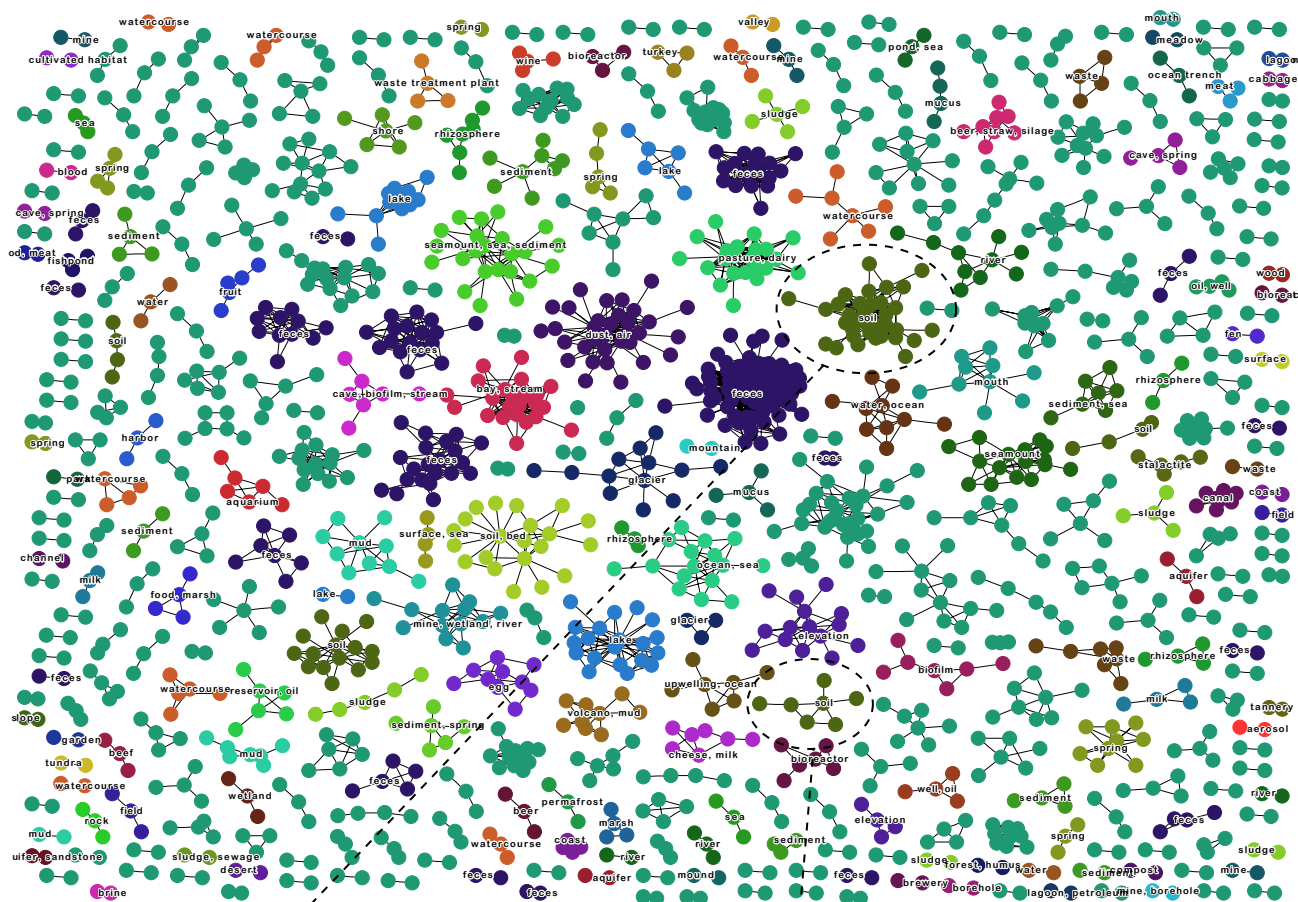


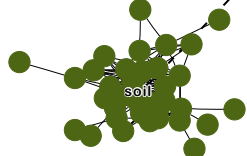
- OTU definition: 97%

- Co-occurrence corrected p-value cut-off (FDR): 0.001

Note: here, nodes represent 'sampling sites', not OTUs.

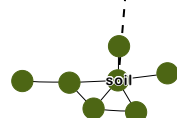


A



Proteobacteria:	80.3%
Actinobacteria:	8.37%
Firmicutes:	6.61%
Bacteroidetes:	3.91%
Verrucomicrobia:	0.270%
Acidobacteria:	0.270%
Planctomycetes:	0.135%
Deinococcus-Thermus:	0.135%

B



Acidobacteria:	35.2%
Proteobacteria:	32.9%
Verrucomicrobia:	15.0%
Actinobacteria:	7.98%
Unclassified:	3.76%
Firmicutes:	2.35%
Gemmatimonadetes:	1.41%
Nitrospira:	0.939%
Planctomycetes:	0.469%

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[...]

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Changes in microbial and metabolic diversity and composition along a hydrogeochemically variable soil profile (unsaturated B horizon soil aggregate)

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