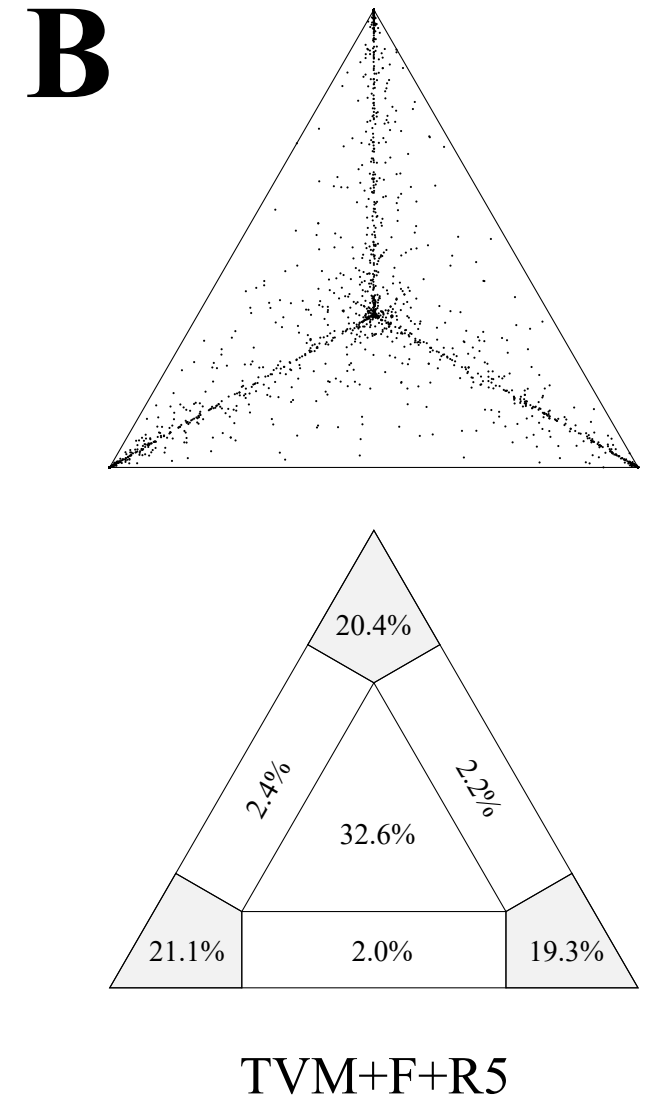
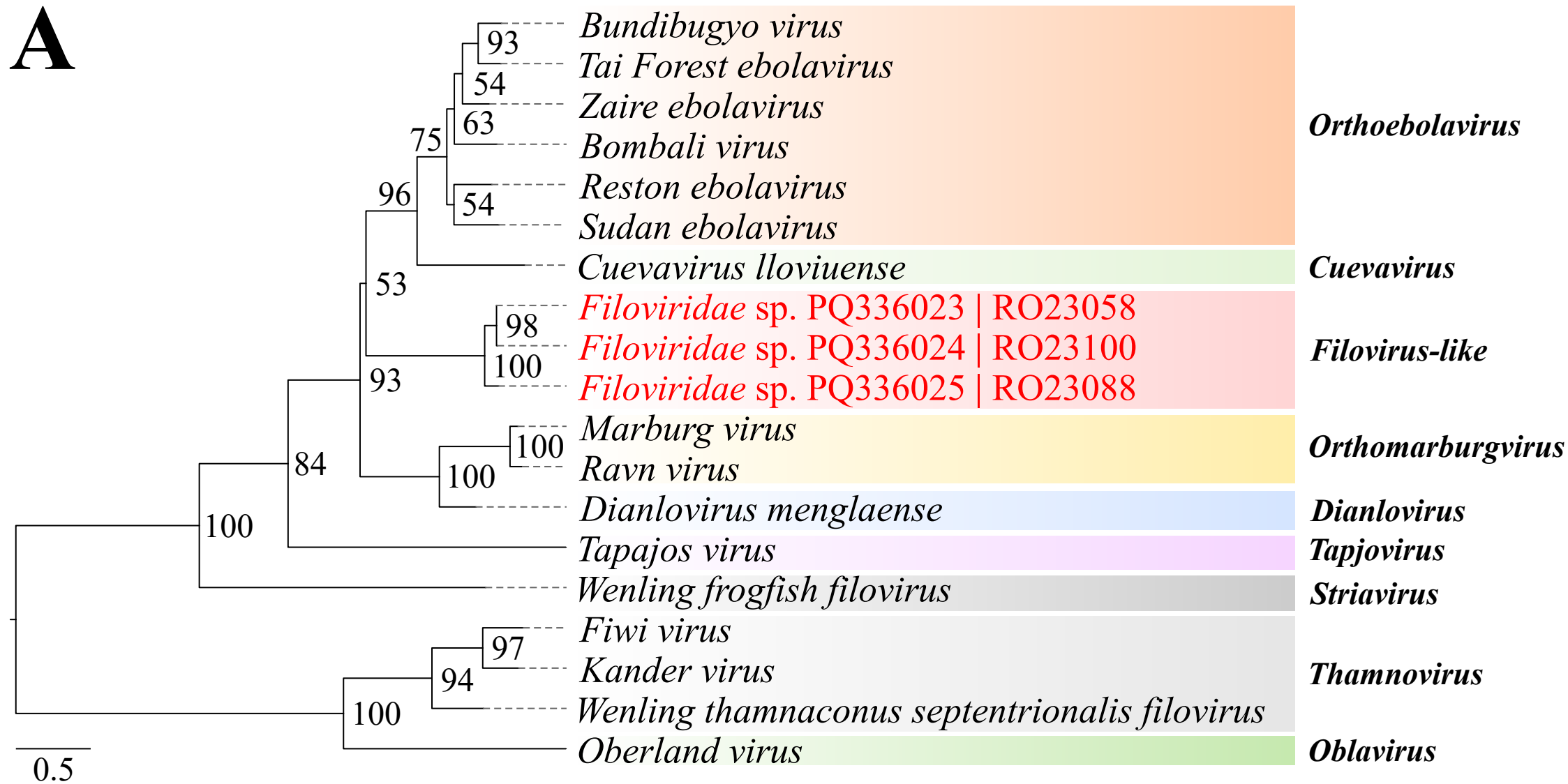




Supplementary Figure 4. (A) Phylogenetic reconstruction using the Maximum Likelihood method based on nucleotide sequences of *Filoviridae*. (B) Maximum Likelihood mapping diagram showing the quality of the phylogenetic signal from the quartet analysis, with 60.8% of quartets resolved