

Appendix 3. Substitution models selected by the JModelTest software for each DNA region included in the analyses, and number of characters in the combined data matrix.

Marker	Substitution models	No. characters
<i>trnL-F</i>	F81+G, HKY+G	1,500
<i>rps16</i>	F81+G, GTR+G, HKY+G	936
ITS	SYM+G, SYM+I	768
18S	K80+G, K80+I	1,601
26S	GTR+G, GTR+I	941
<i>rbcL</i>	HKY+G, HKY+I, K80+G, K80+I	1,303
<i>psaB</i>	HKY+G, HKY+I	1,653