

Bacterial diversity in ferruginous duricrust (canga) and the physicochemical variables affecting their prevalence, distribution and predicted metabolic pathways

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Online Resource 8 Functional prediction heatmap based on the total set of detected ASVs (see Materials and Methods for details). The prediction includes the phylogenetic positioning of the sequences, the reconstruction of ancestral states, and the estimation of the relative abundance of genes and metabolic pathways per sample.

