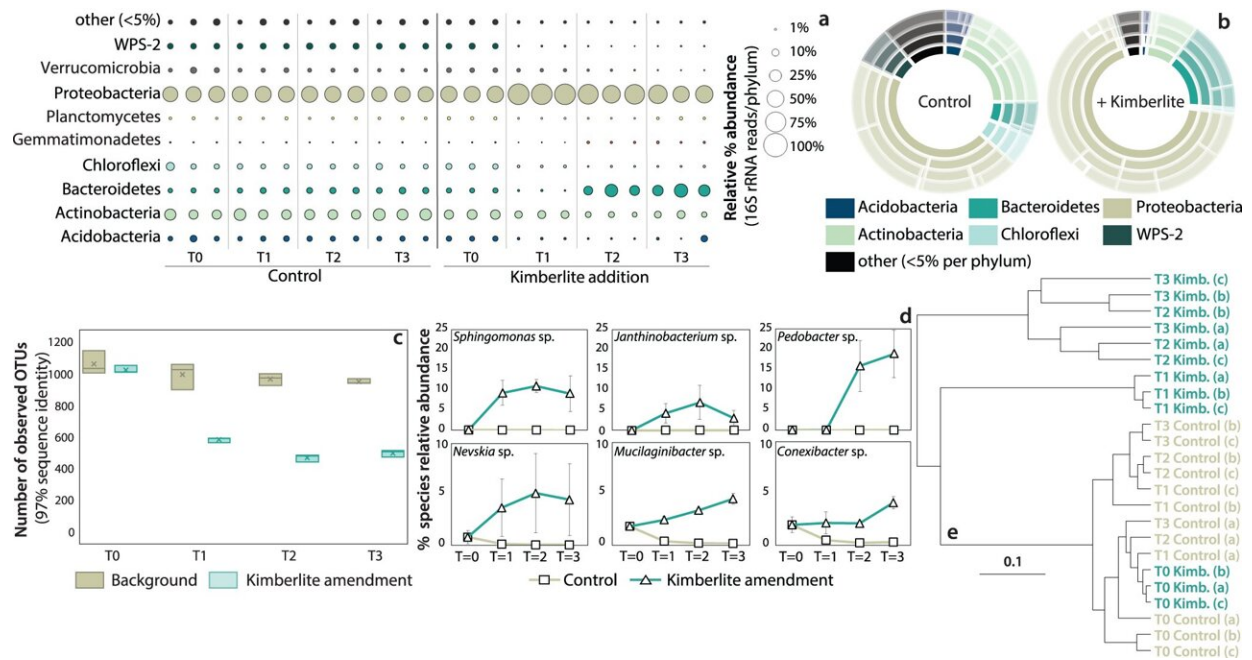


Biological fingerprints in soil show where diamond-containing ore is buried

October 24 2023



Soil microbial community composition, diversity, and indicator species for the kimberlite amendment experiment. a Distribution of 16 S rRNA gene reads per phylum for each sample. The number of reads per phylum is calculated as a percentage of the total reads for each sample. The “other” grouping represents phyla that when summed contributed (on average across all samples)

Citation: Biological fingerprints in soil show where diamond-containing ore is buried (2023, October 24) retrieved 29 April 2024 from <https://phys.org/news/2023-10-biological-fingerprints-soil-diamond-containing-ore.html>

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