

**Supplementary file 3.** Bayesian majority-rule consensus tree of the *Ocotea* complex (Lauraceae) derived from the combined analysis of nuclear ribosomal ITS and plastid *psbA-trnH* sequence data (230 terminals by 1479 aligned sites) showing the placement of the morphologically enigmatic new species *Ocotea bilocellata* within the *Ocotea minarum* group (sensu Trofimov *et al.*, 2019). Posterior probabilities from 0.5 to 1.0 are shown as color gradient from red to black as well as values on the branches.

