Figure S3. Results of Bayesian species tree analysis of the **complete character** matrix. Shown is the maximum clade credibility tree derived from a single MCMC analysis (5·10^8 generations, sampled every 1·10^{-4}, burnin = 5·10^7 generations) using *BEAST v1.6.1 (Drummond et al. 2006, 2007). Nodes are sequentially labelled to match support/conflict graphs in figure S4, and branch lengths are arbitrary.