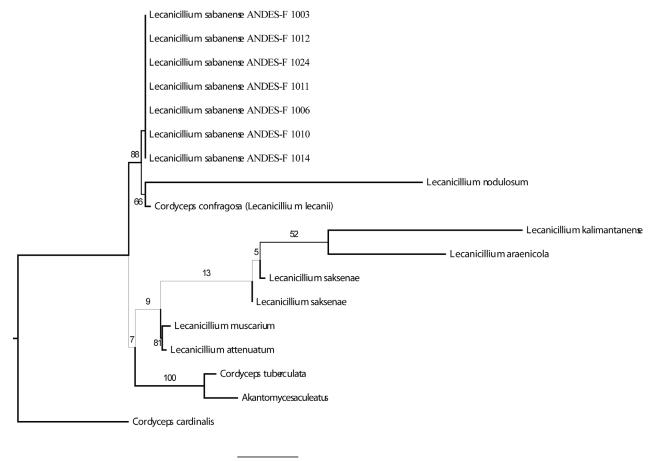
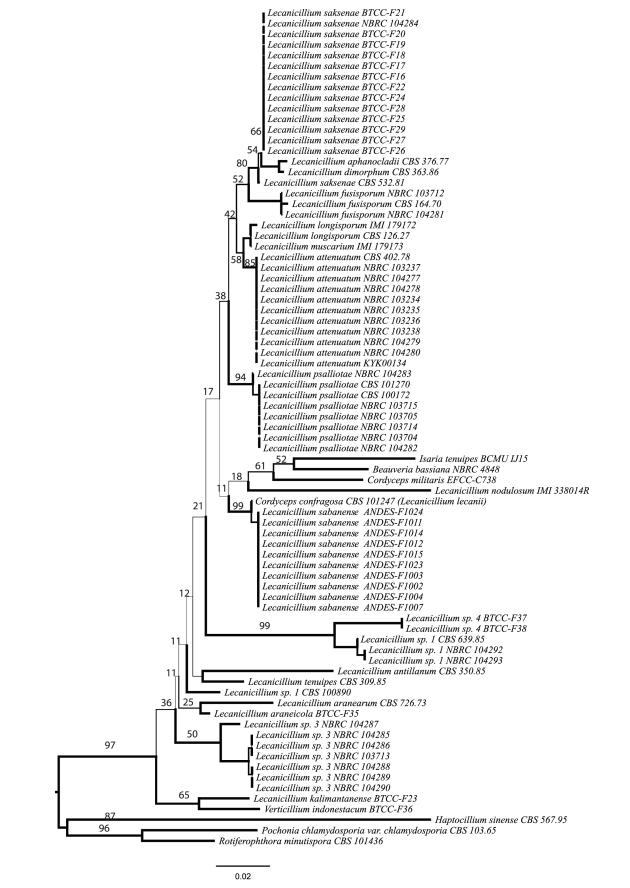


**SUPPLEMENTARY FIGURE 1.** Geolocation, concentration of particulate matter in the air, humidity, and precipitation of each sample site.



0.03

**SUPPLEMENTARY FIGURE 2.** Relationships inferred from Maximum Likelihood analysis of *Cordyceps* species with superficial stromata based on a combined data set of *ITS* and *TEF* gene regions, including *ITS* from *Lecanicillium nodulosum, Lecanicillium araneicola, Lecanicillium kalimantanense*, and *Lecanicillium saksenae*. Numbers above branches indicate Bootstrap support values from 1,000 replications. *Cordyceps cardinalis* was used as outgroup.



**SUPPLEMENTARY FIGURE 3.** Neighbor Joining analysis for the *ITS* region of *Lecanicillium* spp. included in Sukarno *et al.* (2009) as well as *Lecanicillium nodulosum* and *Lecanicillium sabanense*.