**Electronic Supplementary Material:**

**Haemosporidian taxonomic composition, network centrality and partner fidelity among resident and non-resident avian hosts**

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Supplementary Table 3: Parameter estimates, standard errors, and credible intervals for the Bayesian model testing the differences in weighted betweenness among avian hosts from distinct migratory categories. (Residents only = reference category)

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | **Estimate** | **Std. error** | **Cred. Inter (95%)** | |
| Intercept | 0.09 | 0.11 | -0.14 | 0.29 |
| Full migratory host species | 0.03 | 0.22 | -0.50 | 0.37 |
| Partial migratory host species | -0.01 | 0.13 | -0.27 | 0.24 |
| Biomes | 0.09 | 0.09 | 0.01 | 0.32 |
| Avian host phylogeny | 0.16 | 0.08 | 0.02 | 0.32 |

Supplementary Table 4: Parameter estimates, standard errors, and credible intervals for the Bayesian model testing the differences in weighted betweenness between resident and non-resident avian hosts. (Residents only = reference category)

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | **Estimate** | **Std. error** | **Cred. Inter (95%)** | |
| Intercept | 0.07 | 0.13 | -0.29 | 0.28 |
| Non-resident host species | -0.01 | 0.12 | -0.24 | 0.23 |
| Biomes | 0.09 | 0.09 | 0.01 | 0.37 |
| Avian host phylogeny | 0.16 | 0.12 | 0.01 | 0.32 |