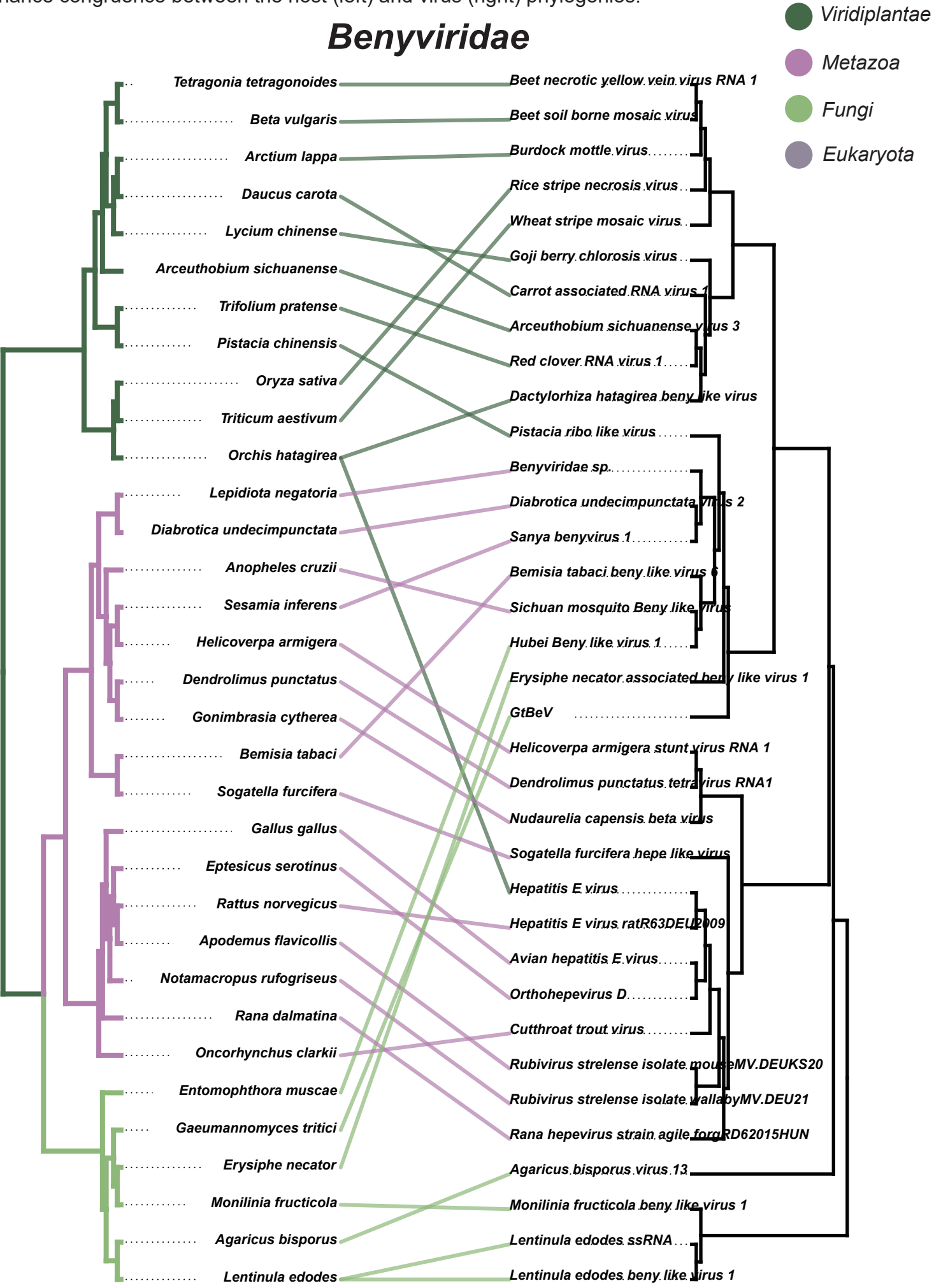
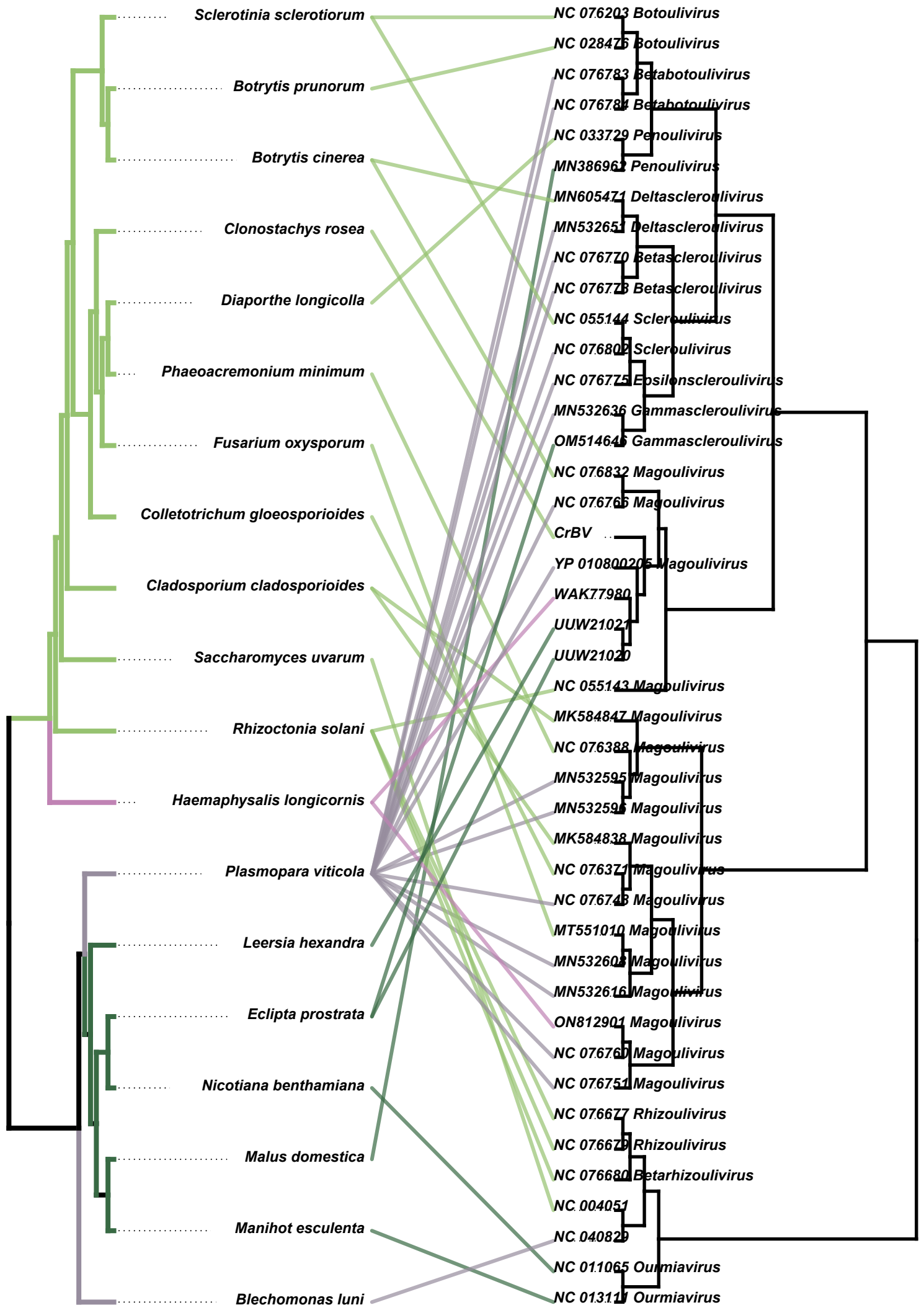


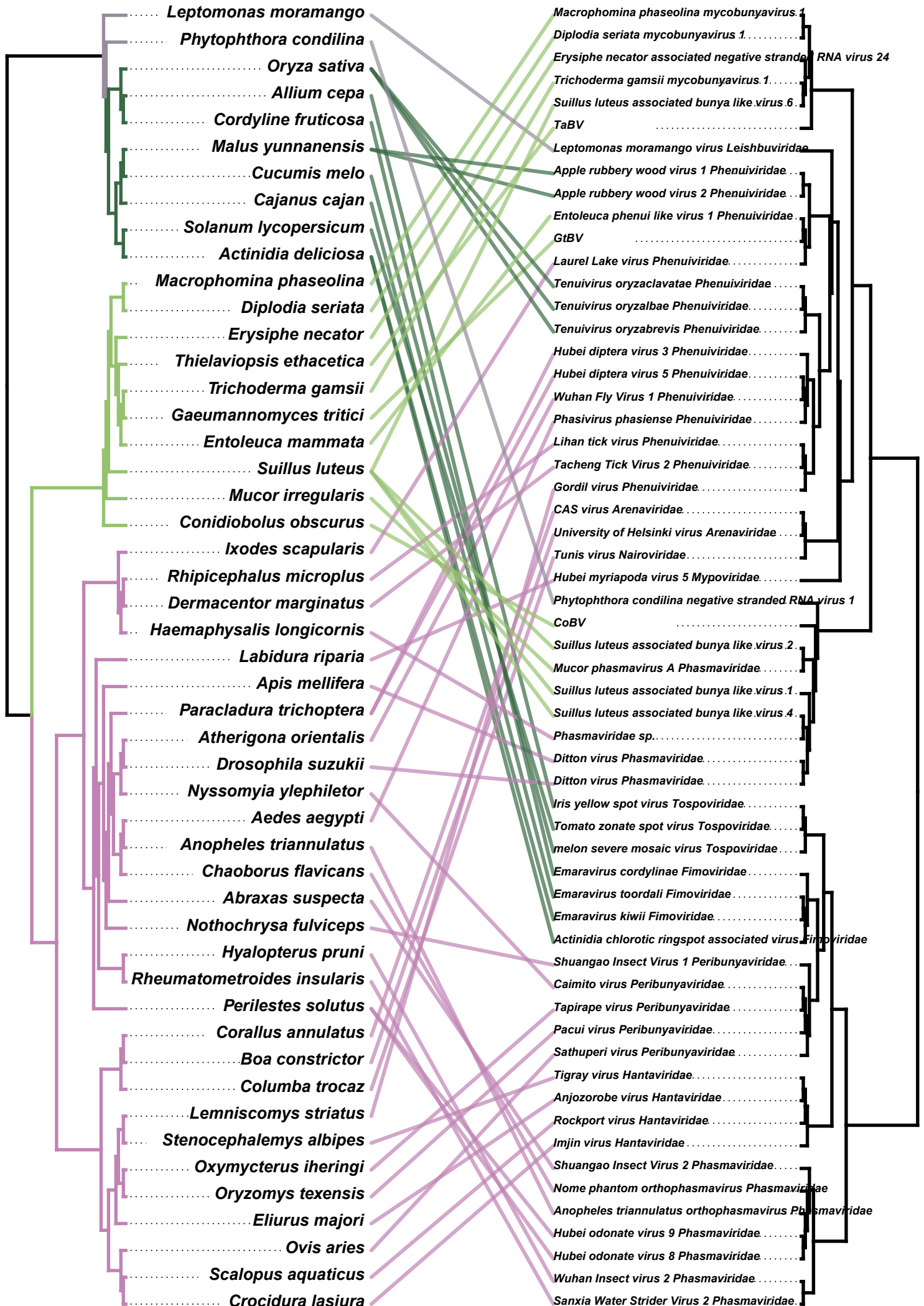
Supplementary Figure 1. Tanglegram of phylogenetic trees for virus orders/families and their hosts. Lines and branches are color-coded to indicate host clades. The cophylo function in phytools was employed to enhance congruence between the host (left) and virus (right) phylogenies.



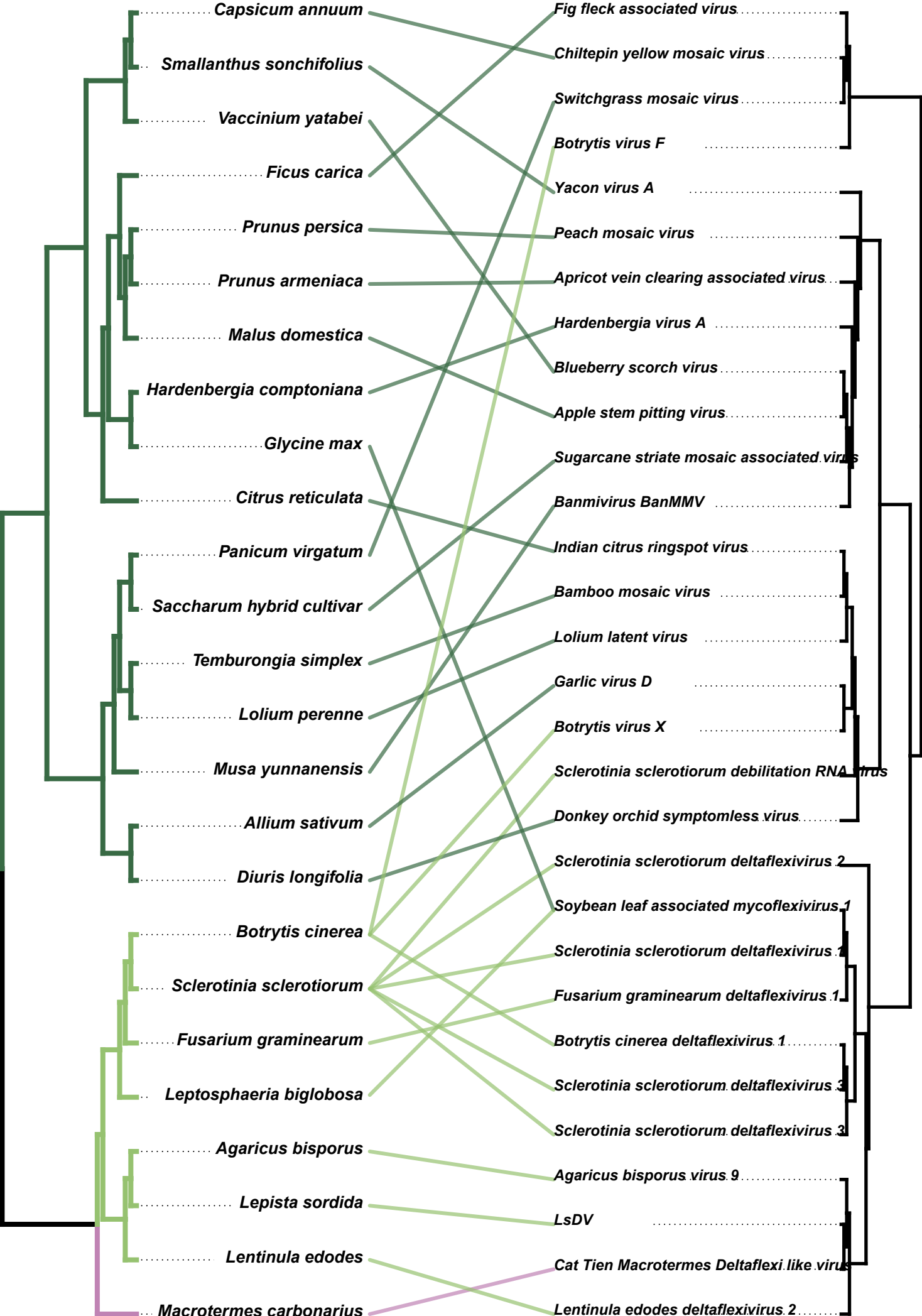
Botourmiaviridae



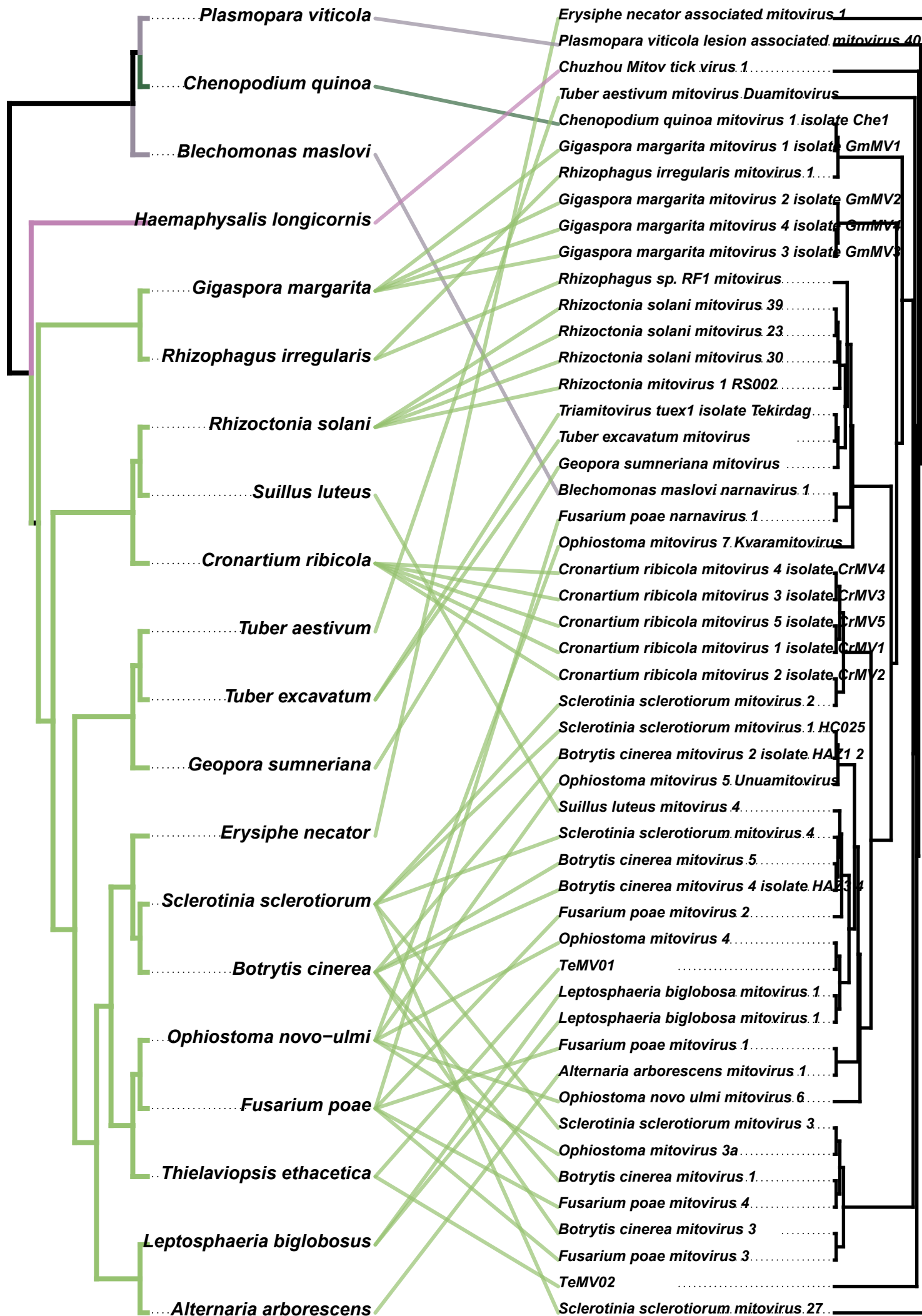
Bunyavirales



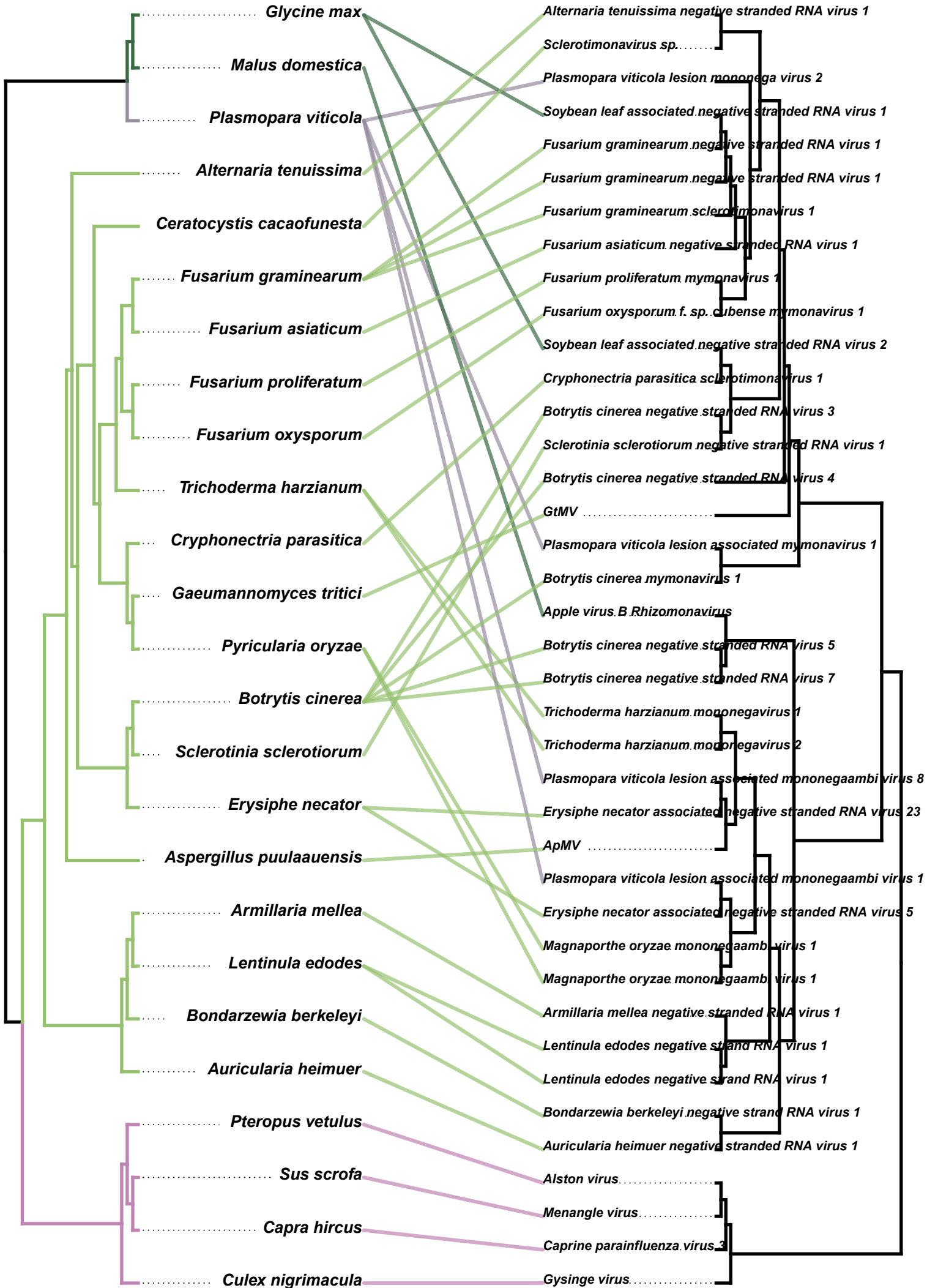
Deltaflexiviridae



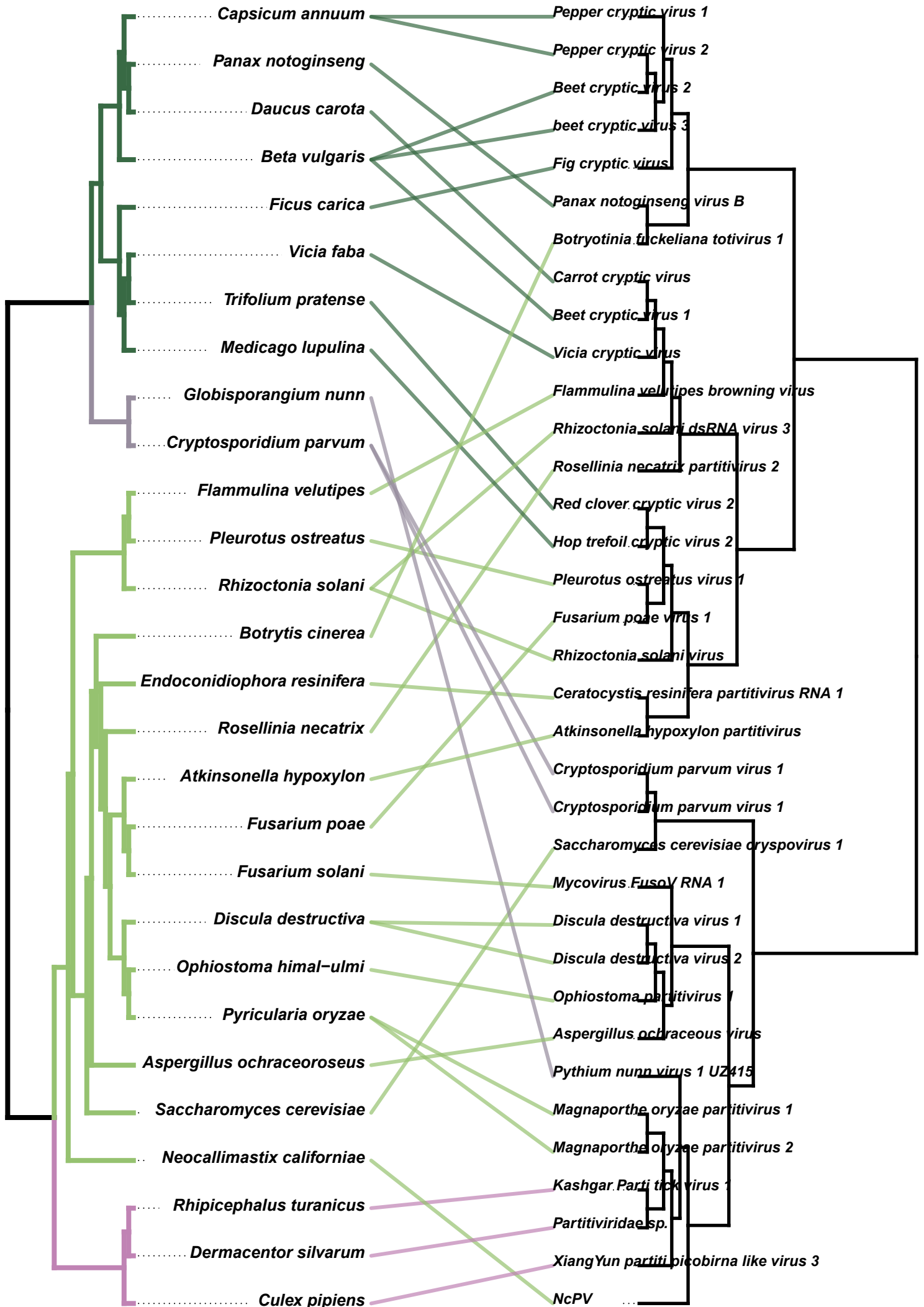
Mitoviridae



Mymonaviridae



Partitiviridae



Togaviridae

