

Majority rule

Soltis, Douglas E., Stephen A. Smith, Nico Cellinese, Kenneth J. Wurdack, David C. Tank, Samuel F. Brockington, Nancy F. Refulio-Rodríguez, Jay B. Walker, Michael J. Moore, Barbara S. Carlswald, Charles D. Bell, Maribeth Latvis, Sunny Crawley, Chelsea Black, Diaga Diouf, Zhenxiang Xi, Catherine A. Rushworth, Matthew A. Gitzendanner, Kenneth J. Sysma, Yin-Long Qiu, Khidir W. Hilu, Charles C. Davis, Michael J. Sanderson, Reed S. Beaman, Richard G. Olmstead, Walter S. Judd, Michael J. Donoghue, and Pamela S. Soltis. Angiosperm phylogeny: 17 genes, 640 taxa. *American Journal of Botany* 98(4): 704-730.

Appendix S22. The maximum parsimony majority-rule consensus resulting from the 17-gene analysis with mtDNA removed for *Polyosma*. Bootstrap values (500 replicates) are given above branches, majority-rule consensus values below branches.

