

Appendix S9. The PCoA 1 and 2 distribution of 1000 PP BEAST trees from *Salvia* nrDNA and the 10 identified grove tree sets (and the MCC tree from Figure 3 and Appendix Figure S8) from Treespace analysis (Kendall, M., and C. Colijn. 2016. *Mapping phylogenetic trees to reveal distinct patterns of evolution. Molecular Biology and Evolution* 33: 2735–2743).

