

Supporting Information

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Bray et al. 2016: The molecular phylogeny of the digenean family Opecoelidae Ozaki, 1925 and the value of morphological characters, with the erection of a new subfamily. Folia Parasitol. 63: 013, doi 10.14411/fp.2016.013

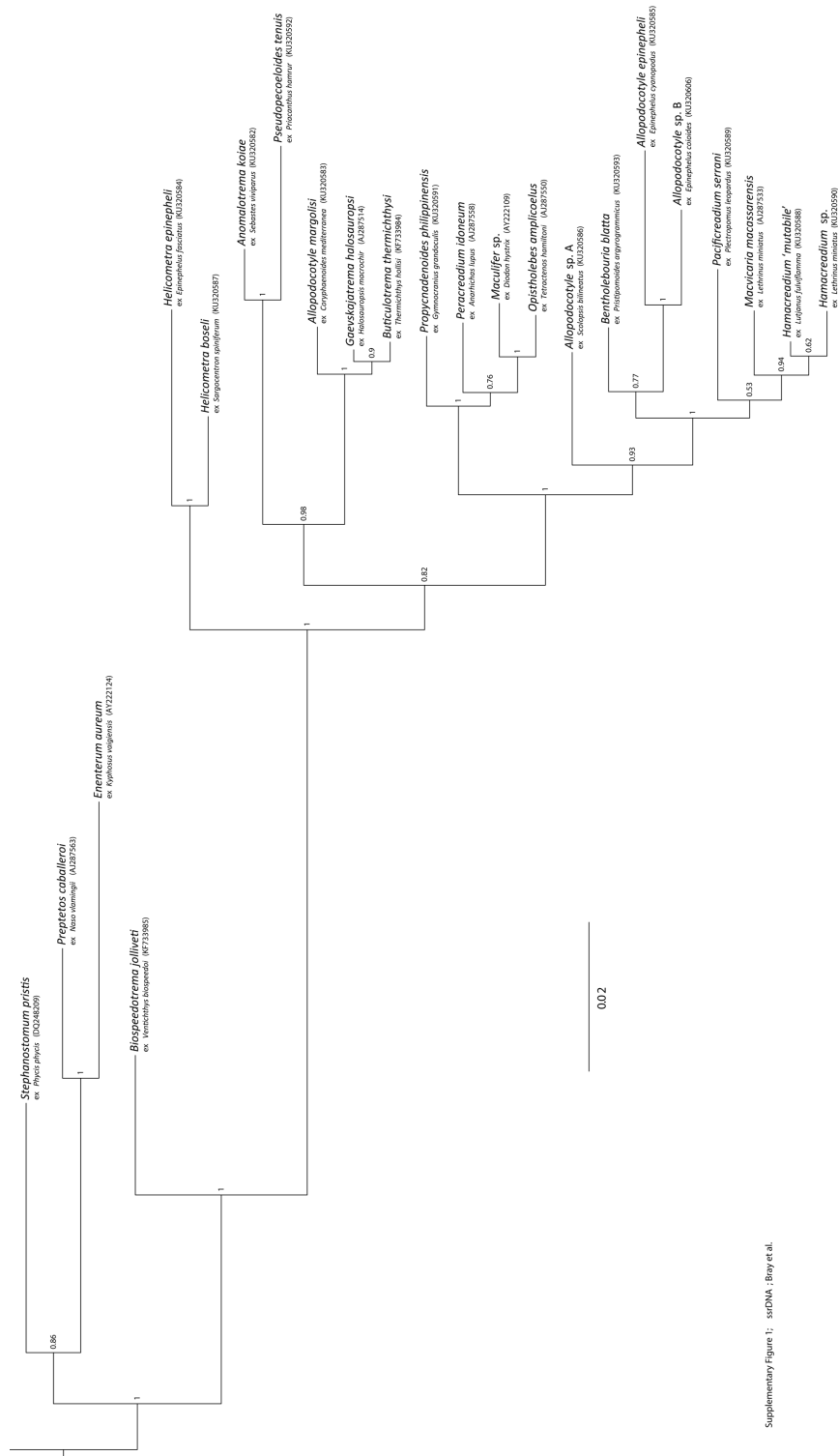
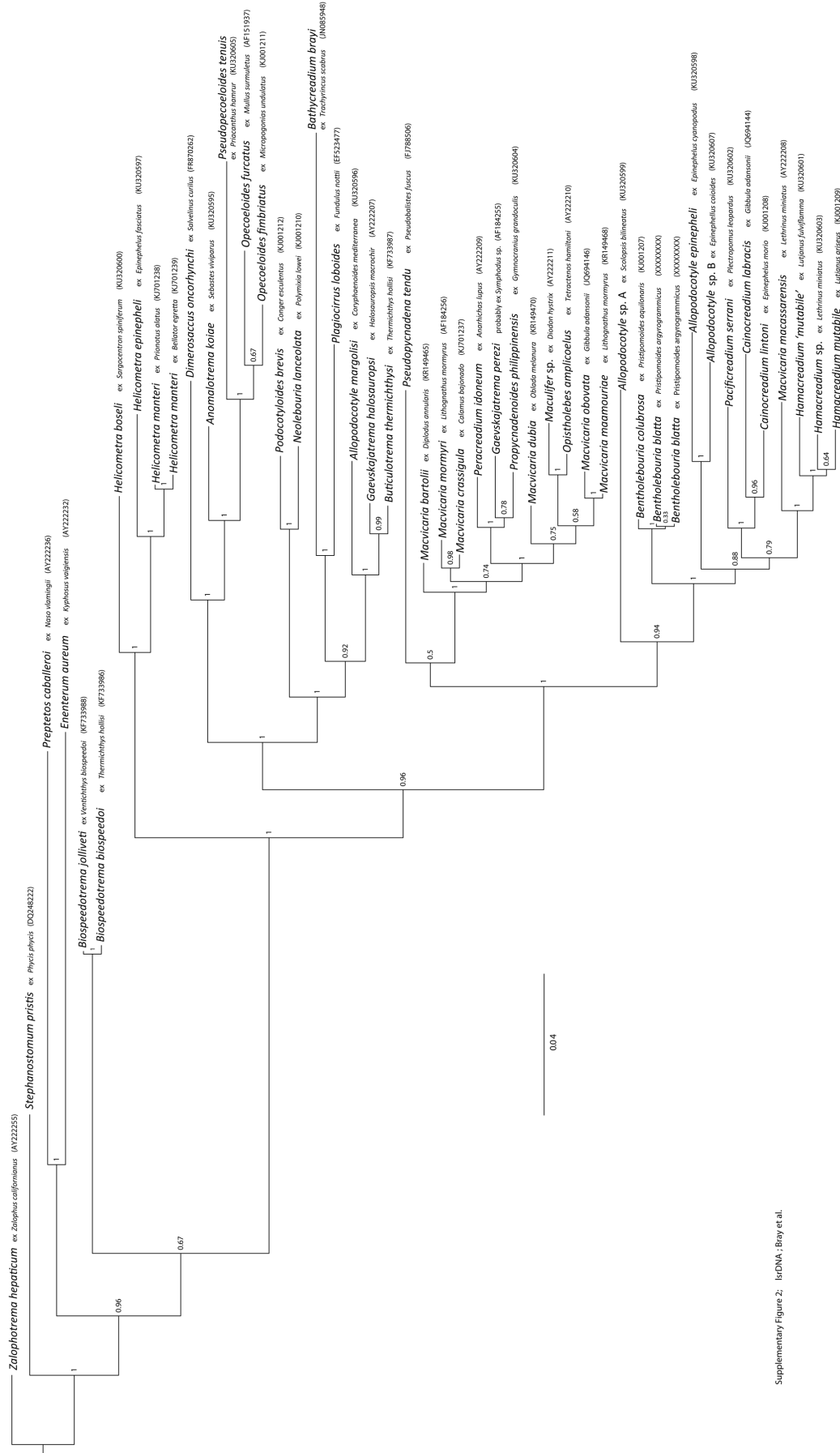


Fig. S1. Bayesian analysis of the *ssrDNA* data constructed using MrBayes v3.2 under the GTR + I + G model; 10 000 000 generations, 6 000 000 generations 'burn-in'. The branch length scale indicates number of substitutions per site.



Supplementary Figure 2: *IsrDNA*: Bray et al.

Fig. S2. Bayesian analysis of the *IsrDNA* data constructed using MrBayes v3.2 under the GTR + I + G model; 10 000 000 generations, 6 000 000 generations ‘burn-in’. The branch length scale indicates number of substitutions per site.