

Supplementary Information

Synergistic genomic mechanisms of adaptation to ocean acidification in a coral holobiont

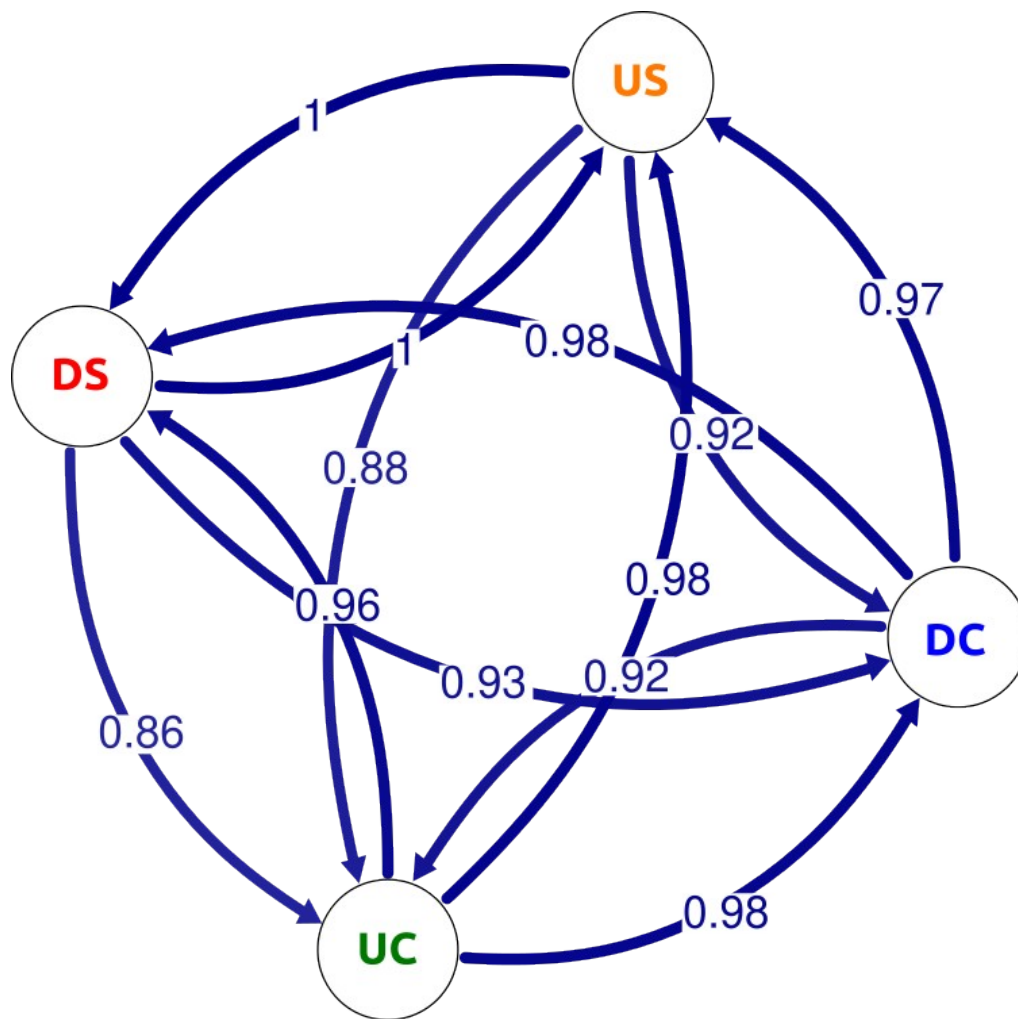
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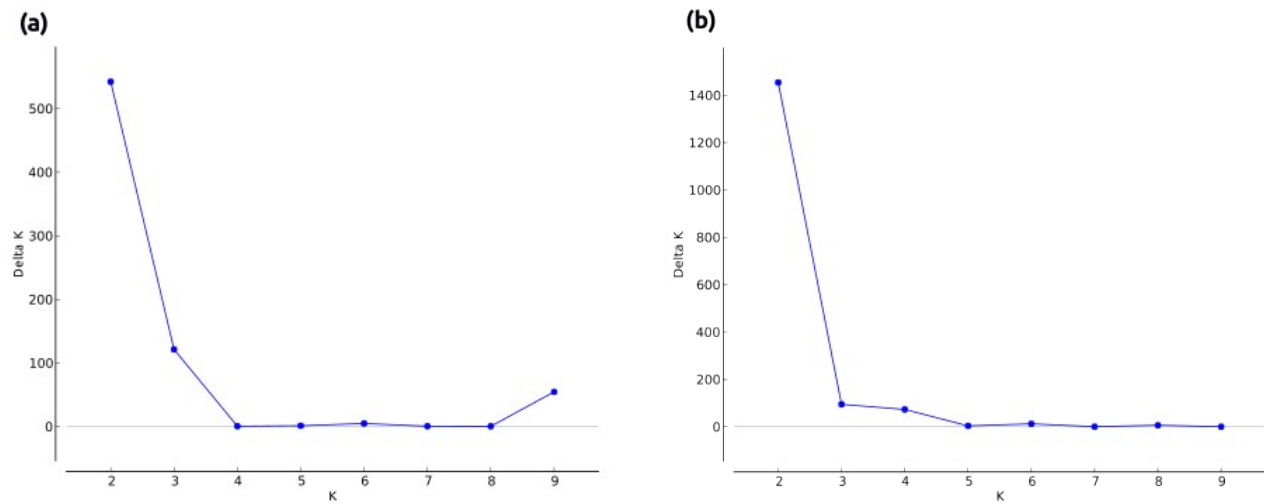
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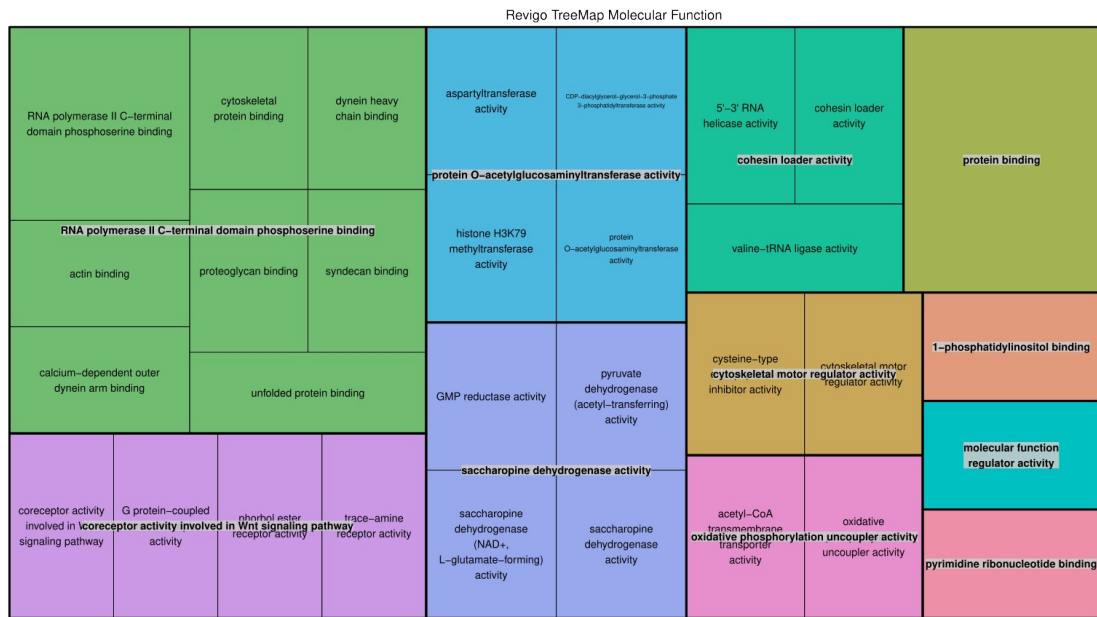
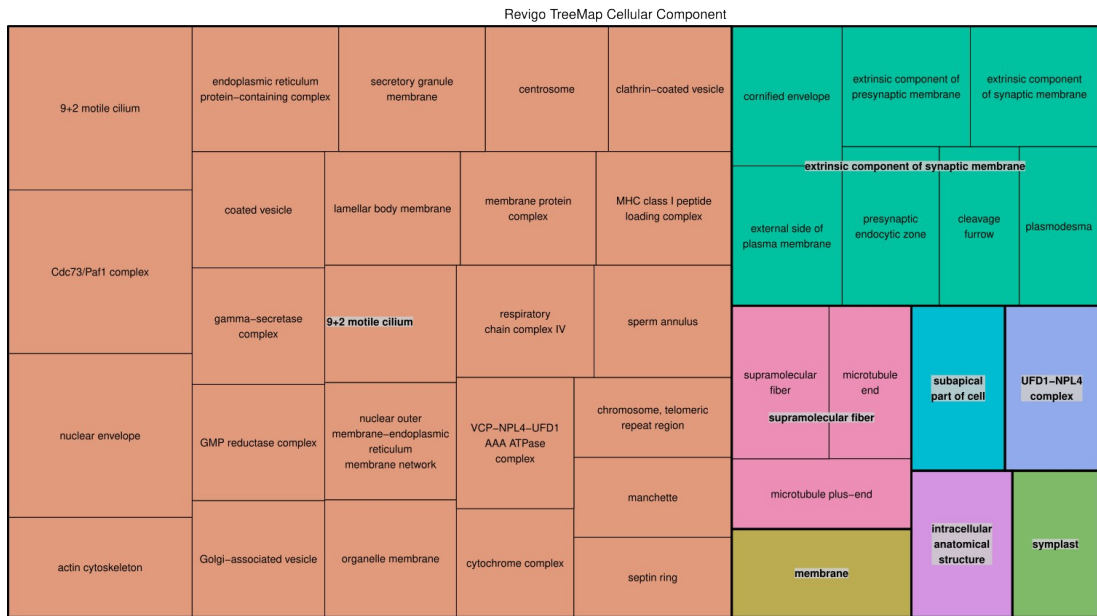
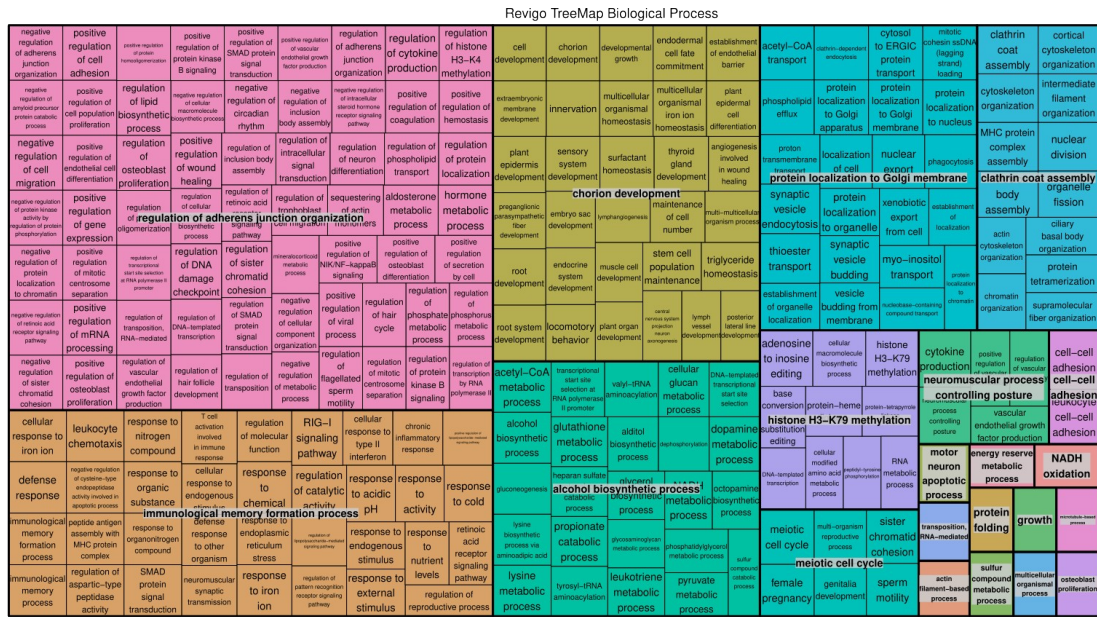
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Supplementary Figure 1. G_{ST} migration network. Circles represent sampling sites: red DS: Dobu Seep; blue DC, Dobu Control; orange US, Upa-Upasina Seep; green UC, Upa-Upasina Control.



Supplementary Figure 2. Delta K plots for the STRUCTURE analyses. (a) Using the 11,169 independent neutral SNP dataset and (b) using the 625 candidate adaptive SNP dataset.



Supplementary Figure 3. REVIGO treemaps for Biological Process, Cellular Component and Molecular Function.