

Table S1. Topological properties of the observed molecular ecological networks (MENs) and their associated random MENs

| | Non-salinity | | Mild-salinity | | Severe-salinity | |
|---|------------------|-----------------|------------------|-----------------|------------------|-----------------|
| Network Indexes | Observed Network | Random Networks | Observed Network | Random Networks | Observed Network | Random Networks |
| Average clustering coefficient (avgCC) | 0.248 | 0.047 +/- 0.002 | 0.207 | 0.055 +/- 0.003 | 0.339 | 0.064 +/- 0.002 |
| Average path distance (GD) | 5.303 | 2.882 +/- 0.010 | 6.094 | 3.091 +/- 0.014 | 3.852 | 2.421 +/- 0.005 |
| Geodesic efficiency (E) | 0.228 | 0.377 +/- 0.001 | 0.2 | 0.350 +/- 0.001 | 0.305 | 0.445 +/- 0.001 |
| Harmonic geodesic distance (HD) | 4.389 | 2.653 +/- 0.007 | 5.002 | 2.856 +/- 0.009 | 3.284 | 2.247 +/- 0.003 |
| Centralization of degree (CD) | 0.06 | 0.060 +/- 0.000 | 0.058 | 0.058 +/- 0.000 | 0.087 | 0.087 +/- 0.000 |
| Centralization of betweenness (CB) | 0.047 | 0.015 +/- 0.002 | 0.118 | 0.020 +/- 0.002 | 0.052 | 0.010 +/- 0.001 |
| Centralization of stress centrality (CS) | 517.933 | 0.163 +/- 0.012 | 648.053 | 0.198 +/- 0.019 | 115.604 | 0.163 +/- 0.011 |
| Centralization of eigenvector centrality (CE) | 0.177 | 0.094 +/- 0.004 | 0.161 | 0.103 +/- 0.007 | 0.109 | 0.061 +/- 0.002 |
| Density (D) | 0.019 | 0.019 +/- 0.000 | 0.014 | 0.014 +/- 0.000 | 0.035 | 0.035 +/- 0.000 |
| Reciprocity | 1 | 1.000 +/- 0.000 | 1 | 1.000 +/- 0.000 | 1 | 1.000 +/- 0.000 |
| Transitivity (Trans) | 0.228 | 0.059 +/- 0.001 | 0.245 | 0.077 +/- 0.001 | 0.297 | 0.069 +/- 0.001 |
| Connectedness (Con) | 0.862 | 0.982 +/- 0.008 | 0.908 | 0.988 +/- 0.007 | 0.965 | 0.995 +/- 0.004 |
| Efficiency | 0.979 | 0.982 +/- 0.000 | 0.986 | 0.987 +/- 0.000 | 0.965 | 0.966 +/- 0.000 |
| Hierarchy | 0 | 0.000 +/- 0.000 | 0 | 0.000 +/- 0.000 | 0 | 0.000 +/- 0.000 |
| Lubness | 1 | 1.000 +/- 0.000 | 1 | 1.000 +/- 0.000 | 1 | 1.000 +/- 0.000 |
| Modularity(fast_greedy) | 0.775 | 0.189 +/- 0.003 | 0.74 | 0.221 +/- 0.003 | 0.677 | 0.125 +/- 0.003 |

Table S2. Pair-wise comparison of network parameters among salinity levels

| | Mild vs Non | Severe vs Non | Severe vs Mild |
|--|-------------|---------------|----------------|
| Average clustering coefficient (avgCC) | 0.0001 | 0.0001 | 0.0001 |
| Average path distance (GD) | 0.0001 | 0.0001 | 0.0001 |
| Geodesic efficiency (E) | 0.0001 | 0.0001 | 0.0001 |
| Harmonic geodesic distance (HD) | 0.0001 | 0.0001 | 0.0001 |
| Centralization of degree (CD) | \ | \ | \ |
| Centralization of betweenness (CB) | 0.0001 | 0.0001 | 0.0001 |
| Centralization of stress centrality (CS) | 0.0001 | 0.0001 | 0.0001 |
| Centralization of eigenvector centrality (CE) | 0.0001 | 0.0001 | 0.0001 |
| Density (D) | \ | \ | \ |
| Reciprocity | \ | \ | \ |
| Transitivity (Trans) | 0.0001 | 0.0001 | 0.0001 |
| Connectedness (Con) | 0.0001 | 0.0001 | 0.0001 |
| Efficiency | \ | \ | \ |
| Hierarchy | \ | \ | \ |
| Lubness | \ | \ | \ |
| Modularity(fast_greedy) | 0.0001 | 0.0001 | 0.0001 |

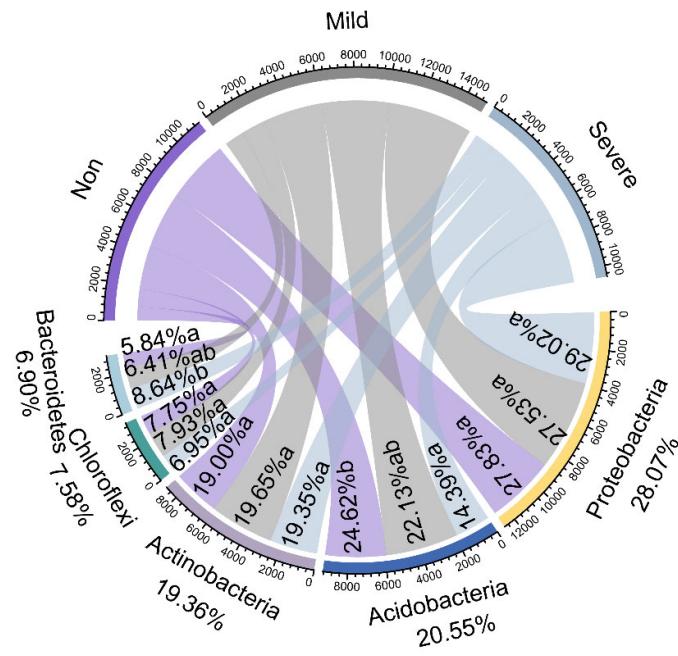


Figure S1. Phylogenetic distributions of dominant bacterial phyla in soils with three salinity levels, Non-salinity, Mild-salinity and Severe-salinity. Different letters denote significant phylogenetic differences ($p<0.05$) among salinity levels.

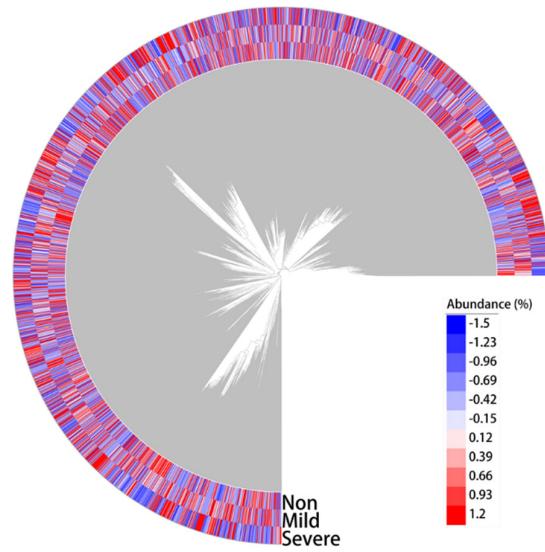


Figure S2. Phylogenetic distribution of the 3123 bacteria and their relative abundance among three salinity soils.