S1 Table: Models of nucleotide evolution for each gene were determined with jModeltest 2.1.4 using the Bayesian information criterion.

|  |  |
| --- | --- |
| Gene | Model of nucleotide evolution |
| COI | GTR+I+G |
| ArgK | HKY+G |
| Pol-II | HKY+I |
| EF1-α | HKY+G |