

Running head: VARGAS ET AL., RECLASSIFICATION OF *BERTHOLLETIA* (LECYTHIDACEAE)

Reclassification of the *Bertholletia* clade of the Brazil nut family (Lecythidaceae), based on a phylogenetic analysis of plastome and target sequence capture data

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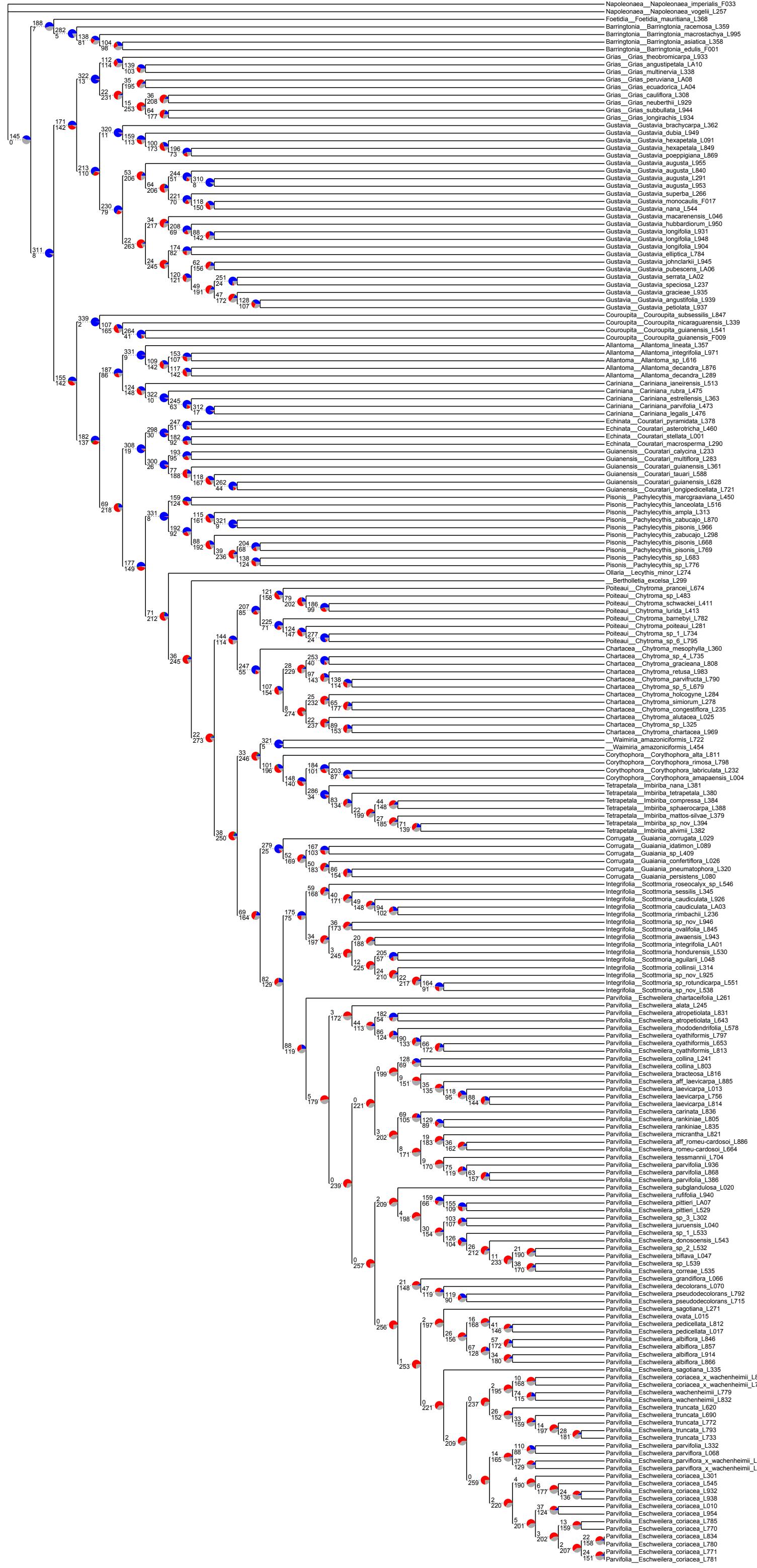
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SUPPLEMENTARY FIGURE 2. Phylogenetic conflict of gene trees with the nuclear tree, generated with phyparts. The proportion of gene trees that are concordant or discordant are represented in blue and red pies respectively. Uninformative gene trees, those with < 70% bootstrap support, are represented in gray. The number of gene trees that are concordant or discordant are also shown above and below individual branches, respectively.



SUPPLEMENTARY FIGURE 3. Plastome tree based on a concatenated matrix of 10 plastome regions. Node labels indicate standard, non-parametric bootstrap support. Branches that agree with the nuclear phylogeny are represented as thick black lines and conflicting branches are lighter gray.

