

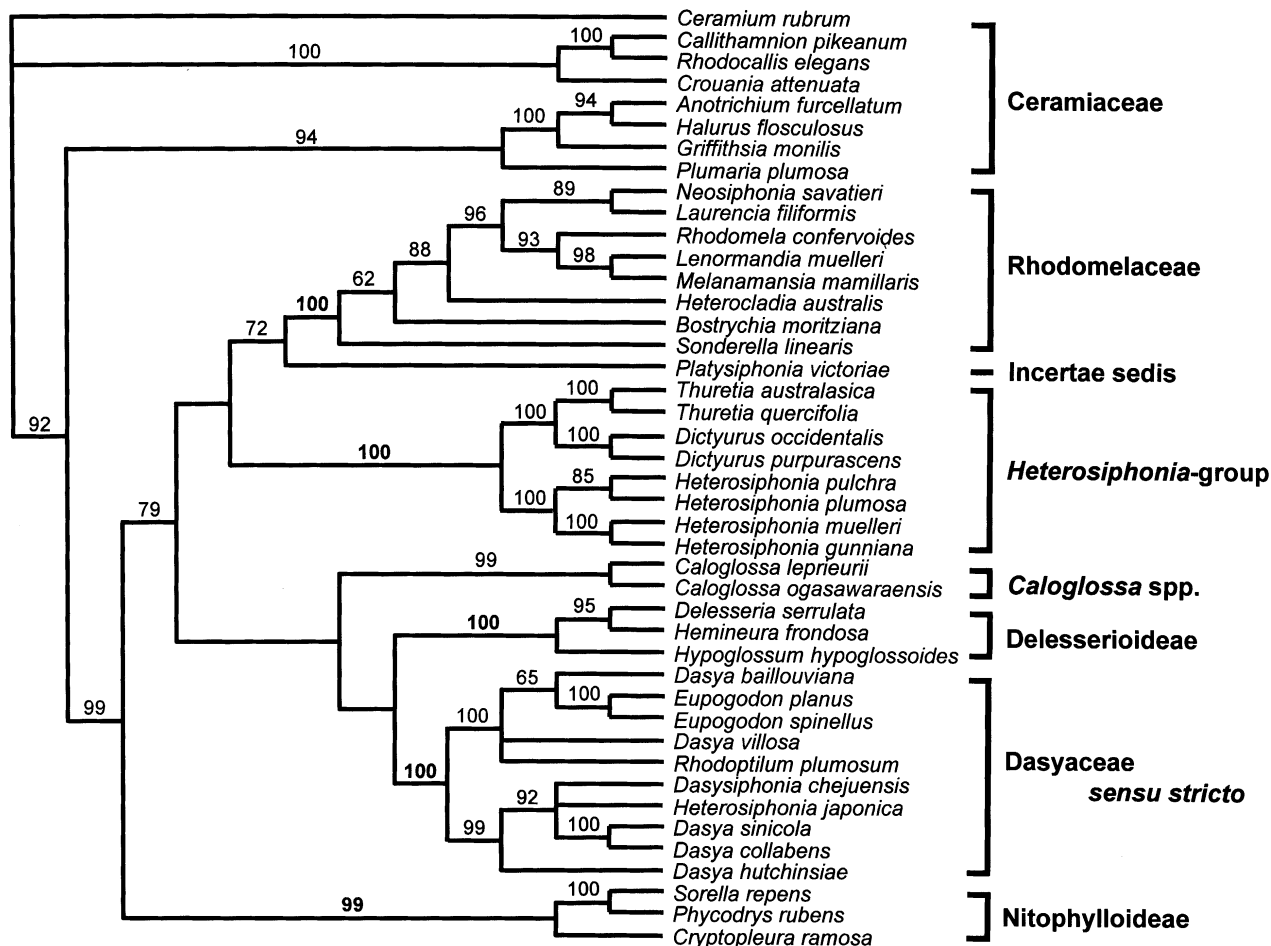


Fig. 4. Strict consensus of four equally parsimonious solutions for combined anatomical and SSU data. Numbers above the branches indicate bootstrap values (% of 2000 replicates). Bold numbers are bootstrap support for the five major lineages of the Dasyaceae/Delesseriaceae/Rhodomelaceae clade resolved. Branches lacking values received less than 50% support.