Supporting Table 1. Population summary statistics for each species.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Species | ΦST | *h* | *π* | NE |
| *H. claudia* | 0.064, **0.123** | 0.572 + 0.049, **0.996 + 0.002** | 0.002 + 0.001, **0.026 + 0.014** | 29,231; **141,790** |
| *H. ornatissimus* | 0.005, **0.009** | 0.407 + 0.036, **0.981 + 0.004** | 0.001 + 0.001, **0.016 + 0.009** | 18,015; **91,196** |
| *H. trimaculatus* | 0.166, **0.095** | 0.754 + 0.017, **0.967 + 0.007** | 0.002 + 0.002, **0.027 + 0.014** | 36,503; **148,634** |
| *H. margaritaceus* | 0.271, **0.241** | 0.821 + 0.030, **0.994 + 0.003** | 0.006 + 0.003, **0.040 + 0.020** | 96,021; **221,791** |

Overall population statistics for each species. Control region values are in bold, following CO1 values. ΦST, Haplotype diversity (*h*), nucleotide diversity (*π*), and effective population sizes (NE) are given for each species.