

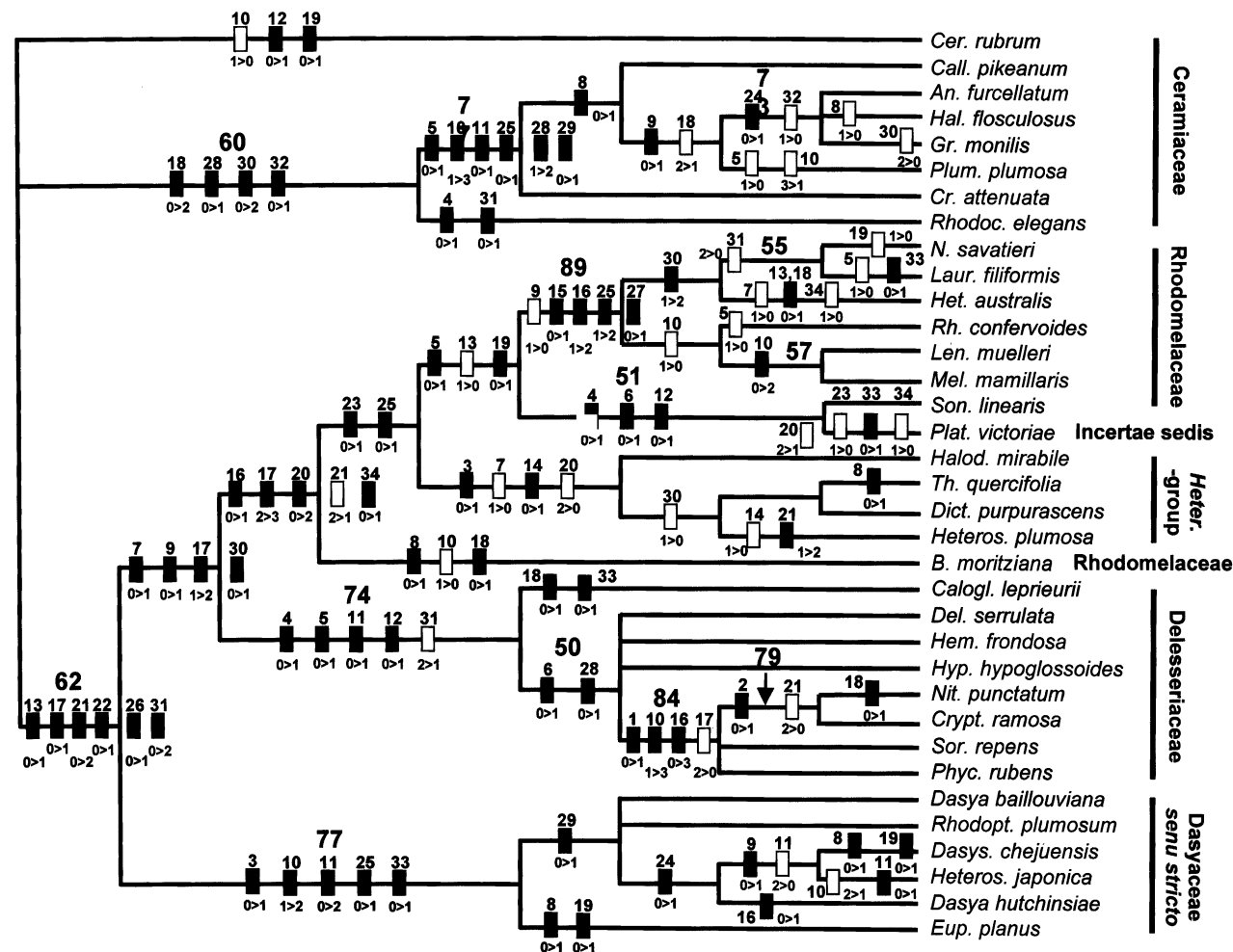


Fig. 3. Cladistic analyses for anatomical data. One of 48 equally parsimonious solutions for anatomical data with 19 reproductive characters weighted 1:3:1 against 15 vegetative characters (weighted parsimony). Bold numbers above the branches indicate bootstrap values (% of 1000 replicates). Branches lacking values received less than 50% support. Numbers above and below the filled and open bars correspond to anatomical characters and character-state changes (Tables 3 and 4), respectively. Filled bars, (syn)apomorphies; open bars, reversals.