

Effect of elevated atmospheric CO₂ on the quantities and communities of microbes and their relationship with soil carbon and nitrogen in a typical summer maize field in North China

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Abstract

Elevated atmospheric CO₂ concentration (eCO₂) has been the most driving factor and characteristic of climate change. To clarify the effects of eCO₂ on the soil microbes and on the concurrent status of soil carbon and nitrogen, an experiment was conducted in a typical summer maize field based on a 10-year mini-FACE (Free Air Carbon Dioxide Enrichment) system in North China. Both rhizospheric and bulk soils were collected for measurement. The soil microbial carbon (MBC), nitrogen (MBN), and soil mineral N were measured at two stages. Characteristics of microbes were assayed for both rhizospheric soil and bulk soils at the key stage. We examined the plasmid copy numbers, diversities, and community structures of bacteria (in terms of 16s rRNA), fungi (in terms of ITS-Internal Transcribed Spacer), ammonia oxidizing bacteria (AOB) and denitrifiers including nirK, nirS, and nosZ using the Miseq sequencing technique. Results showed that under eCO₂ conditions, both MBC and MBN in rhizospheric soil were increased significantly. The quantity of ITS was increased in the eCO₂ treatment compared with that in the ambient CO₂ (aCO₂) treatment, while the quantity of 16s rRNA in rhizospheric soil showed decrease in the rhizospheric soil in the eCO₂ treatment. eCO₂ changed the relative abundance of microbes in terms of compositional proportion of some orders or genera particularly in the rhizospheric soil – in particular, Chaetomium increased for ITS, Subgroup 4 and Subgroup 6 increased for 16s rRNA, Nitrosospora decreased for AOB, and some genera showed increase for nirS, nirK, and nosZ. Nitrate N was the main inorganic nitrogen form at the tasseling stage and both quantities of AOB and denitrifiers, as well as the nosZ/(nirS+nirK) showed an increase under eCO₂ conditions particularly in the rhizospheric soil. The Nitrosospora decreased in abundance under eCO₂ conditions in the rhizospheric soil and some genera of denitrifiers also showed differences in abundance. eCO₂ did not change the diversities of microbes significantly. In general, results suggested that 10 years of eCO₂ did affect the active component of C and N pools (such as MBC and MBN) and both the quantities and relative abundance of microbes which are involved in carbon and nitrogen cycling, possibly due to the differences in both the quantities and component of substrate for relevant microbes in the rhizospheric soils.

Keywords: Elevated CO₂, Microbial Community, Nitrifier, Denitrifier, Ammonia Oxidizing Bacteria, Rhizosphere Soil
