Island Area, Not Isolation, Drives Taxonomic, Phylogenetic and Functional Diversity of Ants on Land-bridge Islands

Running title: Taxonomic, phylogenetic and functional diversity of island ants

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File List:

JBi-19-0424_Ant_community_matrices_on_islands.xlsx (ant species presence absence on islands)

JBi-19-

0424_Ant_community_matrices_on_transects_of_seven_largest_islands.xlsx (ant species presence absence on transects)

JBi-19-0424 DNA alignments (DNA alignments)

JBi-19-0424_island_data.xlsx (species richness, island area, isolation, phylogenetic diversity, functional diversity, mean pairwise distance, standardized effect size of mean pairwise distance, functional equivalent of mean pairwise distance, standardized effect size of functional equivalent of mean pairwise distance on all islands)

JBi-19-0424_phylogenetic_tree.tre (phylogenetic tree)

JBi-19-0424_Transect_data.xlsx (same to JBi-19-0424_island_data, but on transects)

File descriptions:

JBi-19-0424 Ant community matrices on islands.xlsx

33 islands (rows) \times 97 species (columns) community matrix (i.e., island scale in published paper)

JBi-19-

0424_Ant_community_matrices_on_transects_of_seven_largest_islands.xlsx57 transects (rows) × 97 species (columns) community matrix (i.e., transect scale in published paper)

JBi-19-0424 DNA alignments

18S, 28S and wingless DNA alignments of 97 ant species captured in the Thousand Island Lake, China

JBi-19-0424 island data.xlsx

Island ID, species richness (SR), island area (A), isolation (I), phylogenetic diversity (PD), functional diversity (FD), mean pairwise distance (MPD), standardized effect size of mean pairwise distance (sesMPD), functional equivalent of mean pairwise distance (FD_MPD), standardized effect size of functional equivalent of mean pairwise distance (sesFD_MPD) on 33 study islands in the Thousand Island Lake, China.

JBi-19-0424 phylogenetic tree.tre

Phylogenetic tree of 97 ant species captured in captured in the Thousand Island Lake, China. The tree was constructed using BEAST v. 1.10.4 based on three nuclear genes (18S, 28S and wingless; i.e., **JBi-19-0424 DNA alignments**).

JBi-19-0424_Transect_data.xlsx

Transect ID, Island ID, species richness (SR), island area (A), isolation (I), phylogenetic diversity (PD), functional diversity (FD), mean pairwise distance (MPD), standardized effect size of mean pairwise distance (sesMPD), functional equivalent of mean pairwise distance (FD_MPD), standardized effect size of functional equivalent of mean pairwise distance (sesFD_MPD) on 57 transects of 33 study islands in the Thousand Island Lake, China.