



FIG. 1. Phylogenetic relationships for 36 bird species based on DNA hybridization data from Sibley and Ahlquist (1990). The length of vertical branches was scaled according to the genetic distance between the species; the horizontal axis is arbitrary. The position of the nodes marked with triangles had to be assumed arbitrarily, due to the lack of sufficient data. The nodes marked with squares were omitted in the calculations of phylogenetic contrasts, in order to remove the influence of the differences in chicks' modes of development.