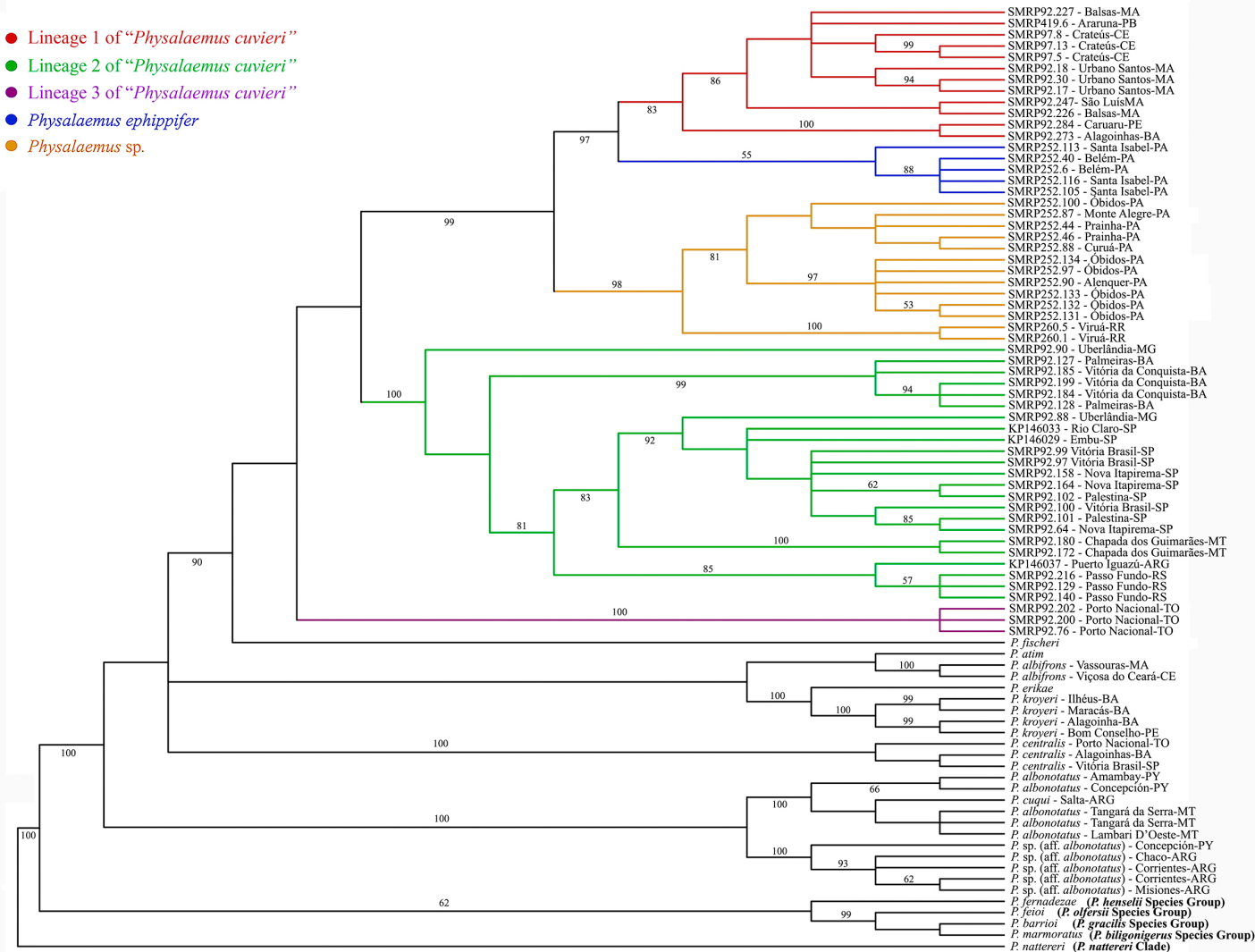




Supplementary Figure S2. Phylogenetic relationships inferred by TNT. Strict consensus cladogram of four most parsimonious trees (2360 steps). Bootstrap values ($\geq 50\%$) are indicated on the branches.