**Supplementary Table S2.** Mean pairwise genetic distance ± standard deviation (%) for all taxa and localities based on concatenated-gene dataset (COI + 12S rRNA + 16S rRNA + 28S rRNA). Yellow, *Indothais gradata,* Green, *I. javanica,* Red, *I. rufotincta*

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | Light BD | Dark BD | Light PB/OR | Dark PB/U | Dark PB/M | BD | PB/OR | PB/OR |
| Light BD |  |  |  |  |  |  |  |  |
| Dark BD | 0.17 ± 0.18 |  |  |  |  |  |  |  |
| Light PB/OR | 0.03 ± 0.02 | 0.19 ± 0.20 |  |  |  |  |  |  |
| Dark PB/U | 0.39 ± 0.02 | 0.35 ± 0.07 | 0.42 ± 0.02 |  |  |  |  |  |
| Dark PB/M | 0.18 ± 0.16 | 0.25 ± 0.15 | 0.20 ± 0.18 | 0.37 ± 0.09 |  |  |  |  |
| BD | 5.07 ± 0.03 | 5.10 ±0.03 | 5.10 ± 0.03 | 5.19 ±0.02 | 5.08 ± 0.04 |  |  |  |
| PB/OR | 4.98 ± 0.03 | 5.02 ± 0.02 | 5.02 ± 0.02 | 5.10 ± NA | 5.00 ± 0.05 | 0.15 ± 0.05 |  |  |
| PB/OR | 6.76 ± 0.03 | 6.83 ± 0.06 | 6.79 ± 0.03 | 6.98 ± 0.02 | 6.85 ± 0.04 | 6.30 ± 0.03 | 6.27 ± 0.02 |  |
| T | 6.79 ± 0.04 | 6.87 ± 0.06 | 6.83 ± 0.04 | 7.01 ± 0.04 | 6.88 ± 0.05 | 6.33 ± 0.03 | 6.30 ± 0.02 | 0.07 ± 0.04 |