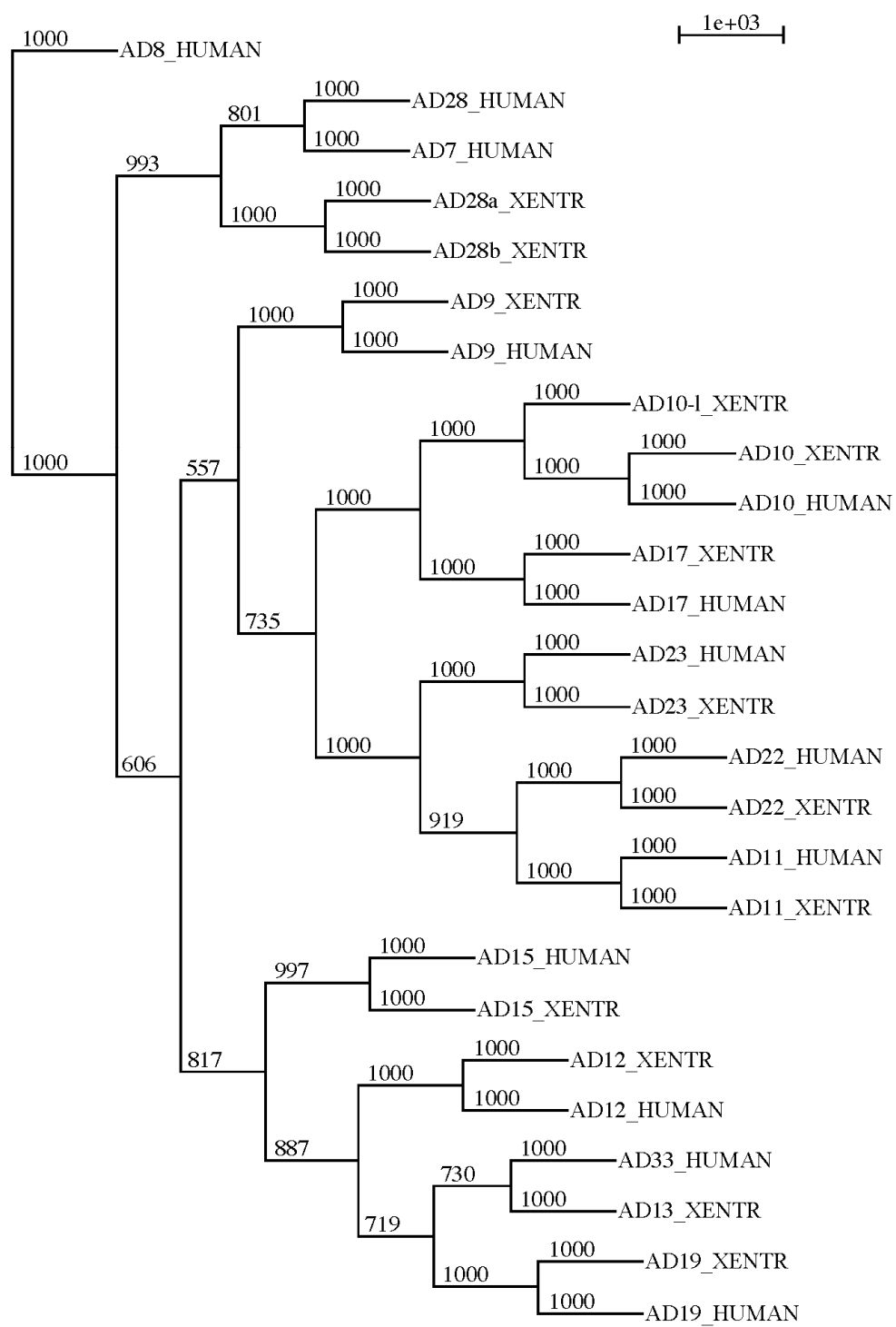
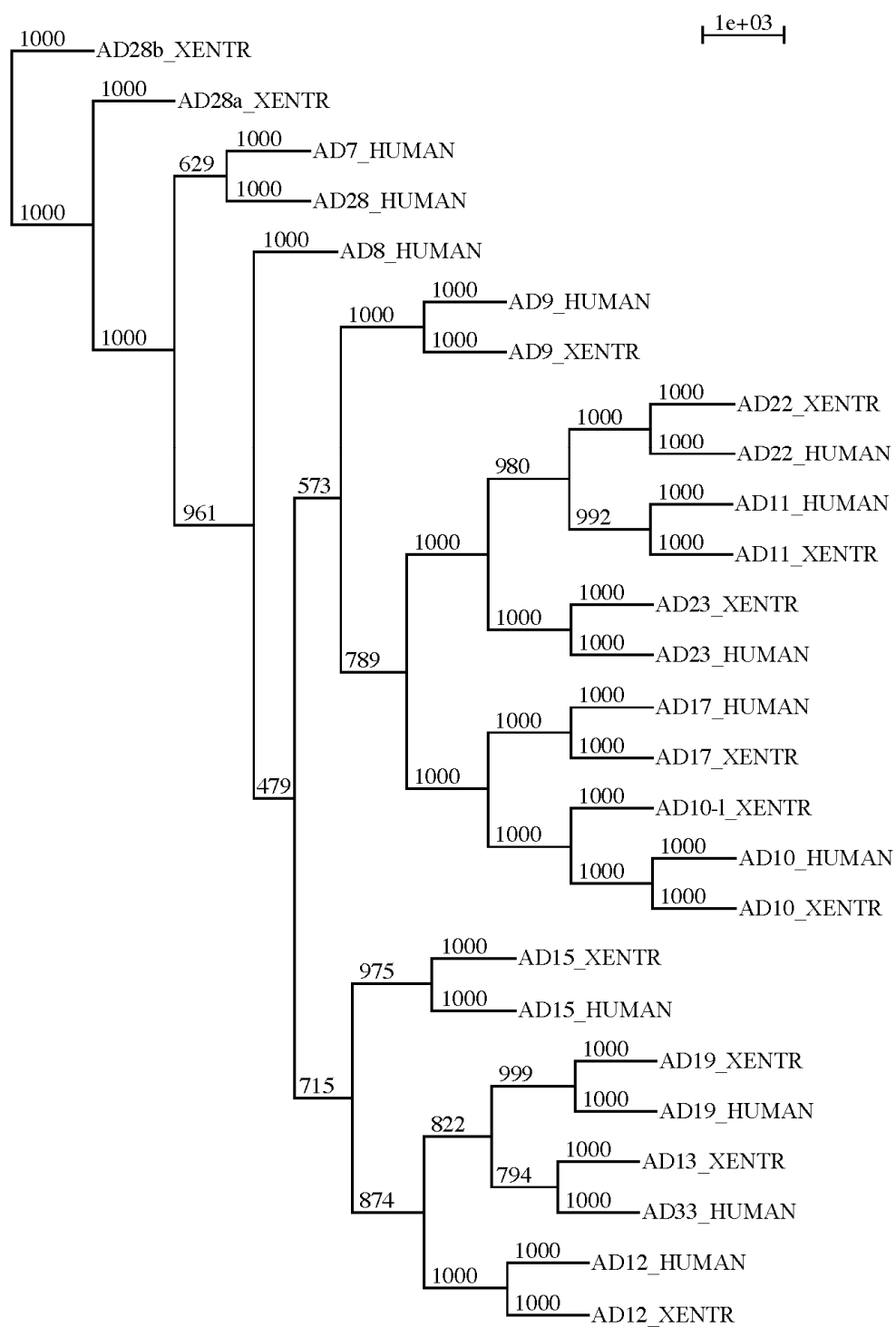


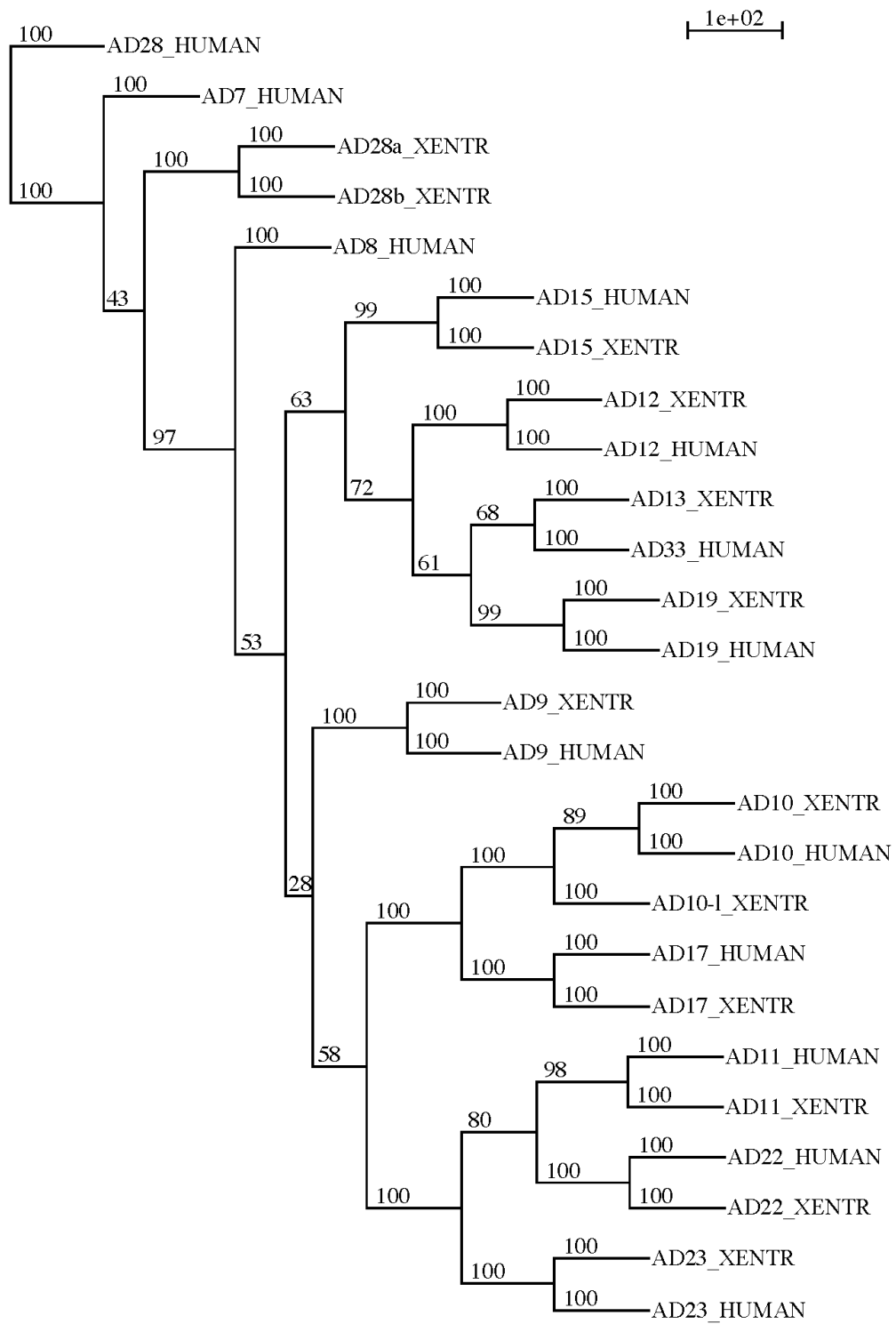
A



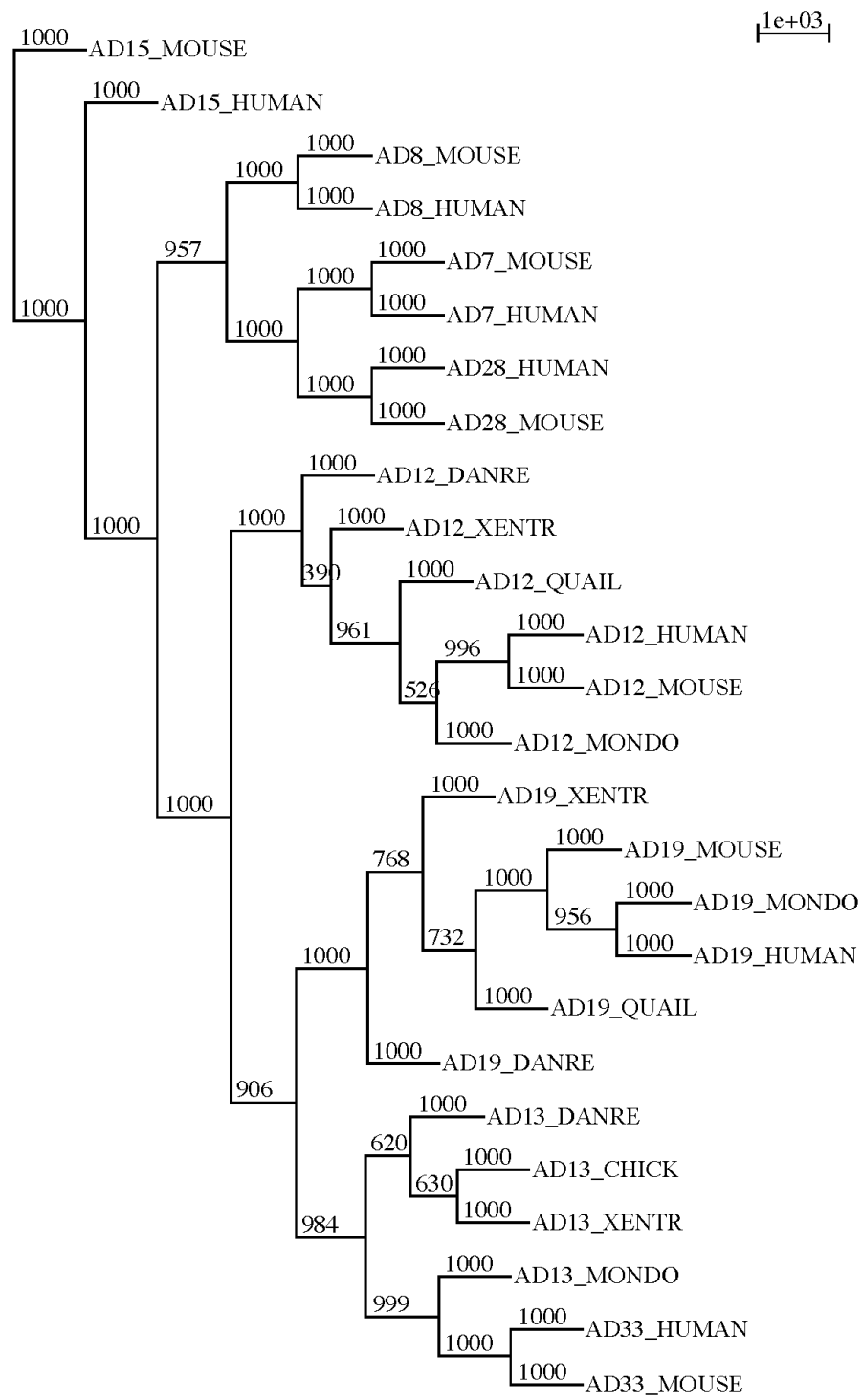
B



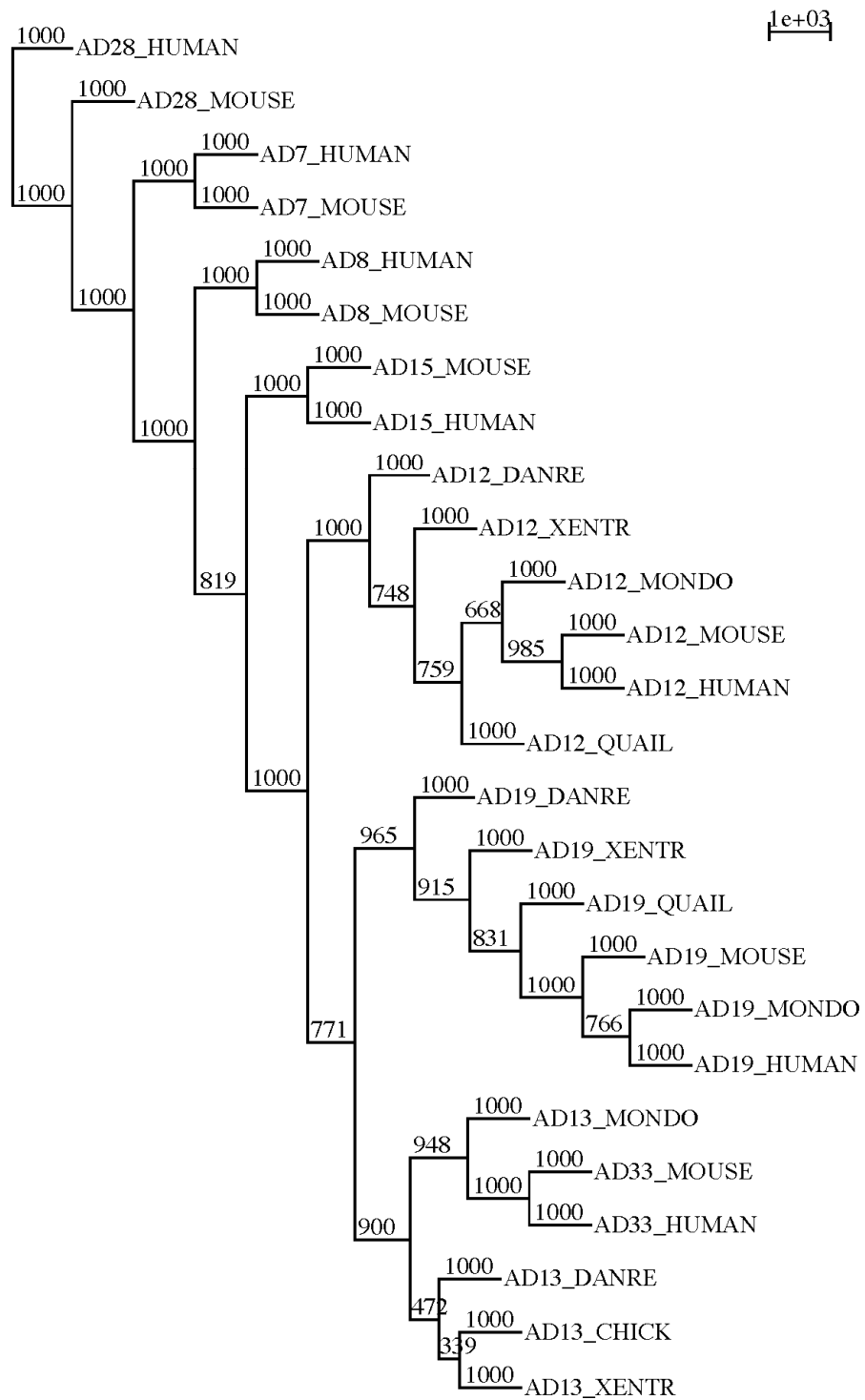
C



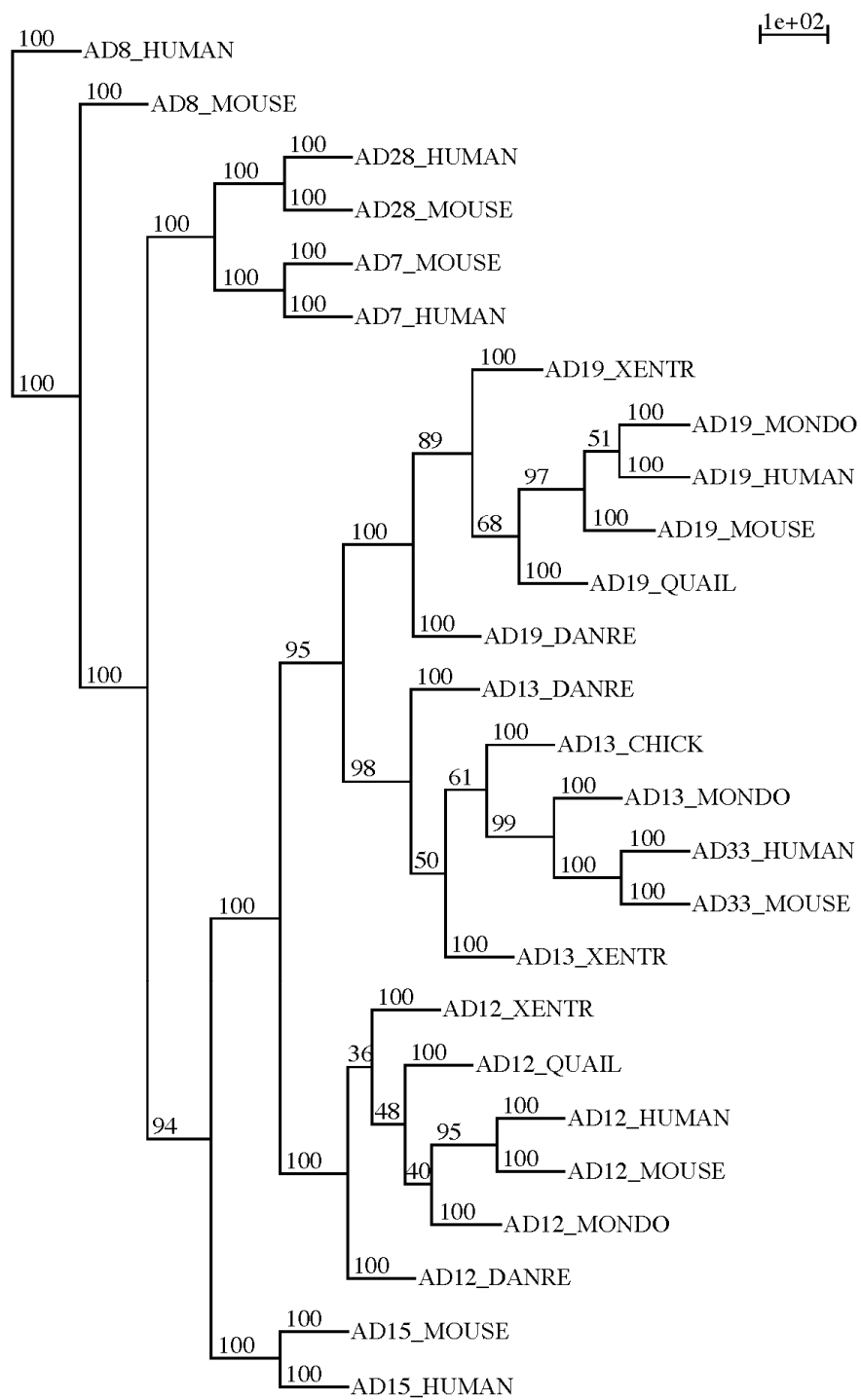
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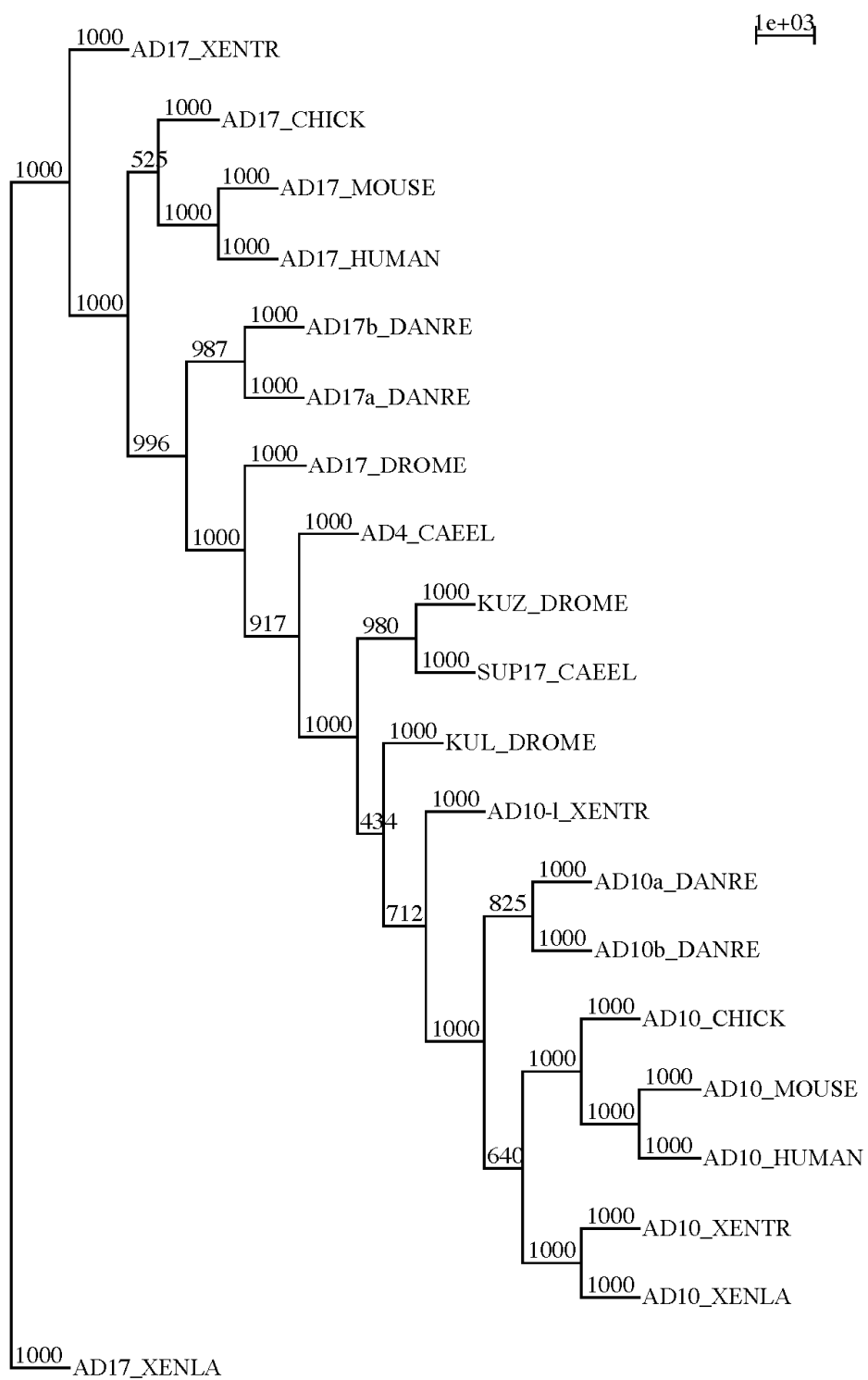
E



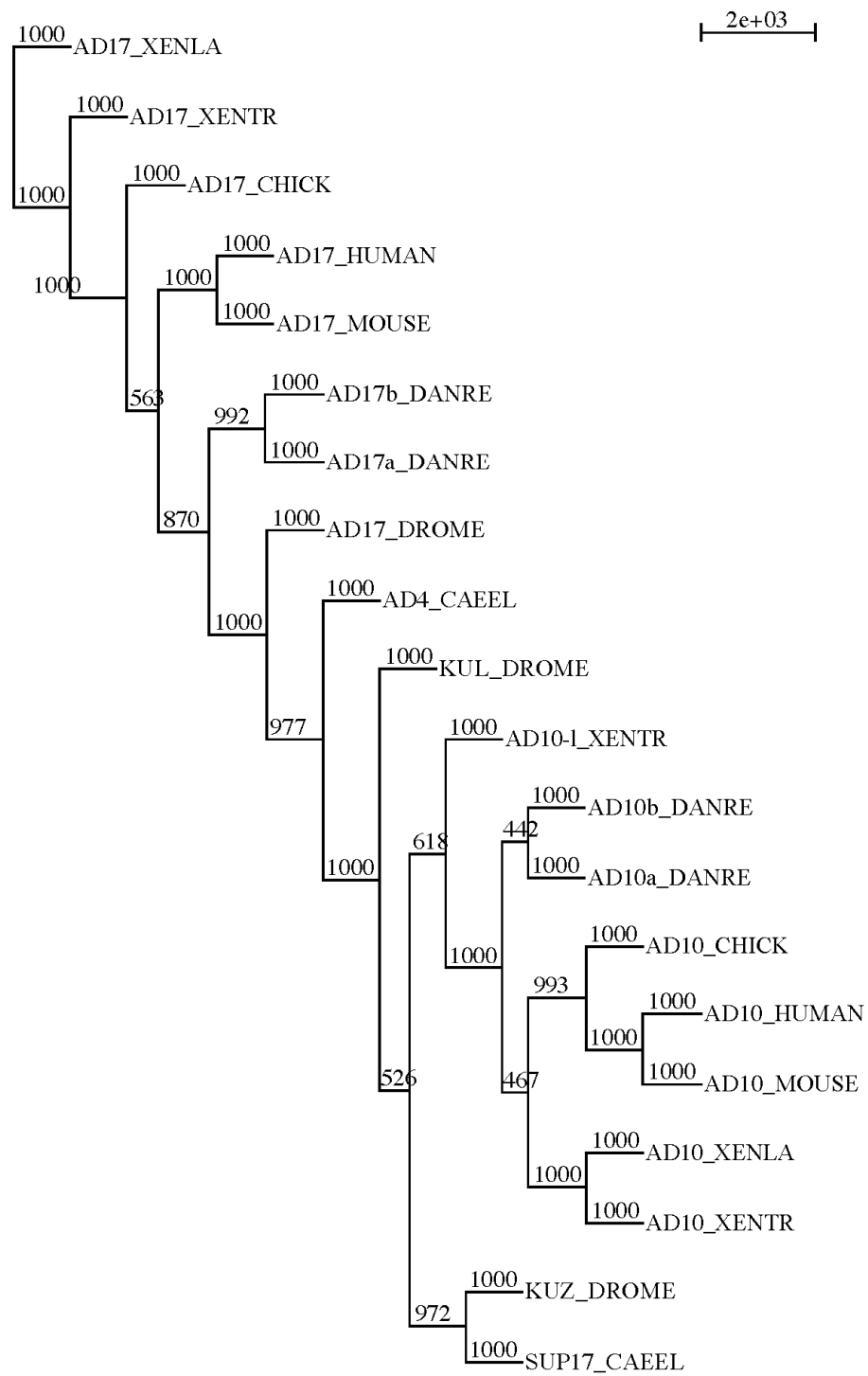
F



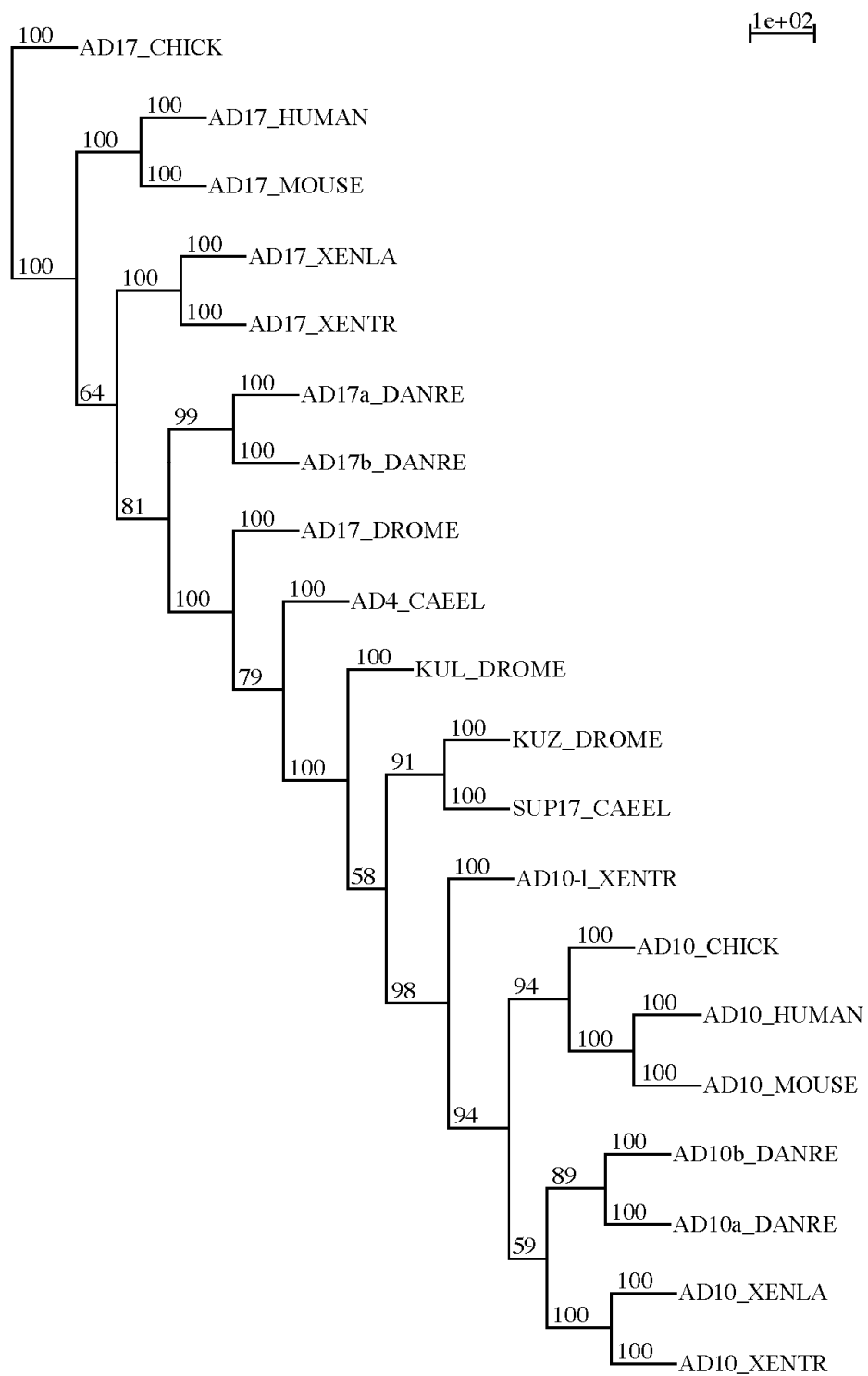
G



H



I



Additional File 6. Phylogenetic trees generated using alternative models. Trees shown in Figs. 1A, 3A and 8, respectively, were rebuilt using Phylip Neighbor (A, D, G), Protpars (B, E, H), and Proml (C, F, I) analyses. Bootstrap values (out of 1000 for Phylip Neighbor and Protpars, and out of 100 for Proml) are displayed on the branches.