

Microbial diversity and biotechnological potential of terrestrial and aquatic environments from Trindade Island, Brazil: Trends and Perspectives

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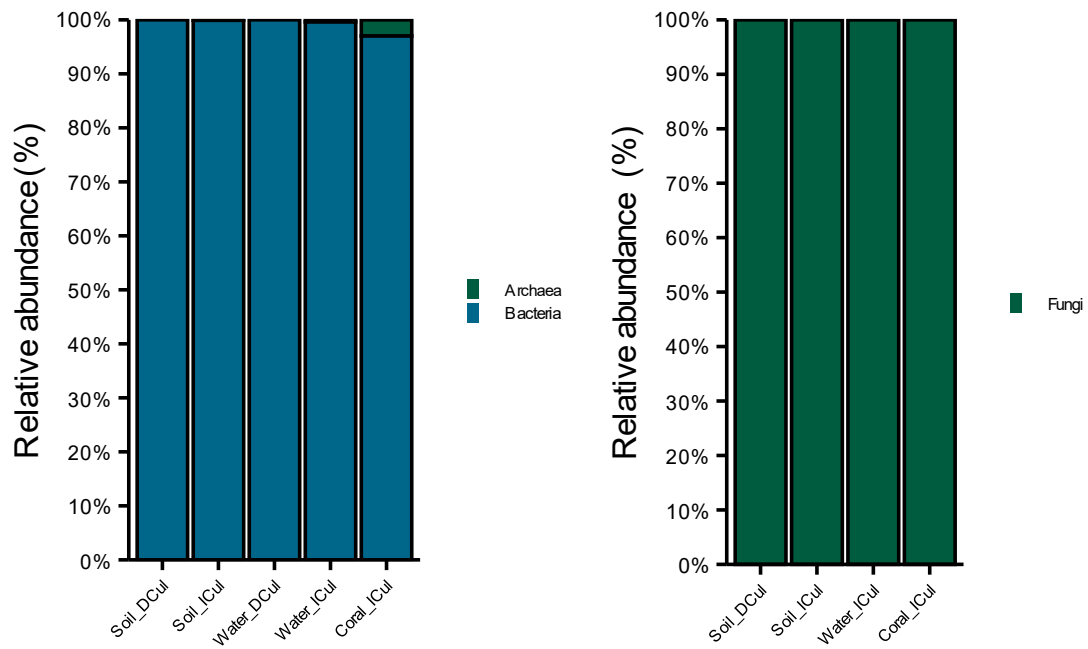


Figure S1. Bar graph of the superkingdom of Archaea and Bacteria and the kingdom of Fungi of the grouped samples.

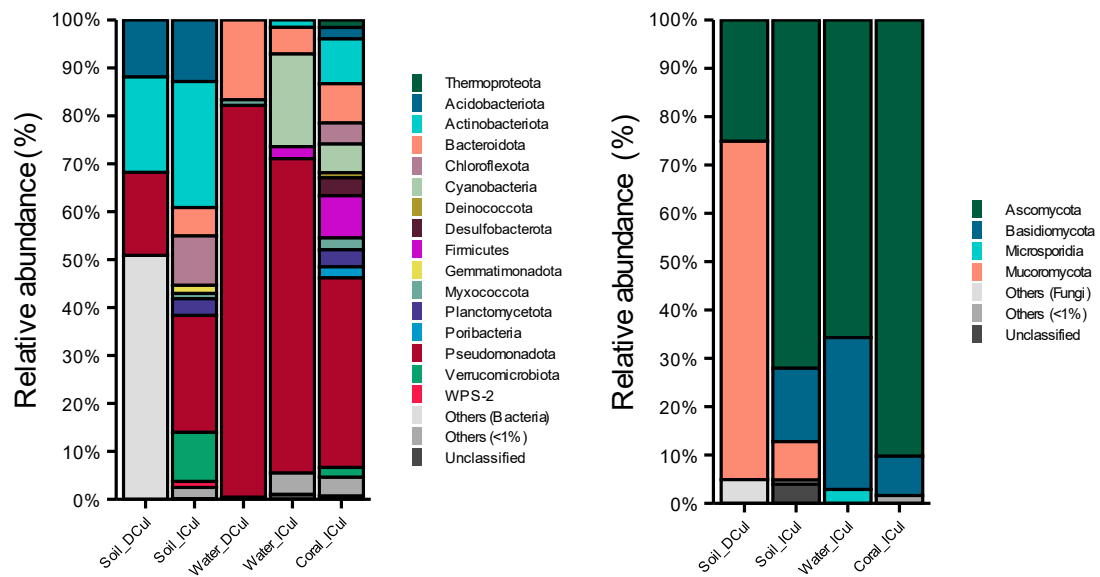


Figure S2. Bar graph of the phyla of Archaea, Bacteria and Fungi of the grouped samples.

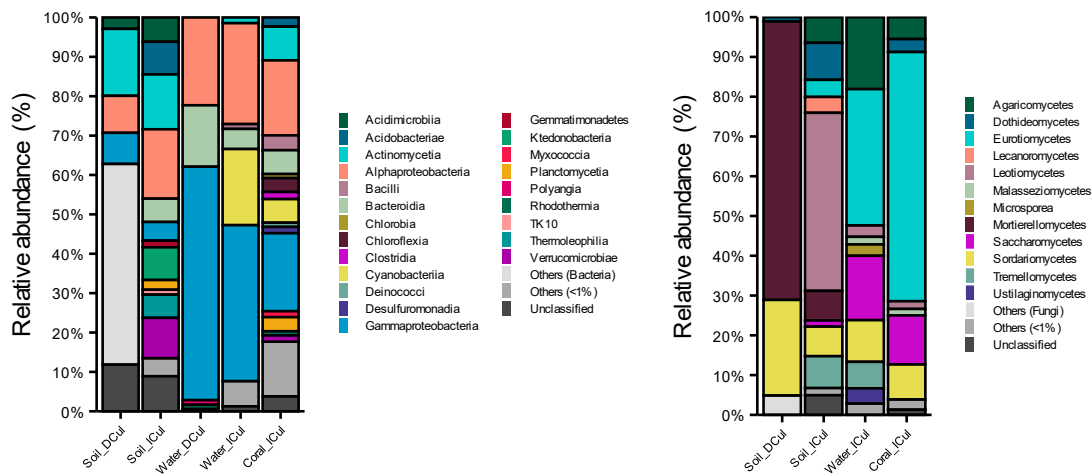


Figure S3. Bar graph of the classes of Archaea, Bacteria and Fungi of the grouped samples.

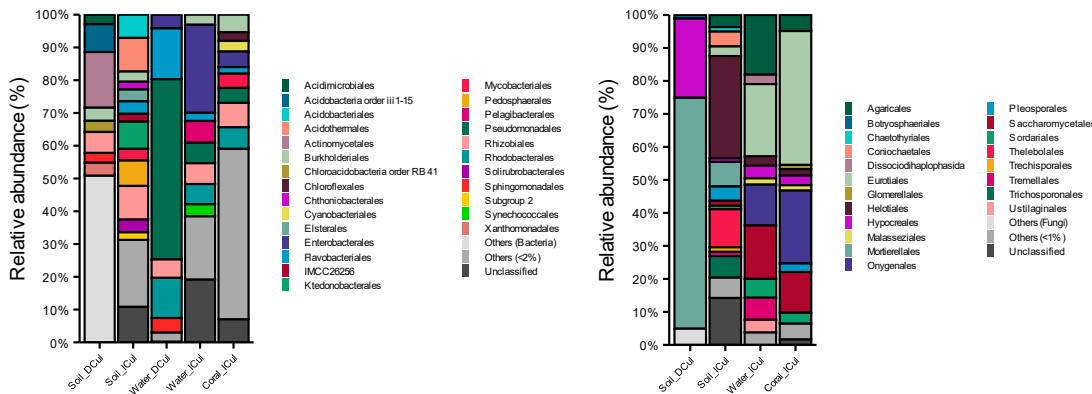


Figure S4. Bar graph of the orders of Archaea, Bacteria and Fungi of the grouped samples.

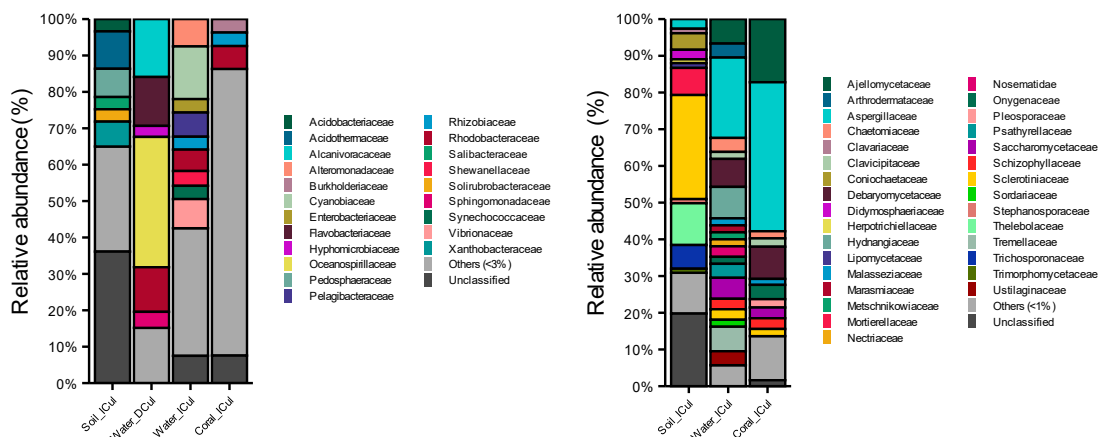


Figure S5. Bar graph of the families of Archaea, Bacteria and Fungi of the grouped samples.

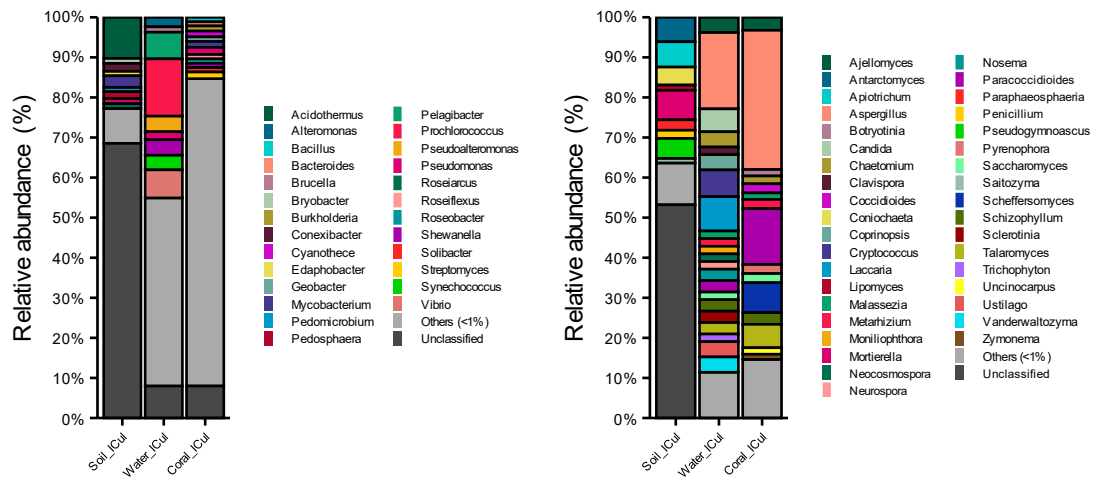


Figure S6. Bar graph of the genera of Archaea, Bacteria and Fungi of the grouped samples.

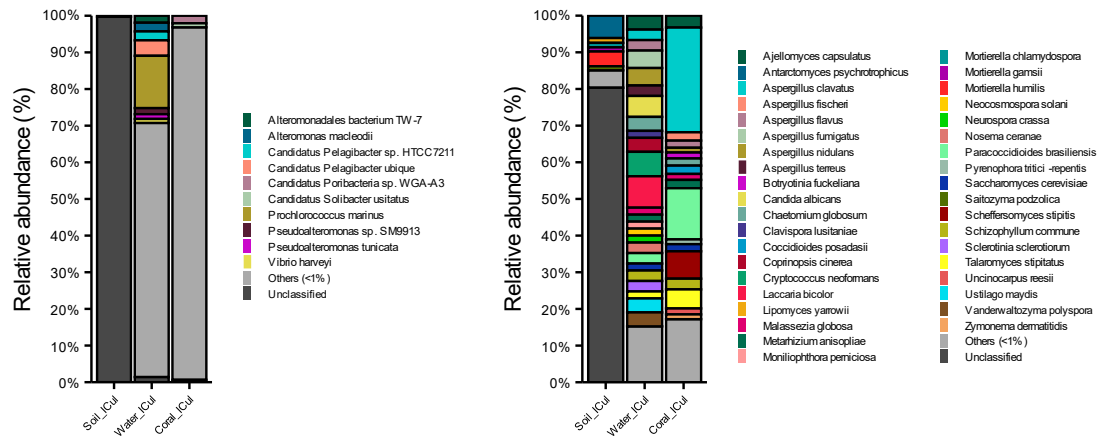


Figure S7. Bar graph of the species of Archaea, Bacteria and Fungi of the grouped samples.

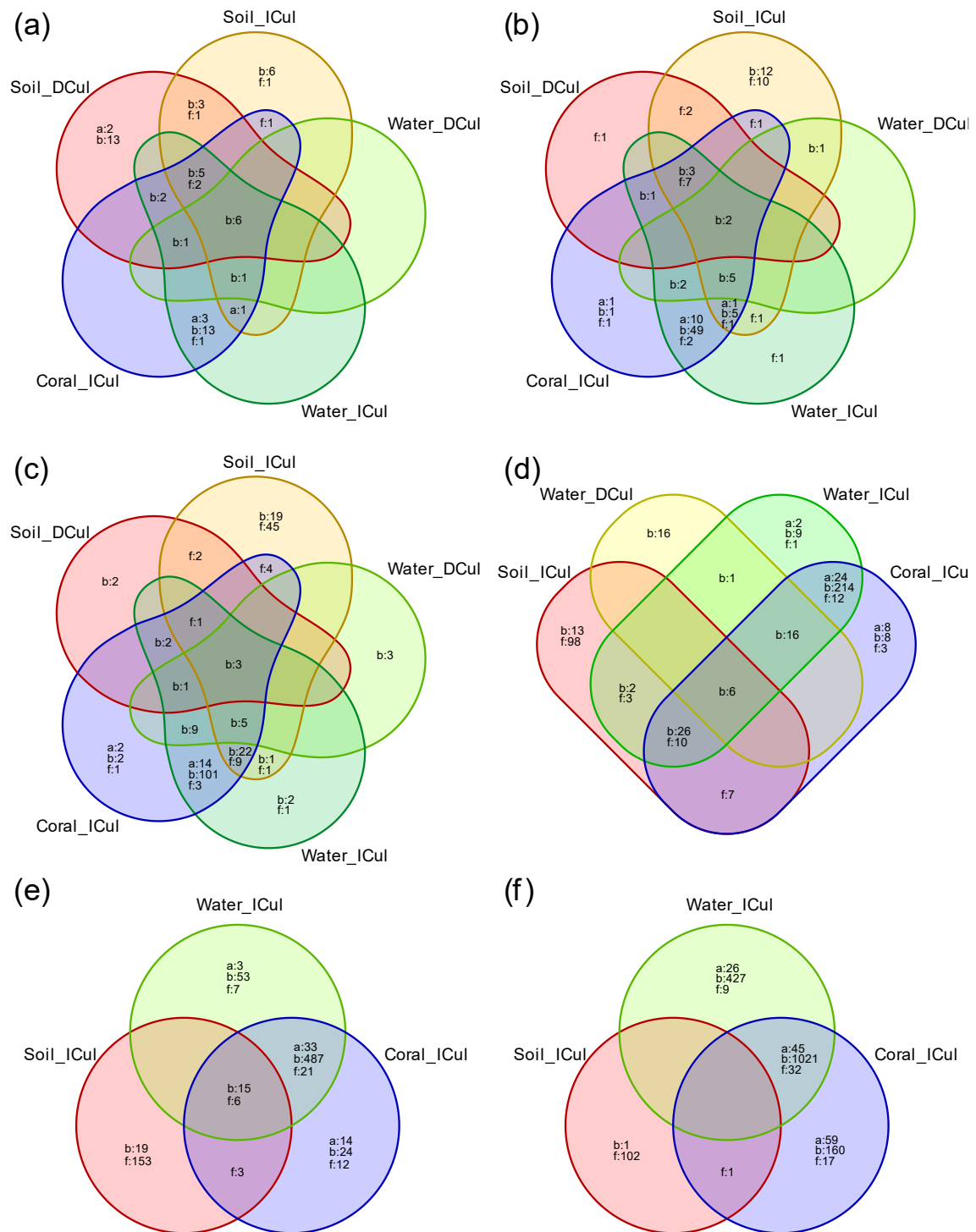


Figure S8. Venn diagram of the categories of Phylum (a), Class (b), Order (c), Family (d), Genus (e) and Species (f) of Archaea, Bacteria and Fungi of the grouped samples.

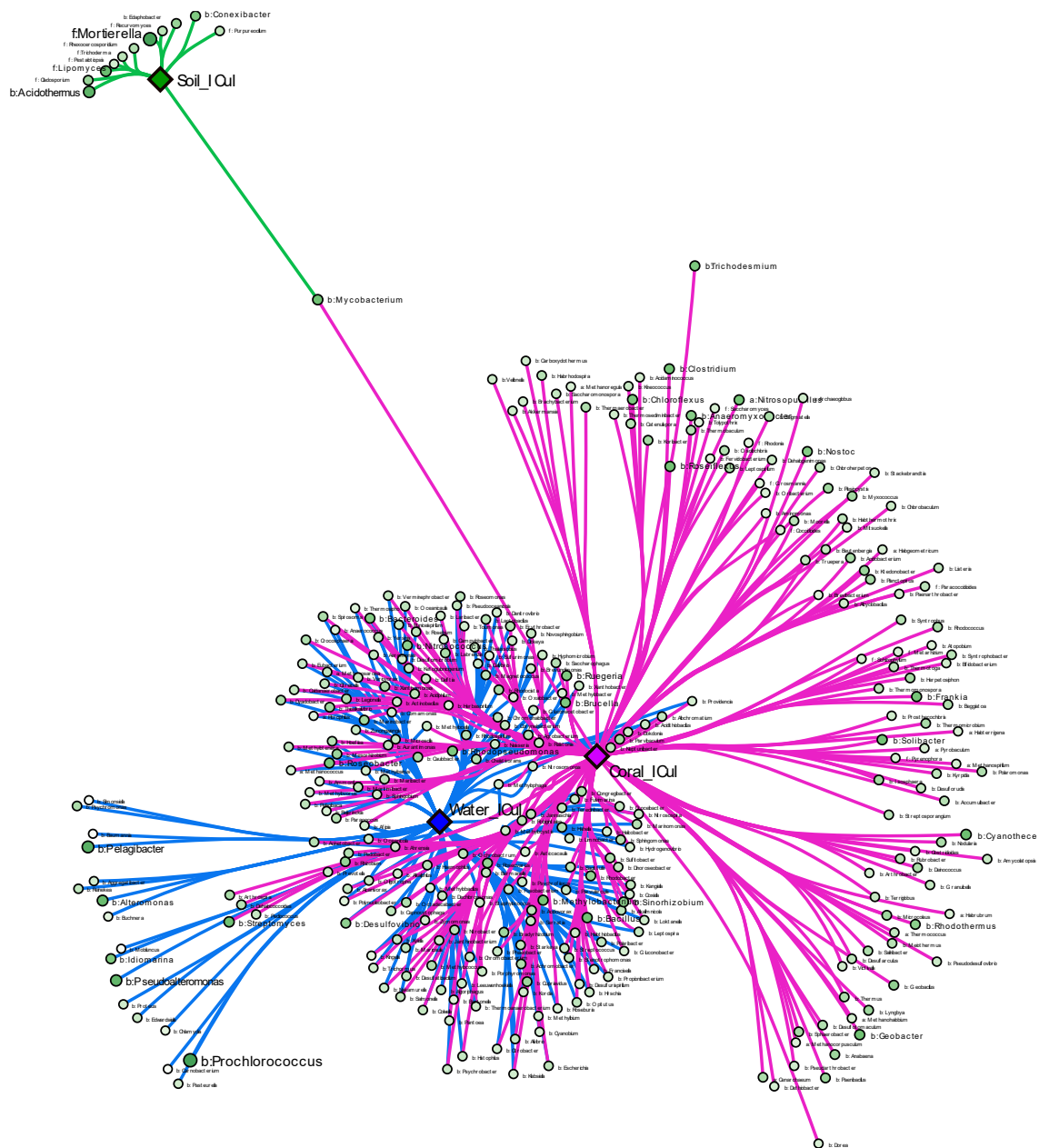


Figure S13. Strong association network between Archaeal, Bacterial and Fungal genera and grouped samples.

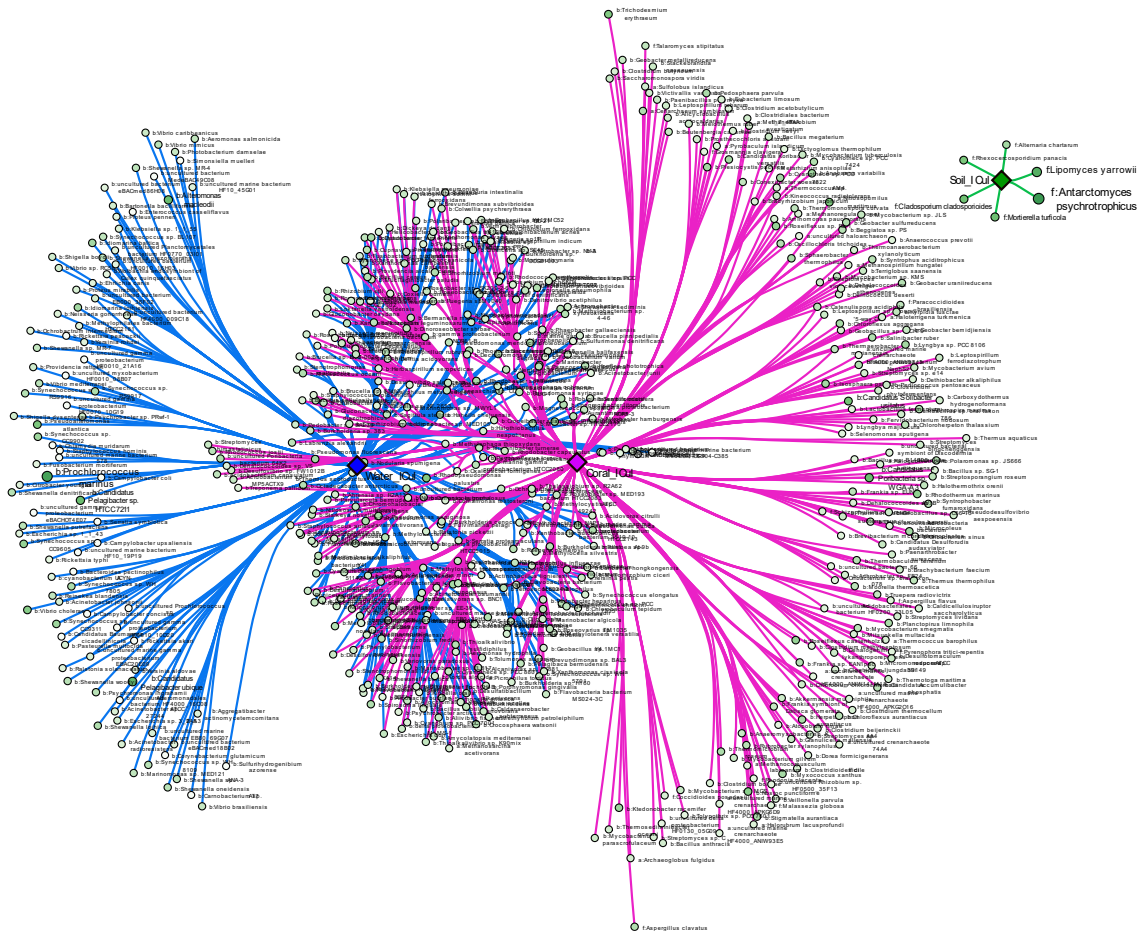


Figure S14. Strong association network between Archaeal, Bacterial and Fungal species and grouped samples.