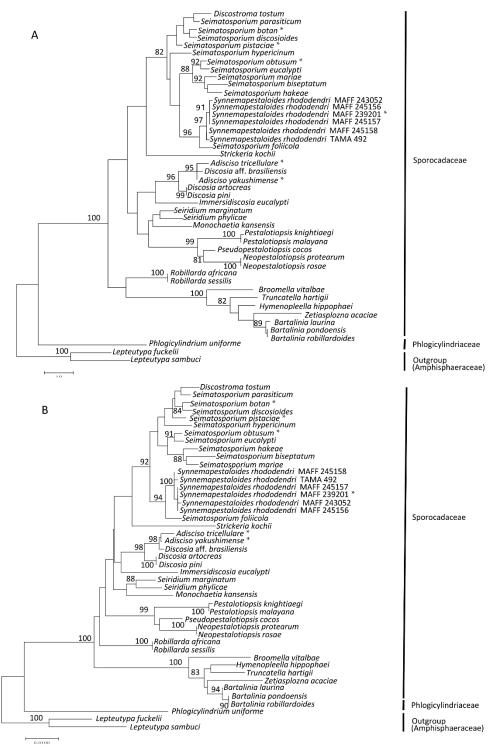
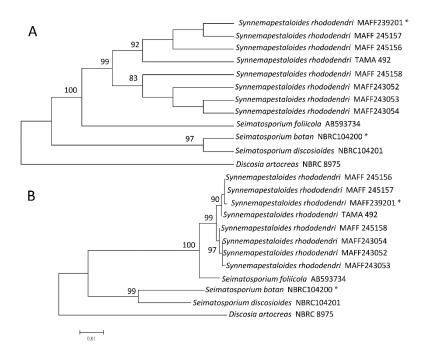
## Supplementary Materials: Phylogenetic Analysis of the Synnema-Producing Genus Synnemapestaloides

Kyoko Watanabe, Mao Sekiguchi, Toyozo Sato, Tom Hsiang, Shigeru Kaneko, Kazuaki Tanaka, Masaru Kanda, Naoko Fujita and Shunsuke Nozawa



**Figure S1.** Phylogenetic trees based on analysis of ITS and LSU (D1–D2) sequences. Numbers on the branches indicate the bootstrap values (%) for each node, calculated from 1,000 replicates. (**A**), MP tree; (**B**), NJ tree. \*: Ex-holotype cultures.



**Figure S2.** Phylogenetic trees based on ITS and  $\beta$ -tubulin sequences. The numbers on the branches indicate bootstrap values (%) for each node, calculated from 1,000 replicates. (**A**), MP tree; (**B**), NJ tree. \*: Ex-holotype cultures.