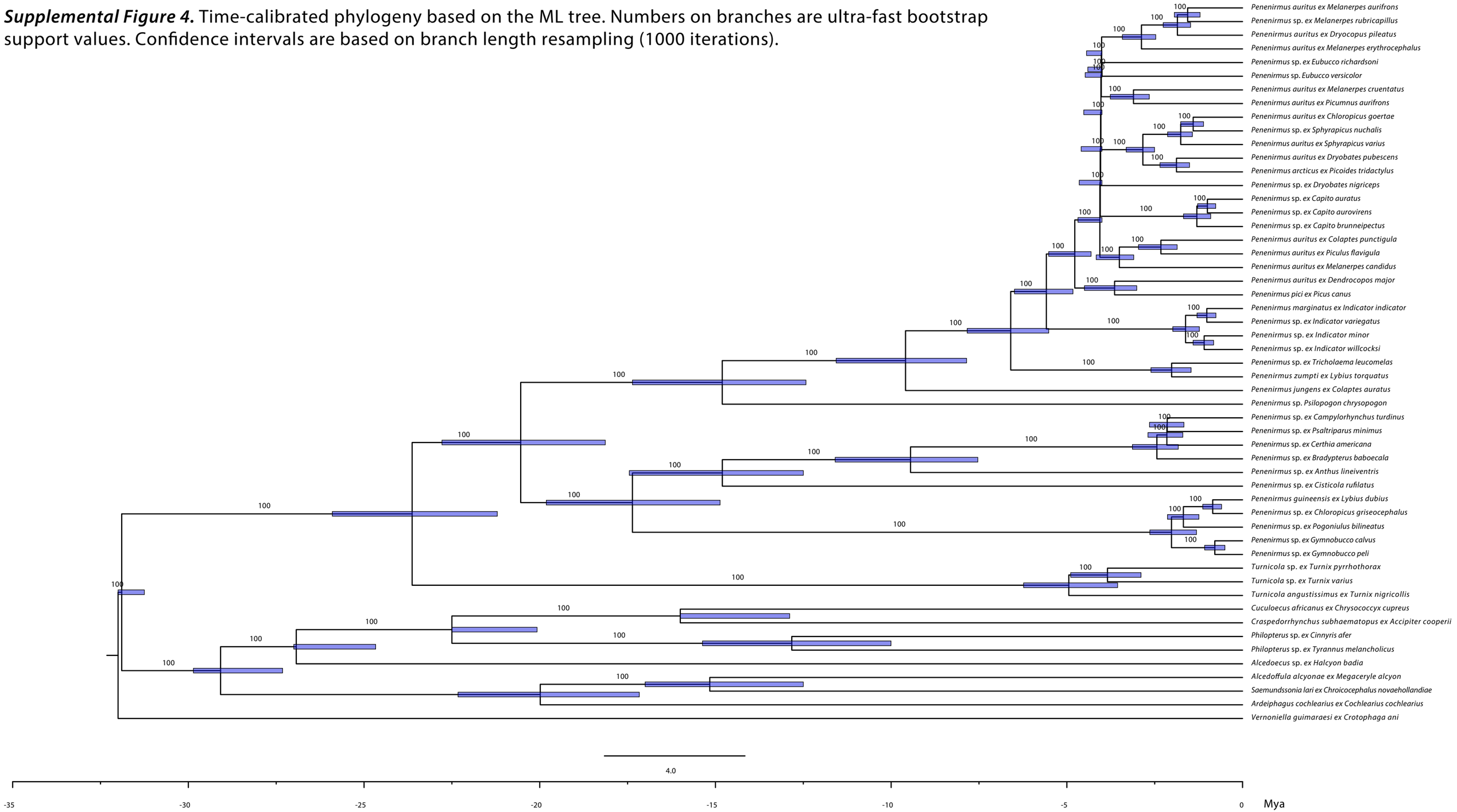Supplemental Figure 4. Time-calibrated phylogeny based on the ML tree. Numbers on branches are ultra-fast bootstrap support values. Confidence intervals are based on branch length resampling (1000 iterations).