Supplementary figures

Salinity gradient controls microbial community structure and assembly in coastal solar salterns

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Running title: Microbial community assembly in the solar saltern

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Conflict of Interest

No conflict of interest exists in the submission of this manuscript, and the manuscript has been approved by all authors for publication.

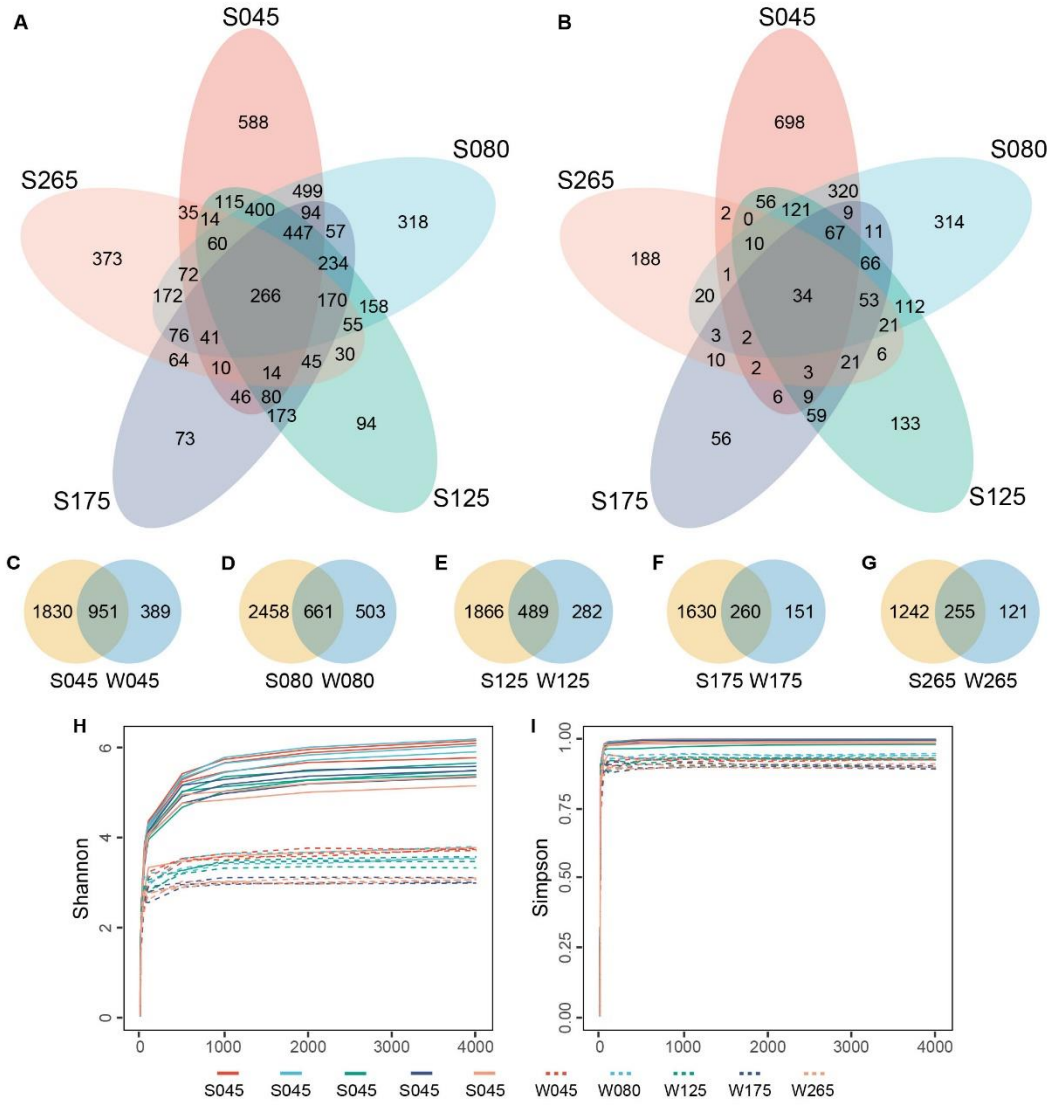

Figure S1. Venn diagrams showing the shared and unique OTUs.

A-B, Visualization of venn diagrams in sediment and water groups. **C-G**, Visualization of venn diagrams at same sampling site, comparing the shared and unique OTUs between sediment and water samples. **H-I**, Visualization of rarefaction curve on basis of Shannon and simpson index.

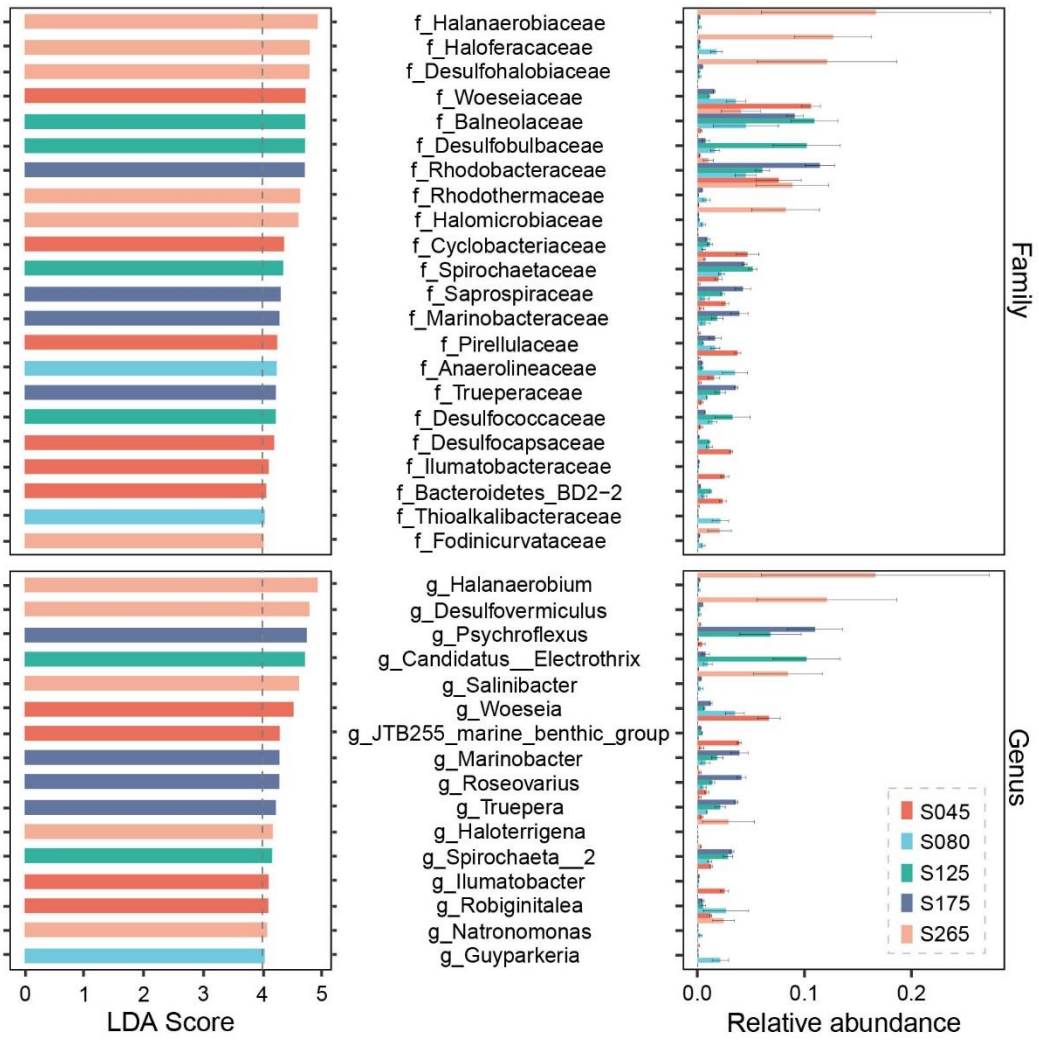

Figure S2. Differential abundance test in sediment of salterns.

The significant family and genus taxa, determining the community differences across groups, were confirmed by R package {microeco} on basis of LEFSe method.

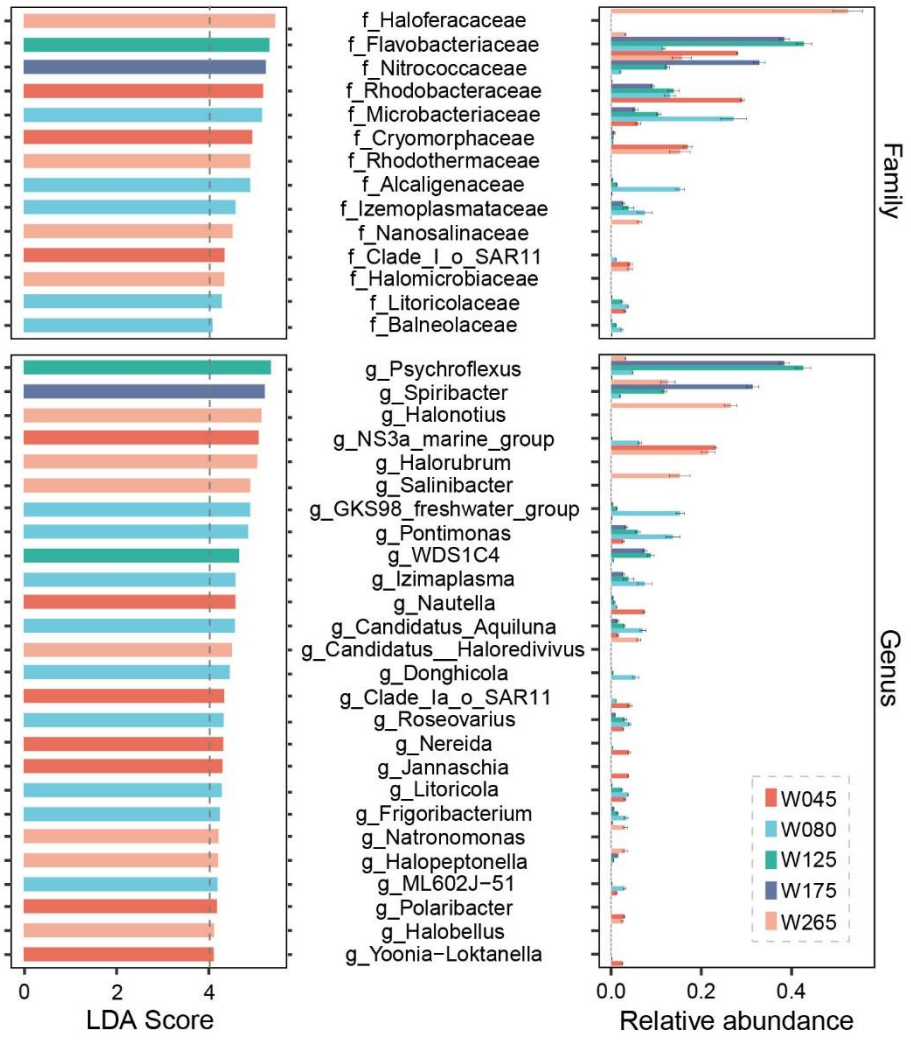

Figure S3. Differential abundance test in water of salterns.

The significant family and genus taxa, determining the community differences across groups, were confirmed by R package {microeco} on basis of LEFSe method.

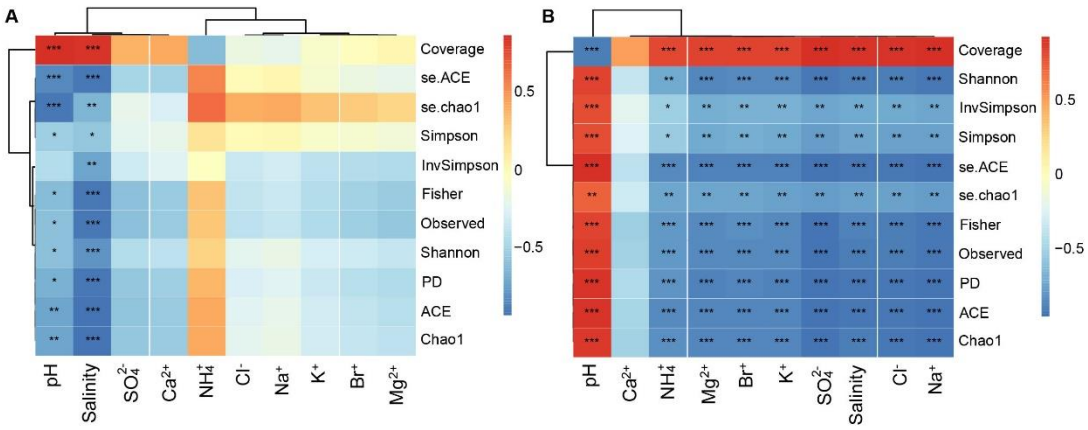

Figure S4. The relationships between physicochemical factors and alpha diversities of prokaryotic community.

A-B, The strong-related genus taxa with physicochemical factors were shown (A, sediment groups; B, water groups). Significant differences were marked. (***** $0.01 < P \leq 0.05$, ****** $0.001 < P \leq 0.01$, ******* $P \leq 0.001$)