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

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