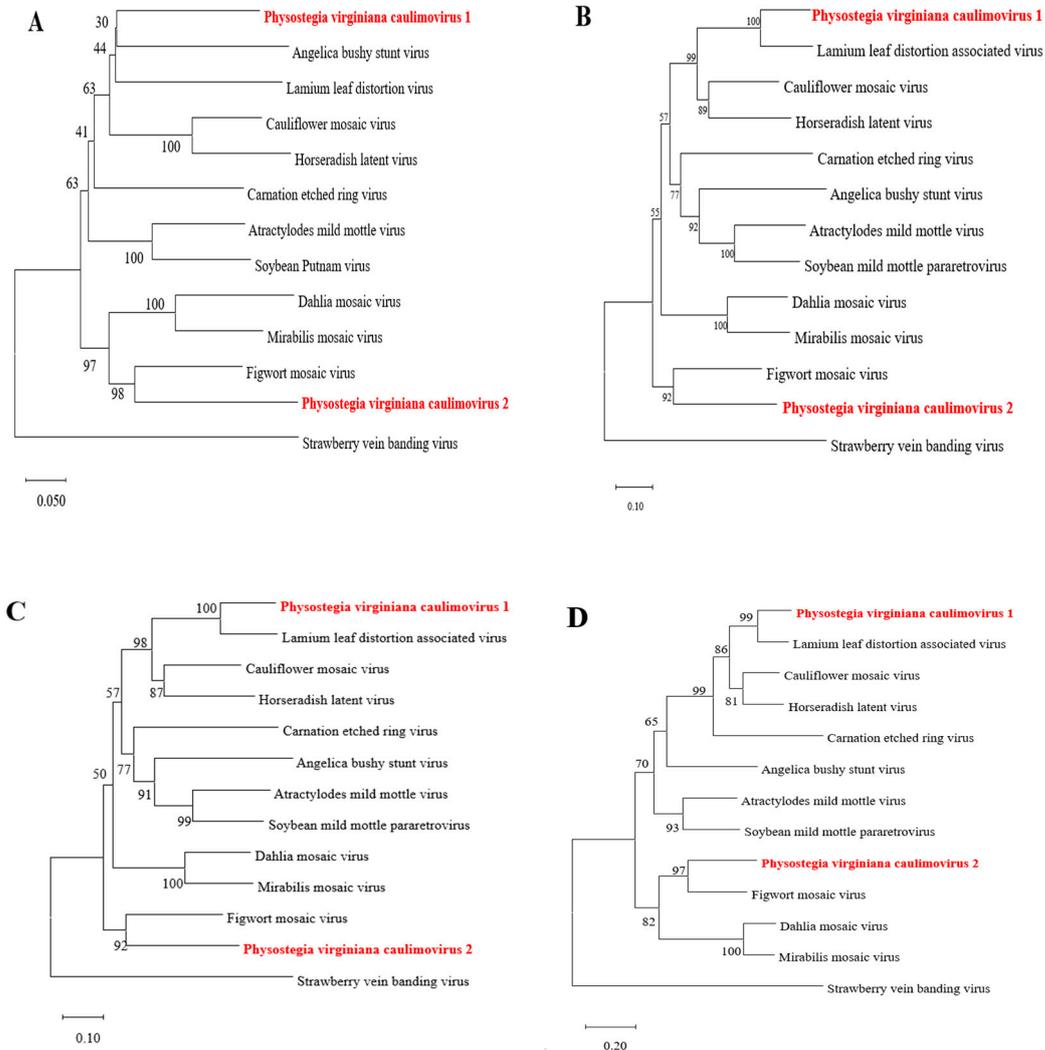
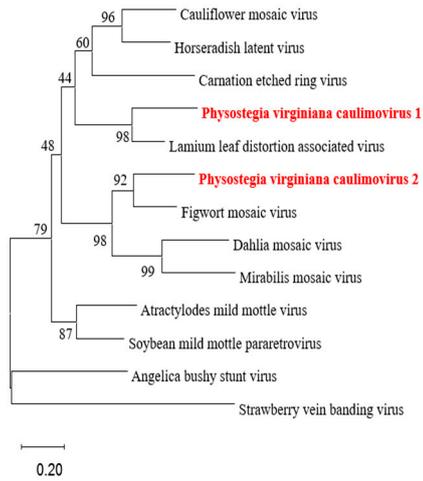


Figure S1. Phylogenetic trees of PVCaV1 and 8 representative viruses from the genus *Fabavirus* based on RdRp (A), LCP (B) and SCP (C). The phylogenetic tree was constructed using the Neighbor-Joining method with 1000 bootstraps. PVCaV1 is shown in red.



E**F**