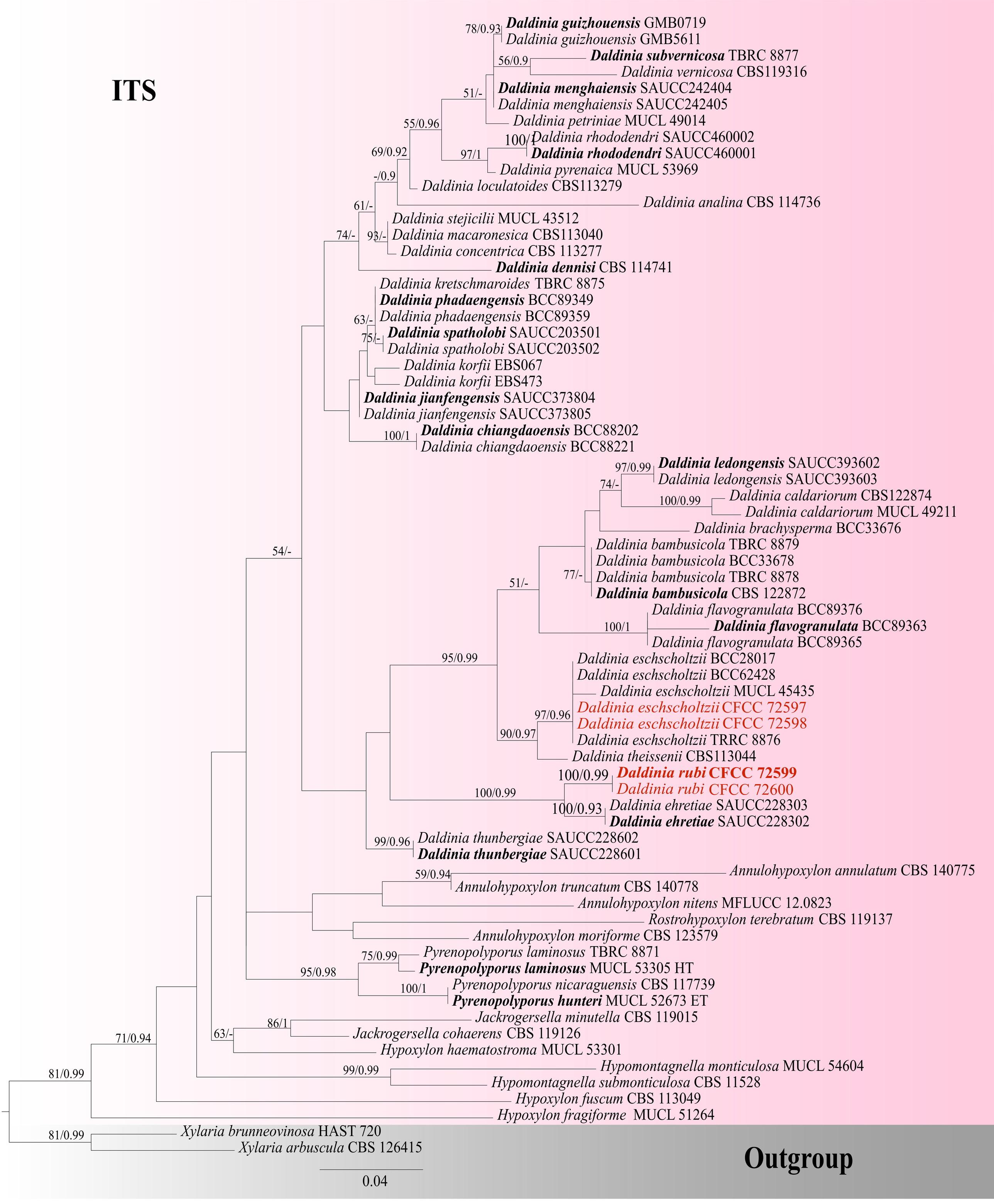

Figure S1-1. Phylogram generated from RAxML analysis based on ITS sequence data of *Daldinia* isolates. The ML (≥ 50%) and BI (≥ 0.9) bootstrap supports are given near the nodes, respectively.

28S

Figure S1-2. Phylogram generated from RAxML analysis based on 28S sequence data of *Daldinia* isolates. The ML (≥ 50%) and BI (≥ 0.9) bootstrap supports are given near the nodes, respectively.

Figure S1-3. Phylogram generated from RAxML analysis based on tub2 sequence data of *Daldinia* isolates. The ML (≥ 50%) and BI (≥ 0.9) bootstrap supports are given near the nodes, respectively.