**S6 Table. Maximum likelihood values and statistics for the alternative phylogenetic hypotheses calculated in GARLI and CONSEL.**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Hypothesis** | **Likelihood scores (lnL)** | **AU** | **KH** | **SH** | **WKH** | **WSH** |
| This study | - 3749.171 | 0.703 | 0.71 | 0.949 | 0.71 | 0.958 |
| Vietnam monophyly | - 3844.787 | 0.297 | 0.29 | 0.559 | 0.29 | 0.486 |
| Sumatra monophyly | - 3796.949 | <0.001\* | <0.001\* | <0.001\* | <0.001\* | <0.001\* |
| Borneo monophyly | - 3755.026 | <0.001\* | <0.001\* | <0.001\* | <0.001\* | <0.001\* |

\*Statistically significant at *P* < 0.05 rejecting the hypothesis tested.