

A

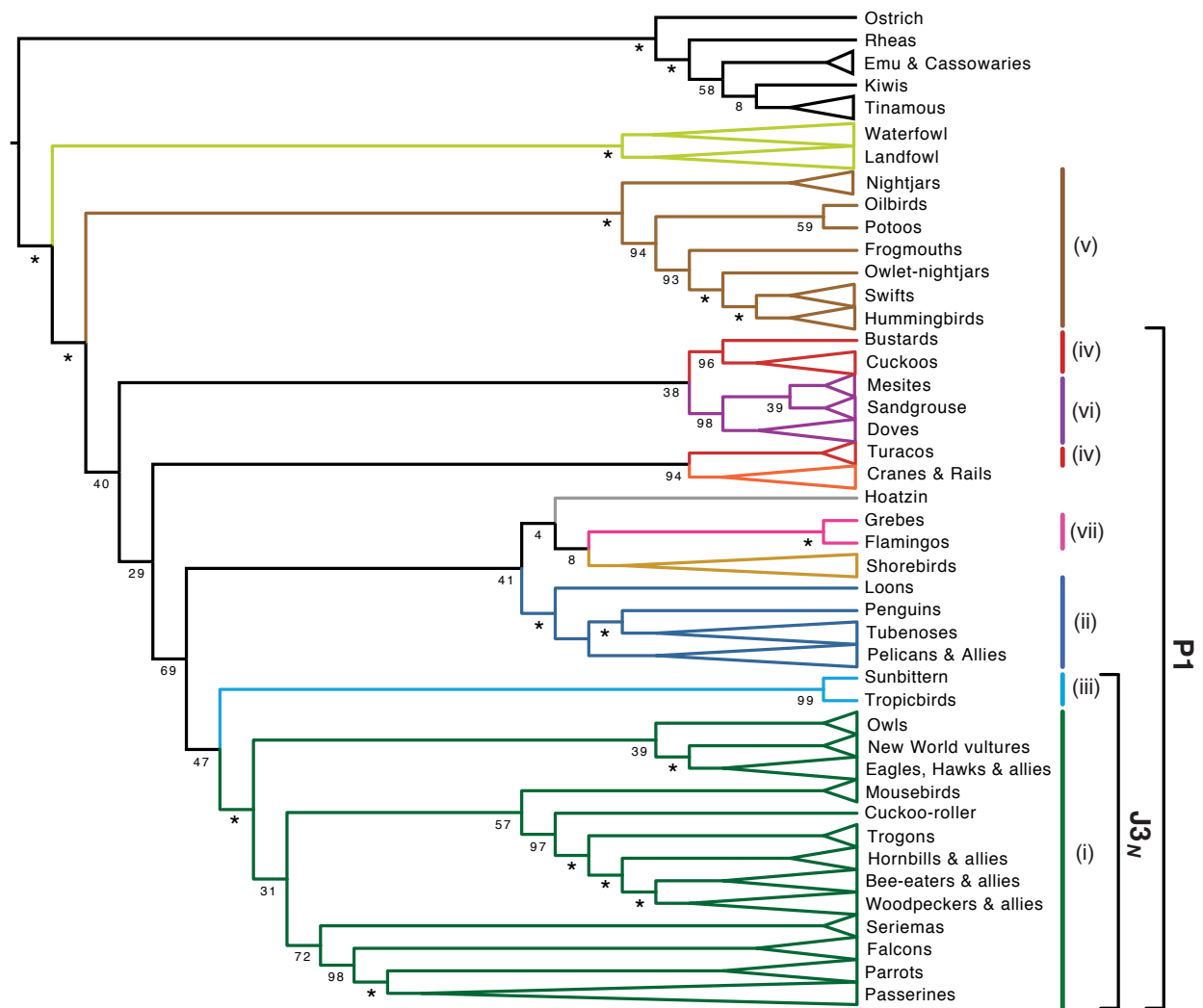


FIGURE S3. The ‘Prum noJAR” ML tree for 198 birds using 104 loci. These analyses used the Prum loci with coding data after excluding the loci that were also included in the Jarvis TENT data. (A) Cladogram based on a RAxML analysis with the maximum number of partitions. The length of the triangles representing each order is proportional to the number of taxa sampled for that taxon. If there is no triangle a single taxon was sampled for the order. Bootstrap values are present below each branch (asterisks indicate 100% bootstrap support). (B; next page) Unlabeled phylogram based on a RAxML analysis with the maximum number of partitions. Colors are identical to those used in part A (and the other phylogeny figures in this Point of View). This phylogeny, along with three additional estimates of phylogeny for the Prum noJAR data (unpartitioned RAxML and partitioned and unpartitioned IQ-TREE), is available as a Nexus format treefile in Supplementary Data S10.

B

