**Table S2.** Summary of nucleotide substitution models of evolution for each partition

|  |  |  |
| --- | --- | --- |
| Subset | Best Model | Partition names |
| 1 | TN+F+I | cyt*b*\_pos (1-2) | |
| 3 | K2P | cyt*b*\_pos 3 |
| - | HKY | Whole partial *cytb* gene |