

5 changes

Figure S5 Gene genealogies of the nucleotide sequences of (A) SLA (898 characters, a maximum parsimony tree), (B) COX13 (788 characters, a neighbor-joining tree), and (C) APN (1444 characters, neighbor-joining tree) genes demonstrating the phylogenetic relationships among G. clavigera and related species. MAT1-2 isolates are indicated in green while MAT1-1 isolates are highlighted in grey.