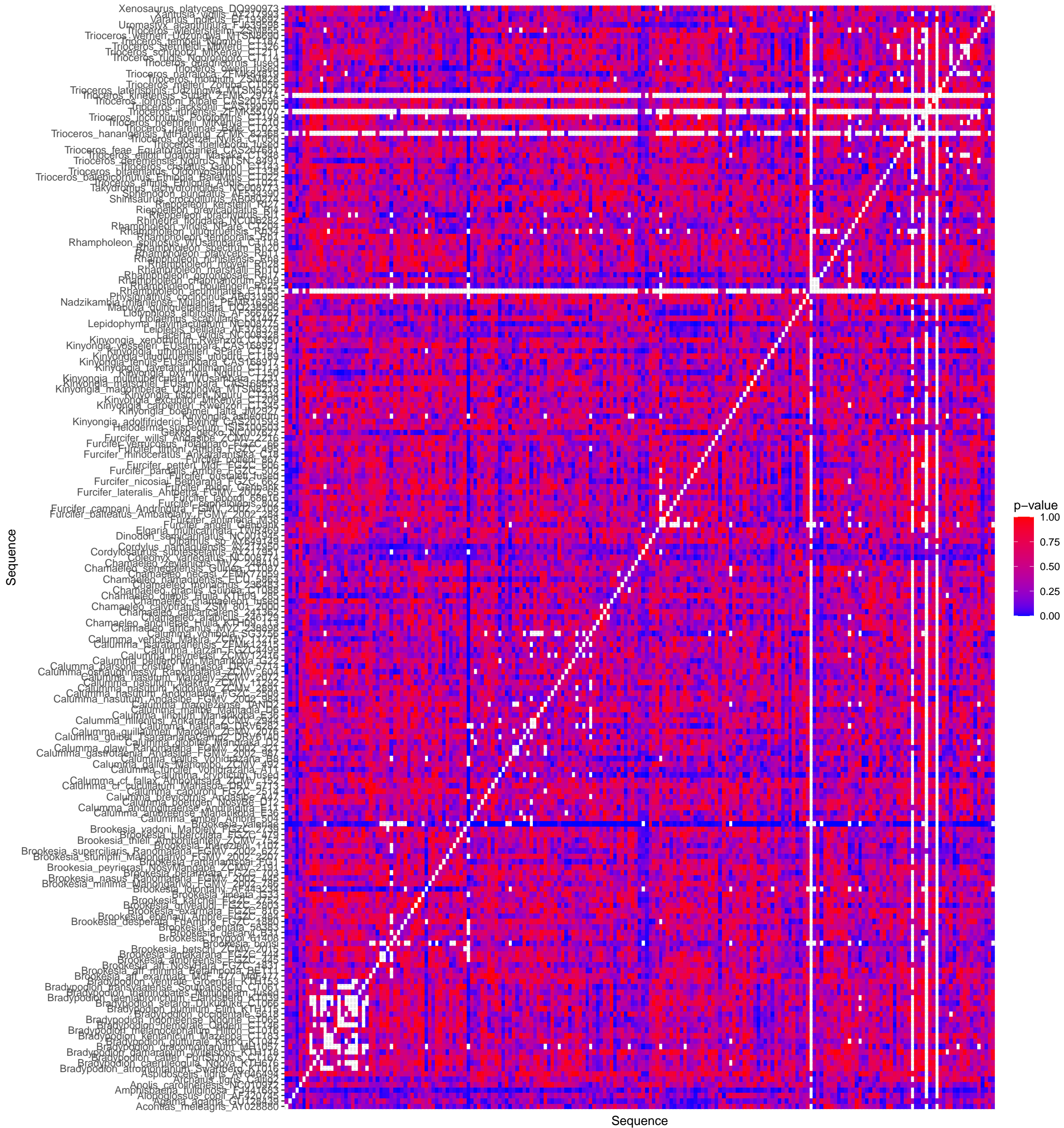
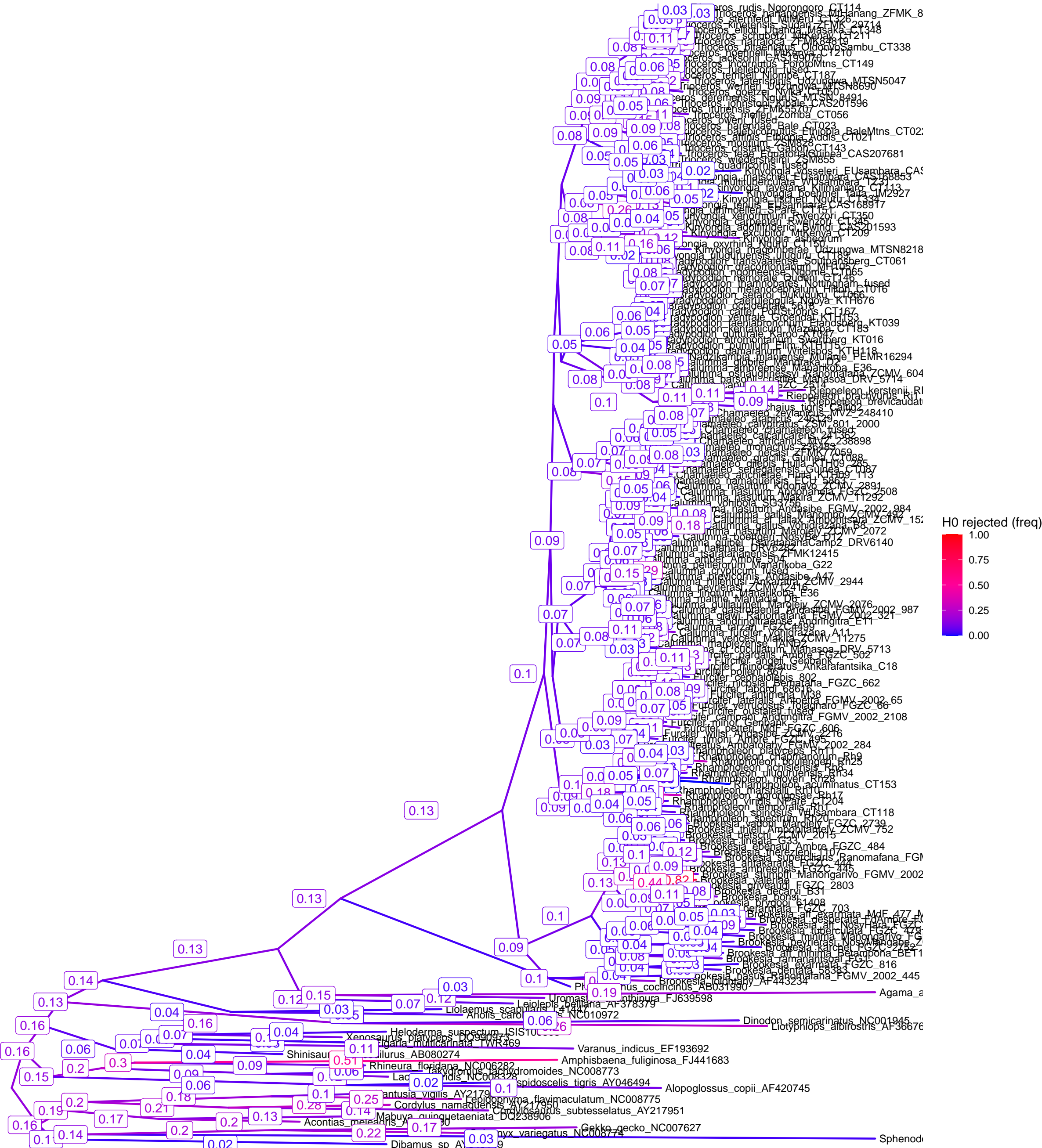


IS Test, omitted pairs: 4.7 %



H0 rejected (freq) on each edge, IS Test, Midpoint root





IS Test, Compressed Tree, Midpoint root



[illegible]

A phylogenetic tree with two main branches. The upper branch leads to the node 'Heloderma\_suspectum\_ISIS100503'. The lower branch leads to the node 'Xenosaurus\_platycephalus\_DQ990973'. The tree is rooted on the left and branches out to the right.

Phylogenetic tree showing the relationships between *Agama agama*, *Uromastix acanthinura*, and *Leiolepis belliana*. The tree is rooted at the bottom left, with *Agama agama* at the top right, *Uromastix acanthinura* in the middle, and *Leiolepis belliana* at the bottom right.