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Author Correction: Plastid phylogenomics resolves ambiguous relationships within the orchid family and provides a solid timeframe for biogeography and macroevolution

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Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-021-83664-5>, published online 25 March 2021

The original version of this Article contained an error in Figure 1 where the image showing "Paphiopedilum villosum" was incorrectly labelled as "Apostasia shenzhenica".

The original Figure 1 and accompanying legend appear below.

The original Article has been corrected.

Published online: 06 July 2021

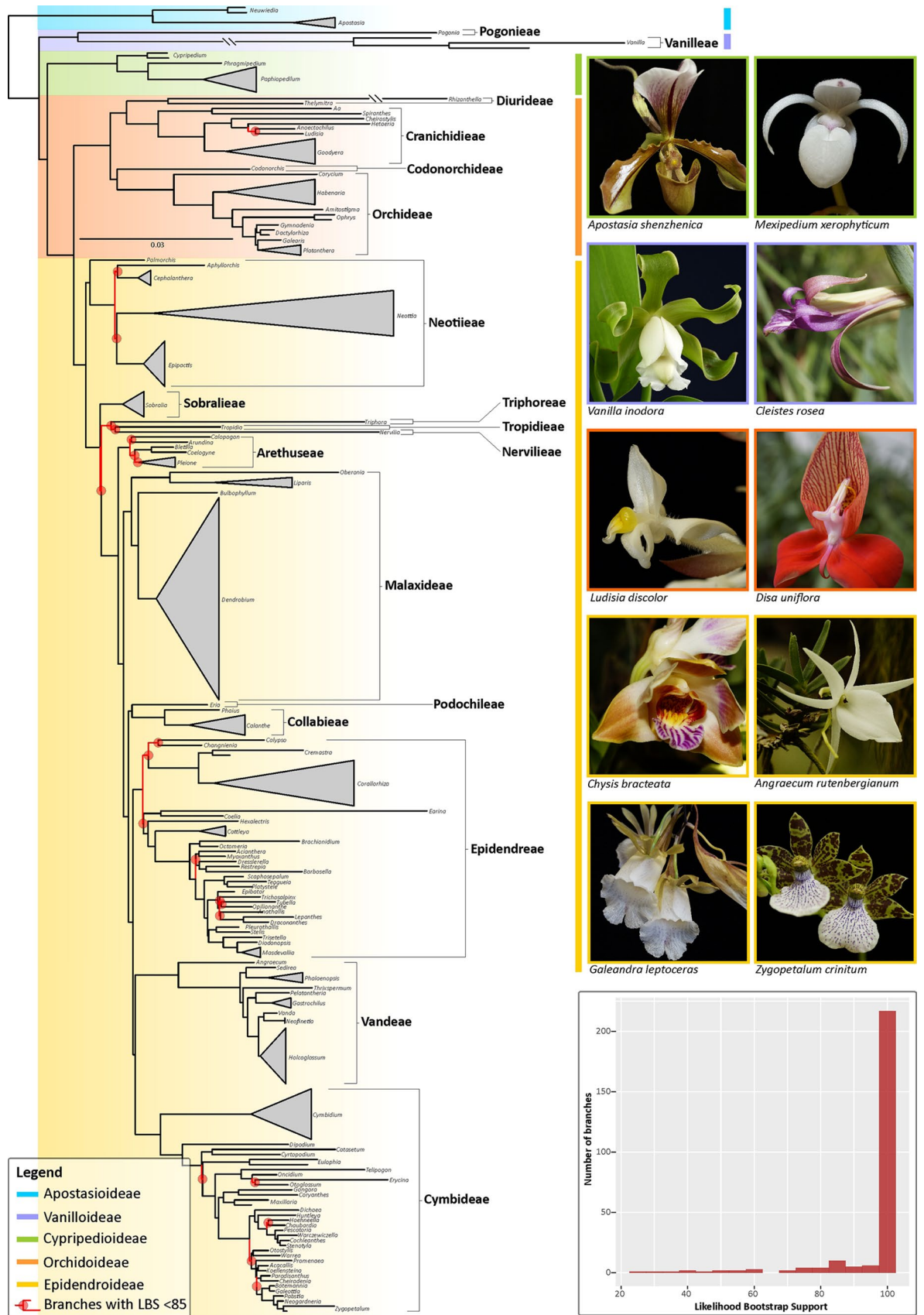


Figure 1. Maximum Likelihood phylogeny of the orchid family inferred from 78 coding plastid genes. Likelihood bootstrap support values (LBS) <85% at nodes are highlighted in red together with their corresponding subtending branches. Orchid genera, tribes and subfamilies are indicated in the phylogeny together with photographs of selected representative species per subfamily. (Inset): Bar plot showing the frequency of LBS values at branches as computed by bin intervals of 5 units. Photos: O. Pérez-Escobar & D. Bogarín.



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