

Table 2 Statistic of *cox2*-3 spacer, ITS region and RuBisCo spacer MP analyses.

Characters	<i>cox2</i> -3 spacer	ITS region	RuBisCo spacer
No. of taxa	12	10	11
Length (bp)	345	1024	259
No. of variable sites (%)	73 (20.98%)	251 (21%)	22 (8.03%)
No. of informative sites (%)	72 (20.69%)	212 (20.70%)	19 (6.93%)
MP tree length	87	218	20
Rescaled consistency index (RC)	0.93	0.98	0.91