

KERR, PETER H. (2010) Phylogeny and classification of Rhagionidae, with implications for Tabanomorpha (Diptera: Brachycera). Zootaxa, 2592, 1–133.

On page 64, the labels of **FIGURE 196** are missing. Below is the corrected **FIGURE 196**.

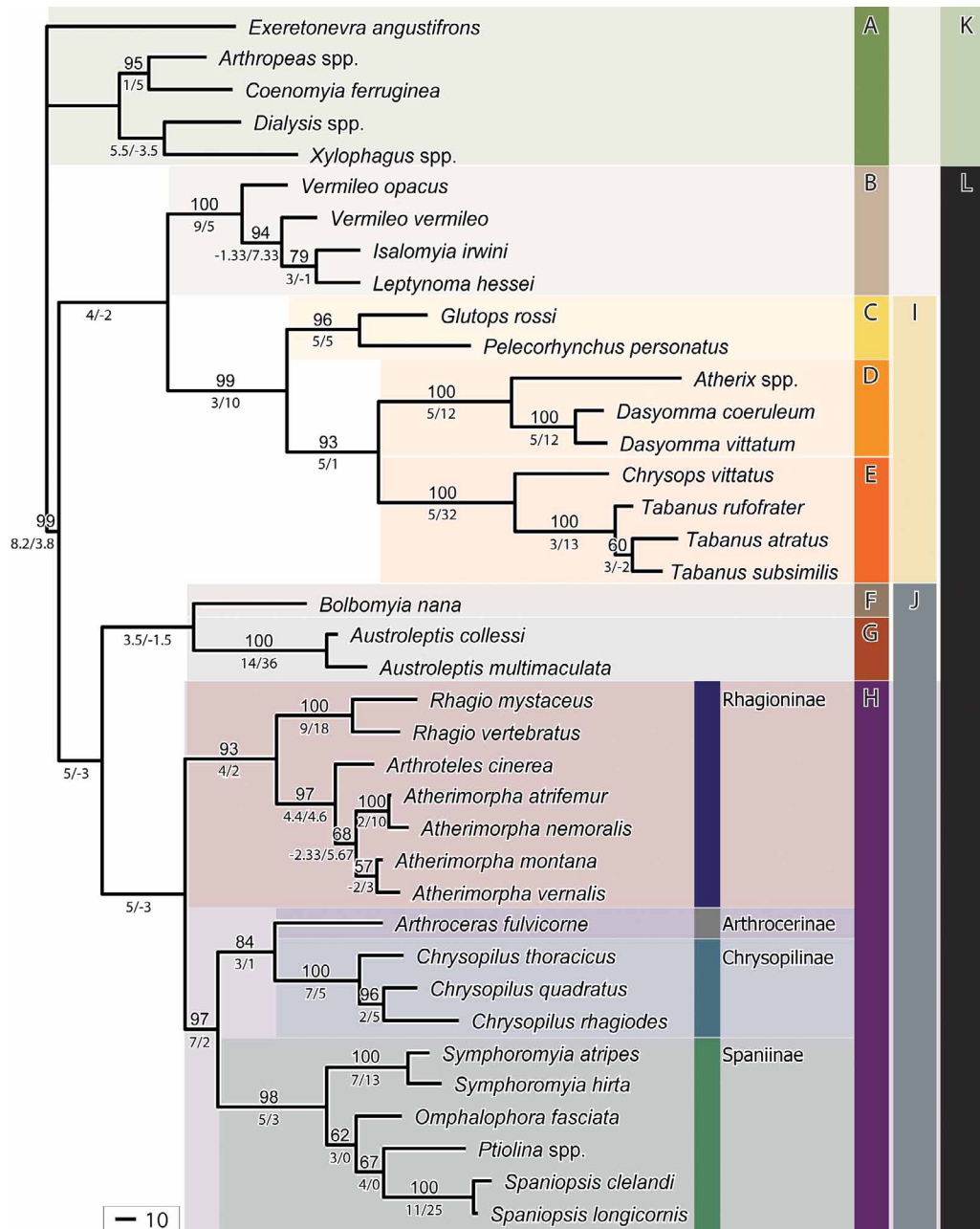


FIGURE 196. The most parsimonious tree of the combined data set, composed of morphological and molecular characters. Bootstrap values greater than 50% are above supported branches (TBR, nreps = 1000 / addseq reps = 5). Partitioned Bremer Support values are below supported branch (morphology partition/molecular partition); the sum of these values equals the overall decay (BS). The Bayesian analysis of the same data set yields the same topology and similar branch lengths for the ingroup, with posterior probabilities of 95% or higher for Tabanomorpha and the early divergences within this group. *Classification along right margin*; A = Xylophagidae, B = Vermileonidae, C = Pelecorhynchidae, D = Athericidae, E = Tabanidae, F = Bolbomyiidae, G = Austroleptidae, H = Rhagionidae, I = Tabanoidea, J = Rhagionoidea, K = Xylophagomorpha, L = Tabanomorpha.