Supplementary Fig. S1. Phylogenetic tree of Ralfsiales based on the sequences of *rbc*L gene. The value above branches = maximum likelihood bootstrap (BS) values in % ≥50, Bayesian posterior probabilities (BPP) ≥0.75. Values of <BS 50 or BPP 0.75 are indicated by hyphens (-). Values of BS 100 or BPP 1.00 are indicated by asterisks (*).