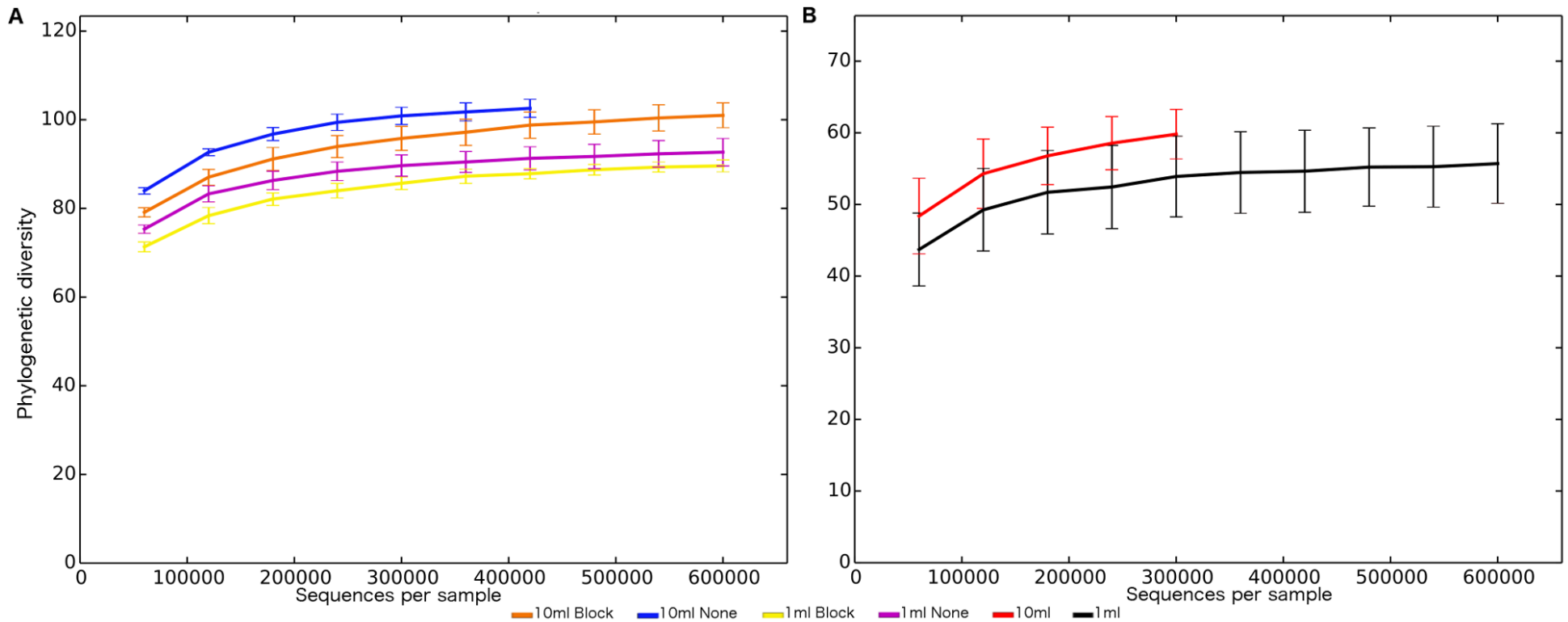


An 18S rRNA workflow for characterizing protists in sewage, with a focus on zoonotic trichomonads;

Microbial Ecology;

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Online Resource 7 Rarefaction curves for protist only OTU tables rarefied from 60,000 to 600,000 sequences with the Phylogenetic Diversity metric. Error bars are standard deviation. Lines that do not extend all the way to the end of the x-axis indicate that at least one sample in that category does not have that many sequences. (A) V9 region and (B) V4 region.