Supplementary material

Transfer of the New Caledonian genus *Boronella* to *Boronia* (Rutaceae) based on analyses of cpDNA and nrDNA

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Fig. S1. Majority-rule consensus tree based on Bayesian inference analysis of combined internal transcribed spacer (ITS) and external transcribed spacer (ETS) datasets, showing mean branch lengths. Numbers above nodes are Bayesian posterior probabilities.
Fig. S2. Majority-rule consensus tree based on Bayesian inference analysis of *trnL–trnF* dataset, showing mean branch lengths. Numbers above nodes are Bayesian posterior probabilities.