

Interactive comment on “Soil microbial community structure and diversity are largely influenced by soil pH and nutrient quality in 78-year-old tree plantations” by Xiaoqi Zhou et al.

Xiaoqi Zhou et al.

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Dear Editor Prof. Yakov Kuzyakov,

We have carefully revised the manuscript according to reviewers' suggestions. The changes we made have been highlighted in yellow throughout the manuscript as shown at the attached file. We hope that it can meet the standard required by your journal Biogeosciences.

Best regards,

Xiaoqi Zhou on behalf of all authors

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Discussion paper



Reviewer #1 This study investigates long-term ecological consequences of forest plantations on the community structure and diversity of soil microorganisms. After initial improvements, the manuscript is well written in a clear way. I suggest that it can be accepted to publish in Biogeosciences after minor revision. R: Thanks.

Please see my comments as follows: 1. Although the authors have mentioned the role of microorganisms in plantations, the significance for elucidating how forest plantations affect microbial community structure and diversity. Plantations are mainly used to supply with timber. Its sustainability will be a target. It will be good if producing more timber and increasing soil organic matter at the same time. To achieve this, a prerequisite is to know how forest plantations affect microbial community structure and diversity. The reason is that microorganisms drive biogeochemical cycles. Here suggest that the authors develop a frame for microbial function to connect with the aim of this study and clarify its importance. R: This has been revised. Lines 13-16 Page 2.

2. Soil CH₄-oxidizing bacteria are one of the important microbial groups, but there are also many others. Why soil CH₄ oxidation processes and CH₄-oxidizing bacterial communities were identified in this study? The authors should clarify this. R: This has been added. Lines 1-4 Page 3.

3. In this study, soil $\delta^{15}\text{N}$ was measured but it seemed that the authors did not mention why measure it. Please clarify what information it can represent. R: Thanks for your kind comments. This has been added. Lines 1-3 Page 9.

4. This study focused on soil microbial community structure and diversity. In the text, the authors used soil bacteria and eukaryotes. It is better to briefly mention the main structure and their main functions in somewhere so that the readers are easy to follow. R: This has been added. Lines 17-19 Page 3.

5. In conclusion, the authors draw a conclusion that soil pH and nutrient quality

indicators such as C:N and EOC:EON ratios were key factors determining the patterns of soil bacterial and eukaryotic communities. If it is possible, it will be better to make this change connected with biogeochemical cycles and thus the consequences. R: This has been added. Lines 29-31 Page 10.

Please also note the supplement to this comment:

<http://www.biogeosciences-discuss.net/bg-2016-552/bg-2016-552-AC1-supplement.pdf>

Interactive comment on Biogeosciences Discuss., doi:10.5194/bg-2016-552, 2017.

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