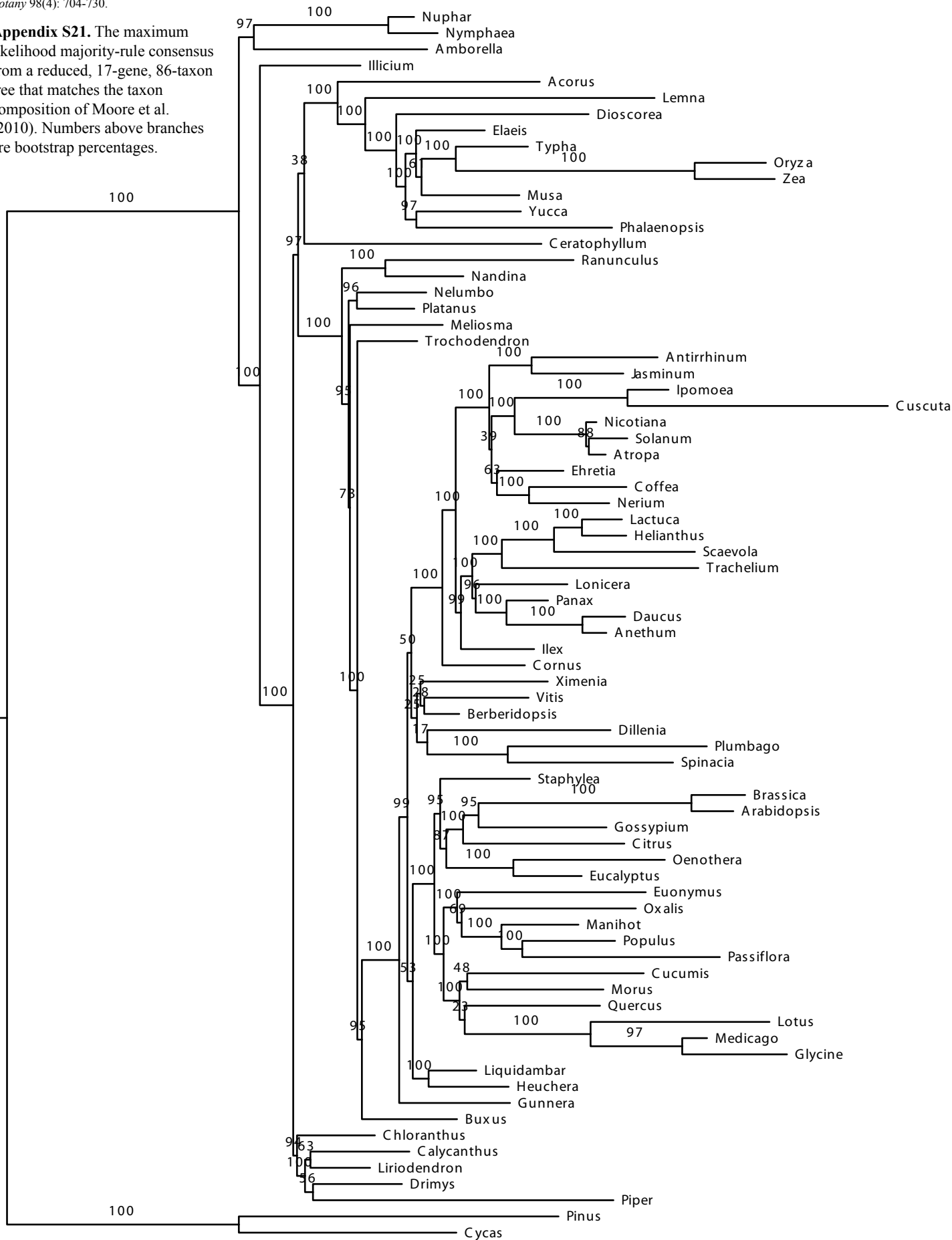


**Appendix S21.** The maximum likelihood majority-rule consensus from a reduced, 17-gene, 86-taxon tree that matches the taxon composition of Moore et al. (2010). Numbers above branches are bootstrap percentages.



0.04