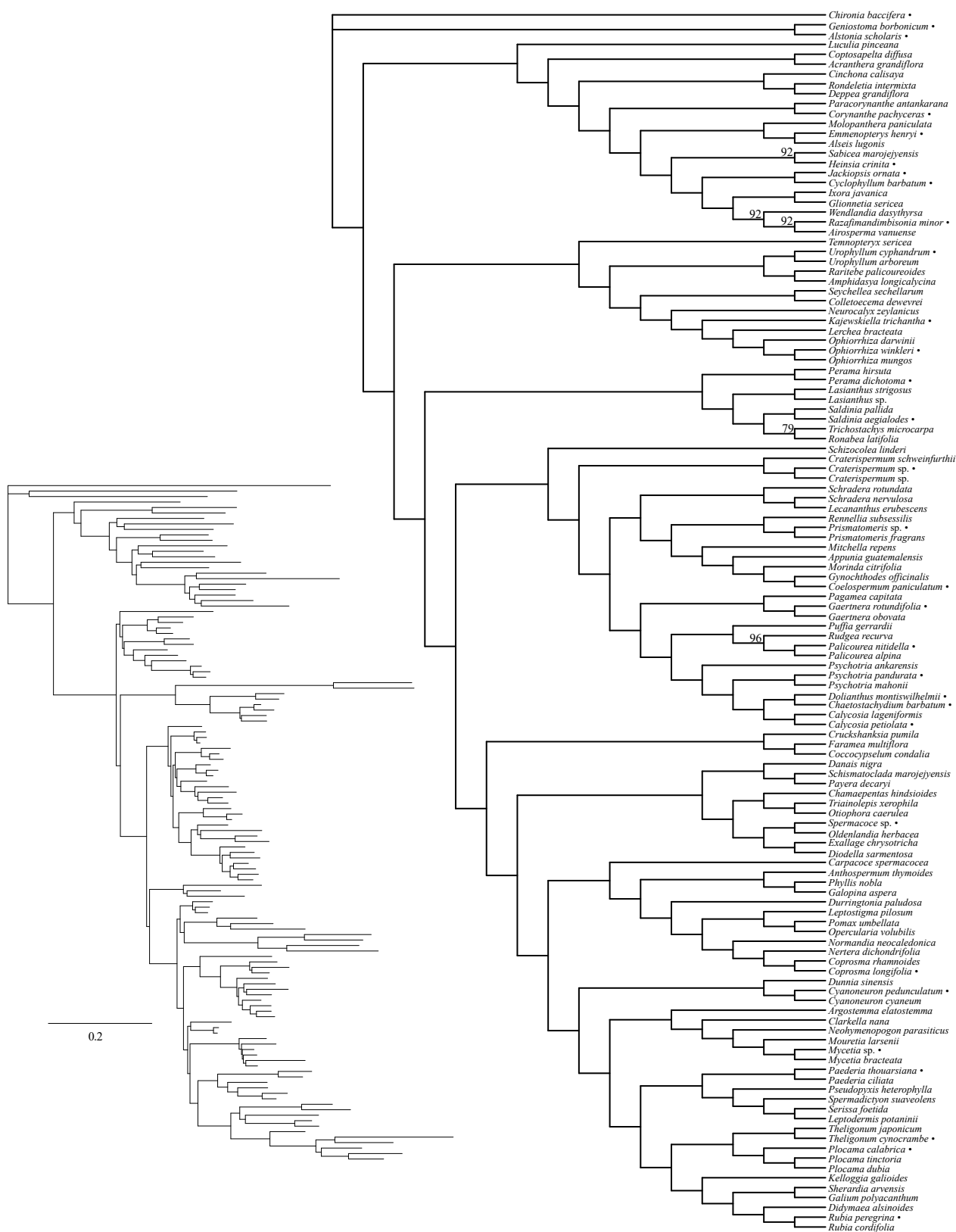




Supplementary Figure 4. Concatenation-based tree estimated using IQ-TREE on the paralog-filtered supercontig dataset. Numbers above branches denote ultrafast bootstrap (BS) support values. Only support values smaller than 100% are shown. Bullets after species names indicate samples downloaded from ENA. Inset shows branch lengths in number of substitutions per site.