

# Tobacco/garlic intercropping system improve chemical properties by changing C,N cycling and plant degradation pathway in rhizosphere soil, thus increase the tobacco plant biomass

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## Abstract

As the mechanism of the microbe-soil-tobacco interaction remains unclear and the contribution of tobacco plant growth is still difficult to predict, the chemical properties and microbes of soil in tobacco/garlic intercropping system, the relevance of the soil chemical properties and the genes involved in C, N cycling and plant degradation (organic matter turnover) were studied by metagenome sequencing. The results showed that the intercropping treatment (T) significantly enhanced the content of organic matter (OM), the available nitrogen (AN), the available phosphorus (AP), the available potassium content (AK), microbe number and the microbial biomass nitrogen, as well as the activity of urease, phosphatase, invertase compared to monocropping treatment (CK). Especially the content of OM, AN, AP, AK increased significantly by 29.46%, 19.75%, 10.37%, 17.42% in rhizosphere of T treatment than CK treatment. The content of polyphenol oxidase activity and microbial biomass carbon significantly decreased in T treatment with by 22.61% and 9.03% relative to CK treatment. Metagenomic analysis showed that the relative abundances of genes related to C cycling (ACA, sdhA, sdhB, sucD, mdh), N cycling (glnA) and plant degradation (bglX) were higher in the T treatment than the CK treatment. Compared to the CK treatment, the relative abundance of ACA, sdhA, sdhB, sucD, mdh, glnA and bglX were respectively 26.06%, 39.37%, 48.27%, 32.44%, 57.55%, 14.28% and 2.39% higher in the T treatment. The intercropping system changed the chemical properties as well as the abundance of microbes, and subsequently regulate genes involved in C, N cycling and plant degradation, these improved the soil environment and led to the increase of tobacco plant biomass.

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