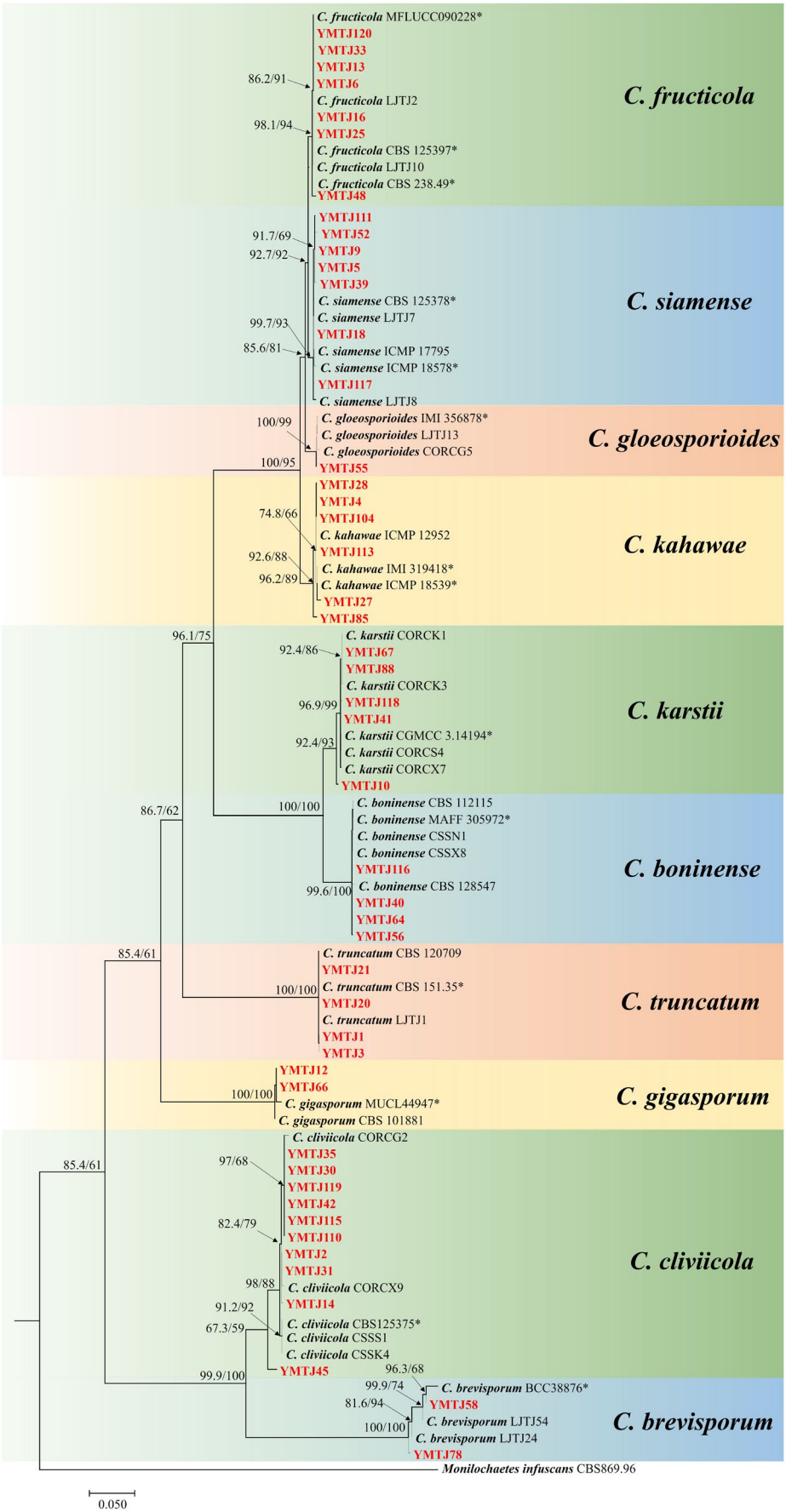

Figure. S2 A Maximum likelihood tree of the *Colletotrichum* species. Phylogenetic relationships of *Colletotrichum* species associated with maize leaves from Sichuan Province, China, based on concatenated sequences of *ACT*, *TUB2*, *CAL*, *GAPDH*, and ITS genes, each analyzed with a separate model of DNA evolution. ML bootstrap and SH-aLRT support values were shown at the nodes (ML ≥ 60%). Isolates from this study are shown in rea and bold. The tree is rooted with *Monilochaetes infuscans*.