



Fig. 1. Tree of *Cryptigata* and *Acanthopneuste* sensu Ticehurst (1938) and Watson et al. (1986), and representatives of all subgenera in *Phylloscopus* recognized by Watson et al., and all main clades in *Phylloscopus* and *Seicercus*, as revealed by previous studies. The tree was estimated by Bayesian analysis of combined cytochrome *b*, 12S, and myoglobin intron II sequences (2172 bp) divided into two partitions analysed under different models (GTR +  $\Gamma$  + I for the mitochondrial genes, GTR +  $\Gamma$  for myoglobin intron II). The bars on the left show *Cryptigata* (black) and *Acanthopneuste* (grey), as circumscribed by Ticehurst (1938) and Watson et al. (1986); asterisks mark the taxa that were not described at the time. The bars on the right refer to clades discussed in the text. All taxa, but only one sample per taxon, were included in this analysis. The names of the taxa in clade A are those of previous authors (e.g., Watson et al., 1986). Nodal support shown in Fig. 2.