

Erratum to: Isolation and molecular characterization of the fungal endophytic microbiome from conventionally and organically grown avocado trees in South Florida

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The original PDF-version of this article was corrected. During the publication process figures and legends were transposed. Please find below the correct sequence of figures with corresponding legends. The publisher apologizes for any inconveniences caused by this mistake (Figs. 1, 2, 3 and 4).

The online version of the original article can be found at <http://dx.doi.org/10.1007/s11557-016-1219-3>.

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Fig. 1 Cultural characteristics and morphology of asexual structures. **a, d** *Cladosporium tenuissimum*, DMW1045 on Potato Dextrose Agar (PDA) at ambient temperature, 14 days. **b** *Alternaria eichhorniae*, DMW1042 on PDA at ambient temperature, 14 days. **c, e, f** *Strelitziana africana*, DMW1044 on PDA at ambient temperature, 14 days. Scale bars for conidia pictured in **d–f** = 10 μ m

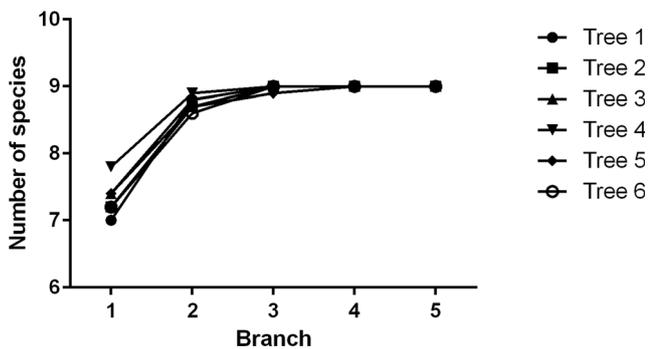
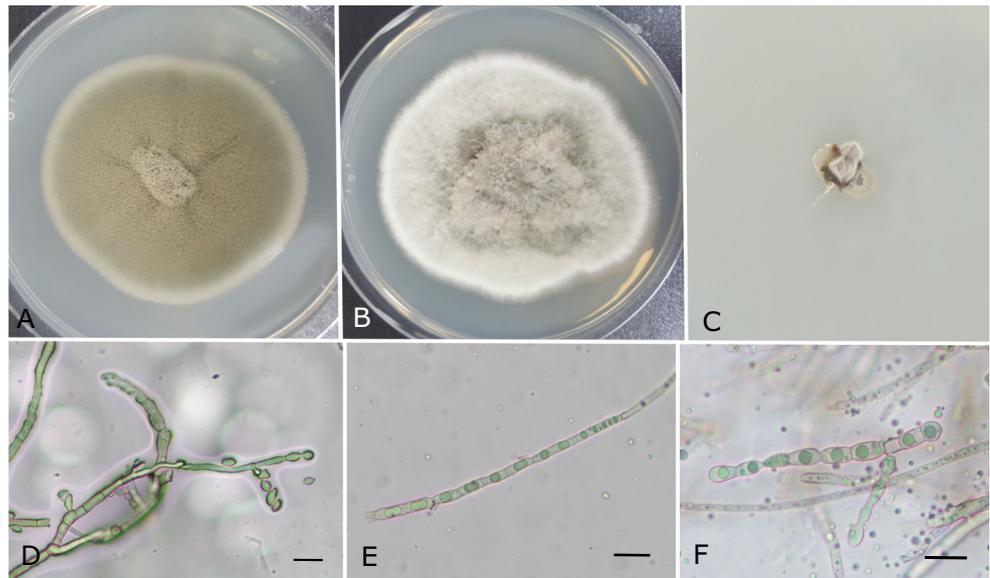


Fig. 2 Organic avocado orchard species accumulation curve of the UGE (Ugland, Gray, and Ellingsen) index of branch sampling effort (x-axis) and number of endophytic species isolated (y-axis)

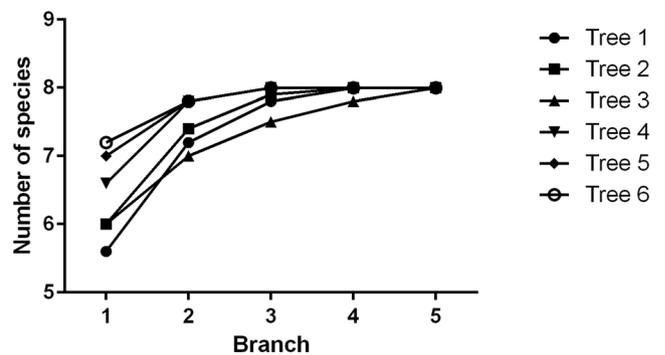


Fig. 3 Conventional avocado orchard species accumulation curve of the UGE (Ugland, Gray, and Ellingsen) index of branch sampling effort (x-axis) and number of endophytic species isolated (y-axis)



Fig. 4 Non-metric multidimensional scaling (nMDS) ordination comparing the β -diversity patterns of endophytic community differences represented as Bray–Curtis distances isolated from five branches from each of six trees ($n=30$ branches) in both organic and conventionally grown orchards. The 2D stress (or measure of

distortion) in the nMDS is low (0.04), indicating that the distance between points is a good representation of the degree of similarity between the fungal communities of each sample. Each point represents a fungal community from a single branch