microbial growth monitored by optical density measurements after 24-48 hours. Soil suspensions were inoculated at dilutions of 10^{-2} , 10^{-4} , and 10^{-6} ; only the latter two allowed discrimination between treatments. Rhizospheric communities preferentially consumed aliphatic non-polar, polar and negative, and polar non-charged groups, while bulk soil communities favored aliphatic non-polar, polar and positive, aromatic, and polar non-charged compounds. Both environments strongly utilized aliphatic non-polar amino acids such as methionine, a cooperative metabolite. We are currently assessing the auxotrophic profiles of bacterial isolates obtained from these enrichment cultures, and subsequent metabolomic analyses will further elucidate their metabolic dependencies and roles in plant-microbe interactions.

KEYWORDS: Soil microbiology, Auxotrophy, Rhizosphere, Microbial dependencies

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