

Review of: "Shotgun metagenomics of soil invertebrate communities reflects taxonomy, biomass and reference genome properties"

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This an interesting study by using the metagenomic approach to explore the soil invertebrates.

In the present work, the authors employed the artificial soil, I was wondering does the author validate the method in field soils?

In addition, I agree with that metagenomics offers major advances such as PCR bias than the metabarcoding sequencing. However, it would be better if the author could compare the results between metagenomics and 18S rRNA or COI gene sequencing, and present their major advantages and disadvantages. Since I believe each method has their own advantages and limitations.