

# TAXON

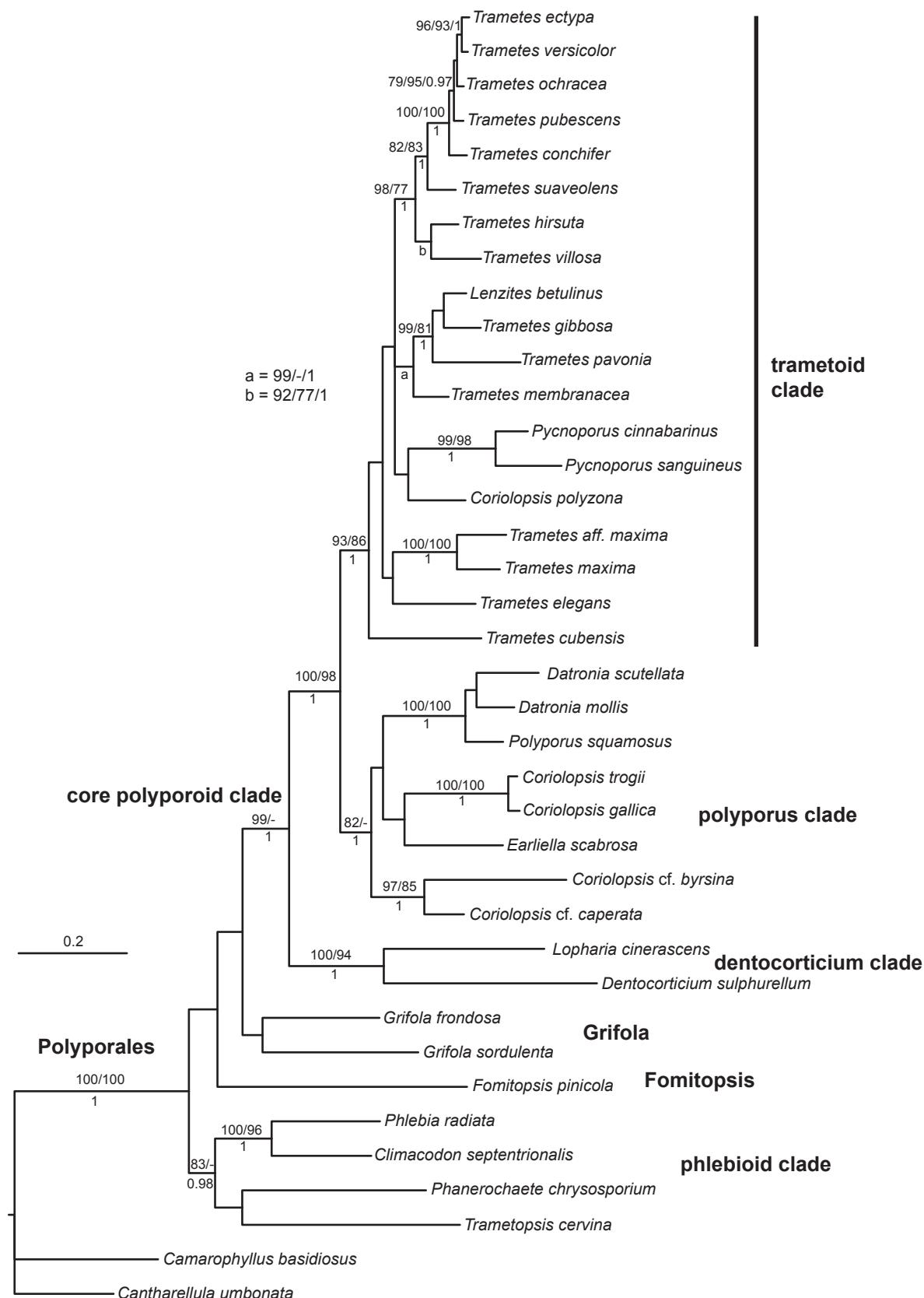
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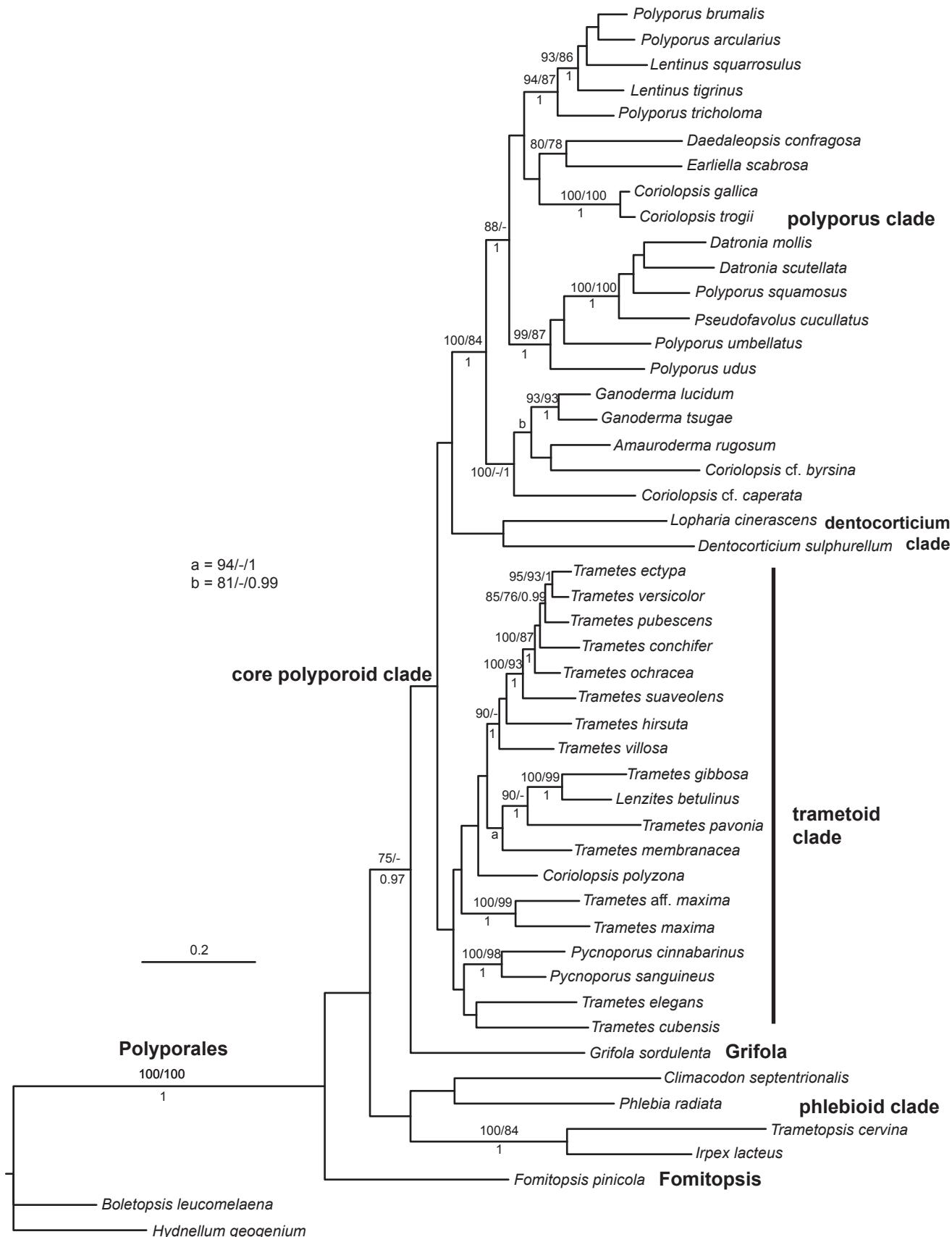
## **Phylogenetic classification of *Trametes* (Basidiomycota, Polyporales) based on a five-marker dataset**

**Alfredo Justo & David S. Hibbett**

***Taxon* 60: 1567–1583**



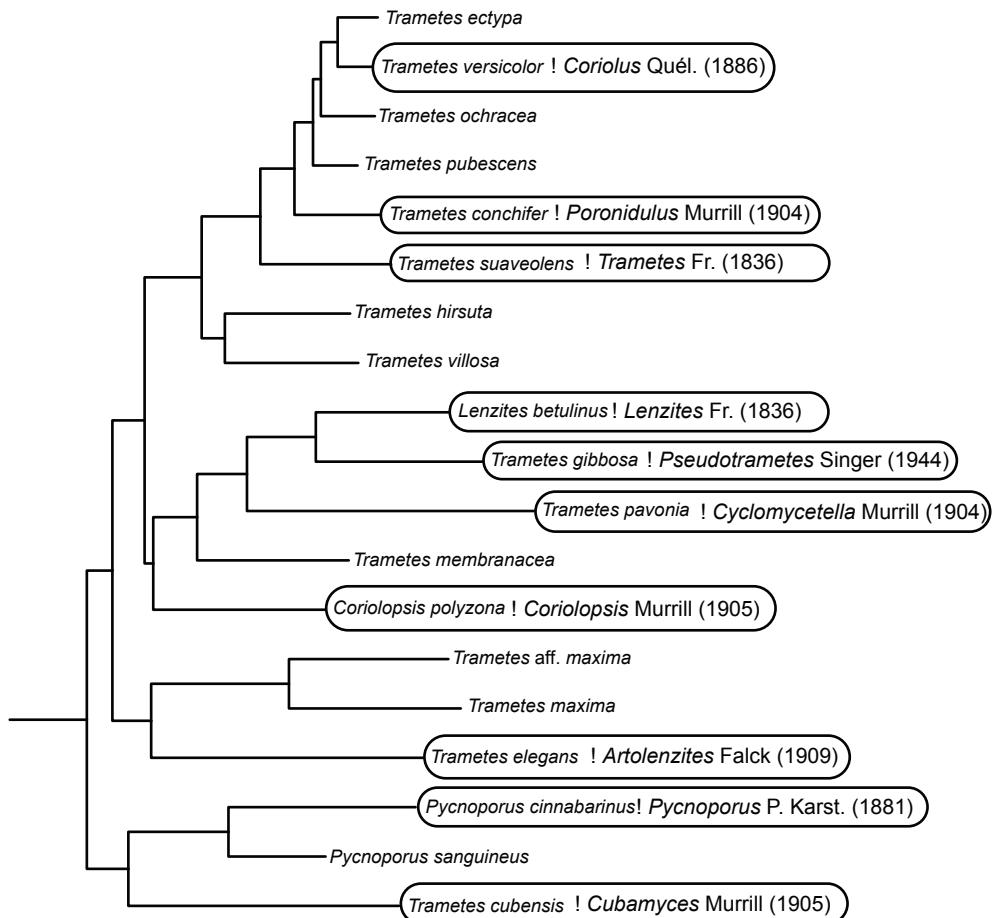
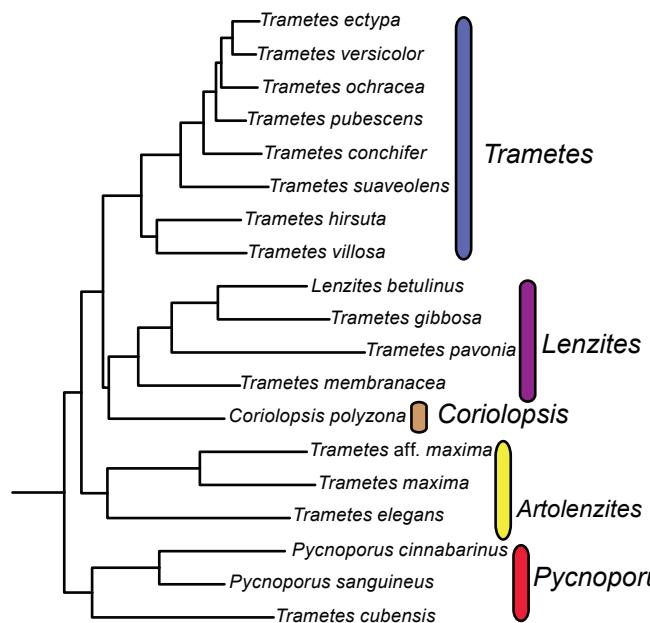
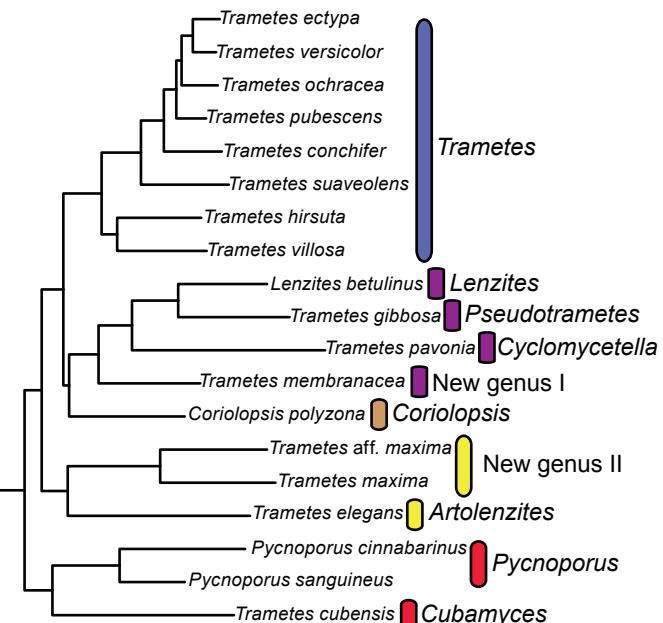
**Fig. S1.** Fifty percent majority-rule consensus tree from the BY analysis of the RPB1 dataset of the Polyporales. BS values from the ML and MP analyses  $\geq 75$  (ML/MP) and PP  $\geq 0.95$  from the BY analysis for nodes supported in at least two of the analyses are given on or below the branches.



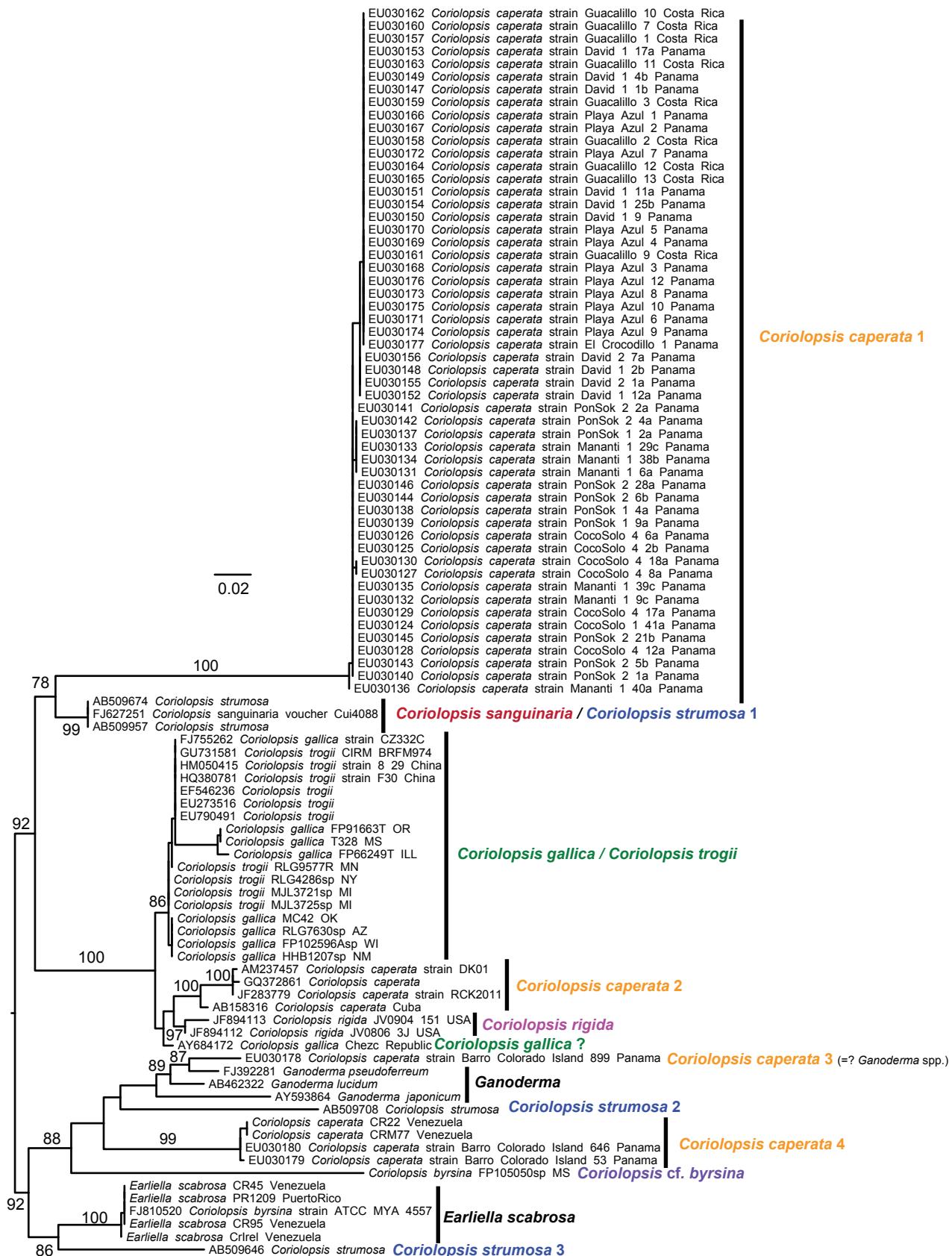
**Fig. S2.** Fifty percent majority-rule consensus tree from the BY analysis of the RPB2 dataset of the Polyporales. BS values from the ML and MP analyses  $\geq 75$  (ML/MP) and PP  $\geq 0.95$  from the BY analysis for nodes supported in at least two of the analyses are given on or below the branches.



**Fig. S3.** Fifty percent majority-rule consensus tree from the BY analysis of the TEF1 dataset of the Polyporales. BS values from the ML and MP analyses  $\geq 75$  (ML/MP) and PP  $\geq 0.95$  from the BY analysis for nodes supported in at least two of the analyses are given on or below the branches.

**The Five genera option****The Ten genera option**

**Fig. S4.** Nomenclatural overview of the trametoid clade based on the results from the five-marker dataset of the Polyporales and the two alternative taxonomic scenarios.



**Fig. S5.** Best tree from the ML analysis of the ITS dataset of *Coriolopsis*. The midpoint rooting option was used. BS values from the ML analysis are given on or below the branches.