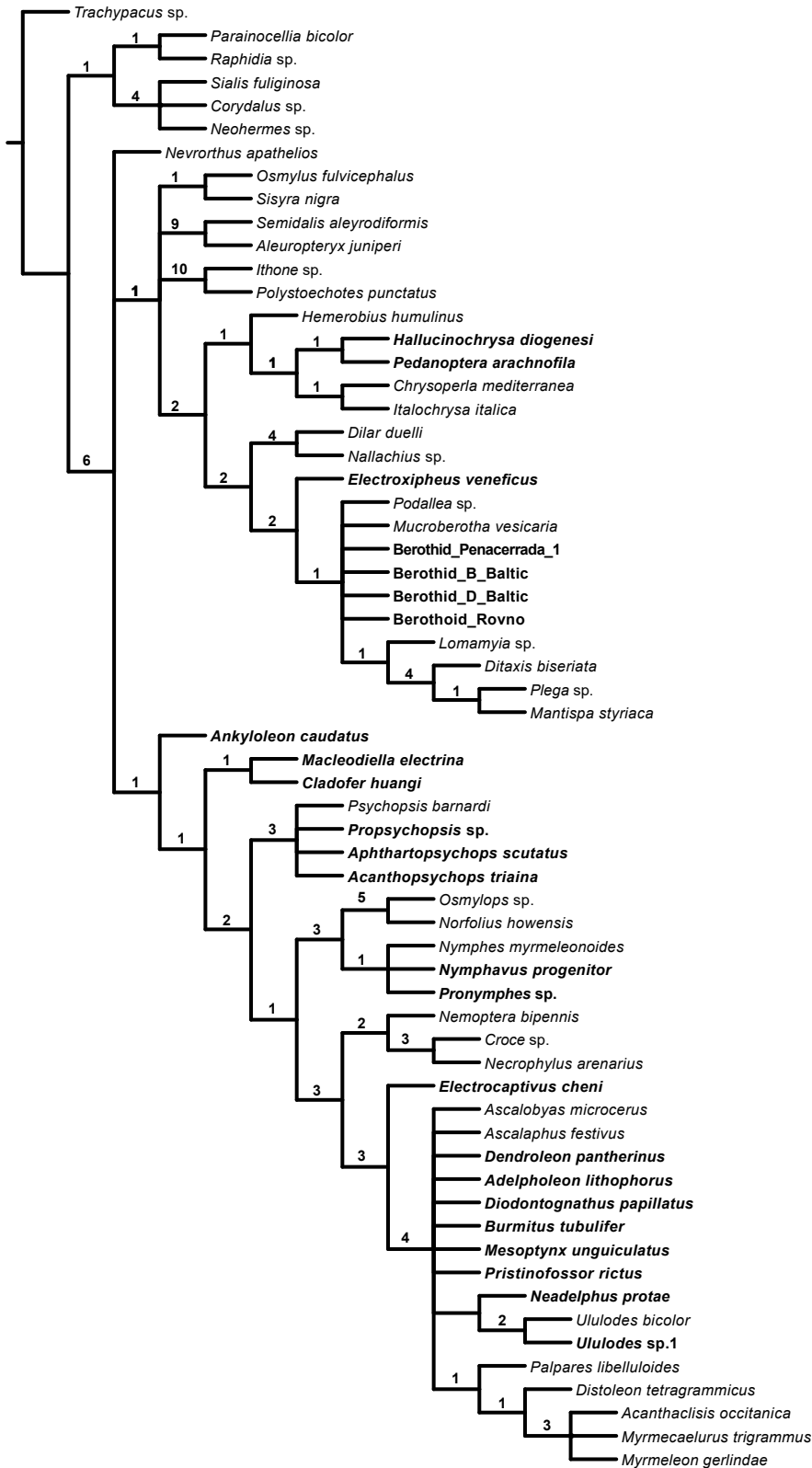


Supplementary Information Fig. S1. Strict consensus cladogram from 936 most parsimonious trees resulting from the maximum parsimony analysis of the morphological dataset under equal weights (CI: 0.535; RI: 0.854). Numbers above branches represent Bremer support values.



Supplementary Information Fig. S2. Cladogram obtained from the maximum parsimony analysis of the morphological dataset under implied weight (IW) with optimized character state changes. Black and white squares indicate unique and homoplasious apomorphies, respectively.

