

Effects of operational taxonomic unit inference method community analysis using long-read metabarcoding

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Citation Report

#	ARTICLE	IF	CITATIONS
1	MinION Sequencing of Yeast Mock Communities To Assess the Effect of Databases and ITS-LSU Markers on the Reliability of Metabarcoding Analysis. <i>Microbiology Spectrum</i> , 2023, 11, .	3.0	3
2	Intragenomic diversity of the V9 hypervariable domain in eukaryotes has little effect on metabarcoding. <i>IScience</i> , 2023, 26, 107291.	4.1	2
3	Considerations for the inclusion of metabarcoding data in the plant protection product risk assessment of the soil microbiome. <i>Integrated Environmental Assessment and Management</i> , 2024, 20, 337-358.	2.9	1