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Correction to: Rhizosphere microbiomes diverge among *Populus trichocarpa* planthost genotypes and chemotypes, but it depends on soil origin



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Correction to: Microbiome 7, 76 (2019) https://doi.org/10.1186/s40168-019-0668-8

Following publication of the original article [1], the authors identified an error in Fig. 2. The correct figure is given below.

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Reference

 Veach AM, Morris R, Yip DZ, et al. Rhizosphere microbiomes diverge among Populus trichocarpa plant-host genotypes and chemotypes, but it depends on soil origin. Microbiome. 2019;7:76 https://doi.org/10.1186/s40168-019-0668-9

The original article can be found online at https://doi.org/10.1186/s40168-019-0668-8.

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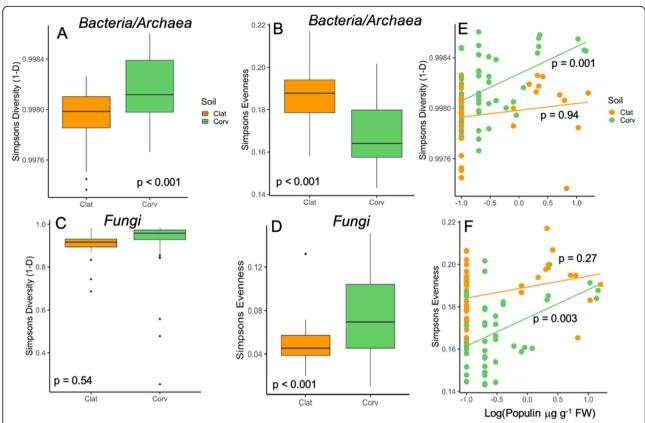


Fig. 2 Bacterial/archaeal and fungal diversity (Simpson's Diversity: 1-D; Panel **a, c**) and Simpsons's Evenness (Panel **b, d**) in Clatskanie and Corvallis soil origins. Orange boxplots and points denote Clatskanie and green denotes Corvallis soils. Bacterial/archael diversity and evenness was correlated with populin concentration in Corvallis soils (Panel **e, f**). Type-1 error rates given were generated by stepwise regression model analyses