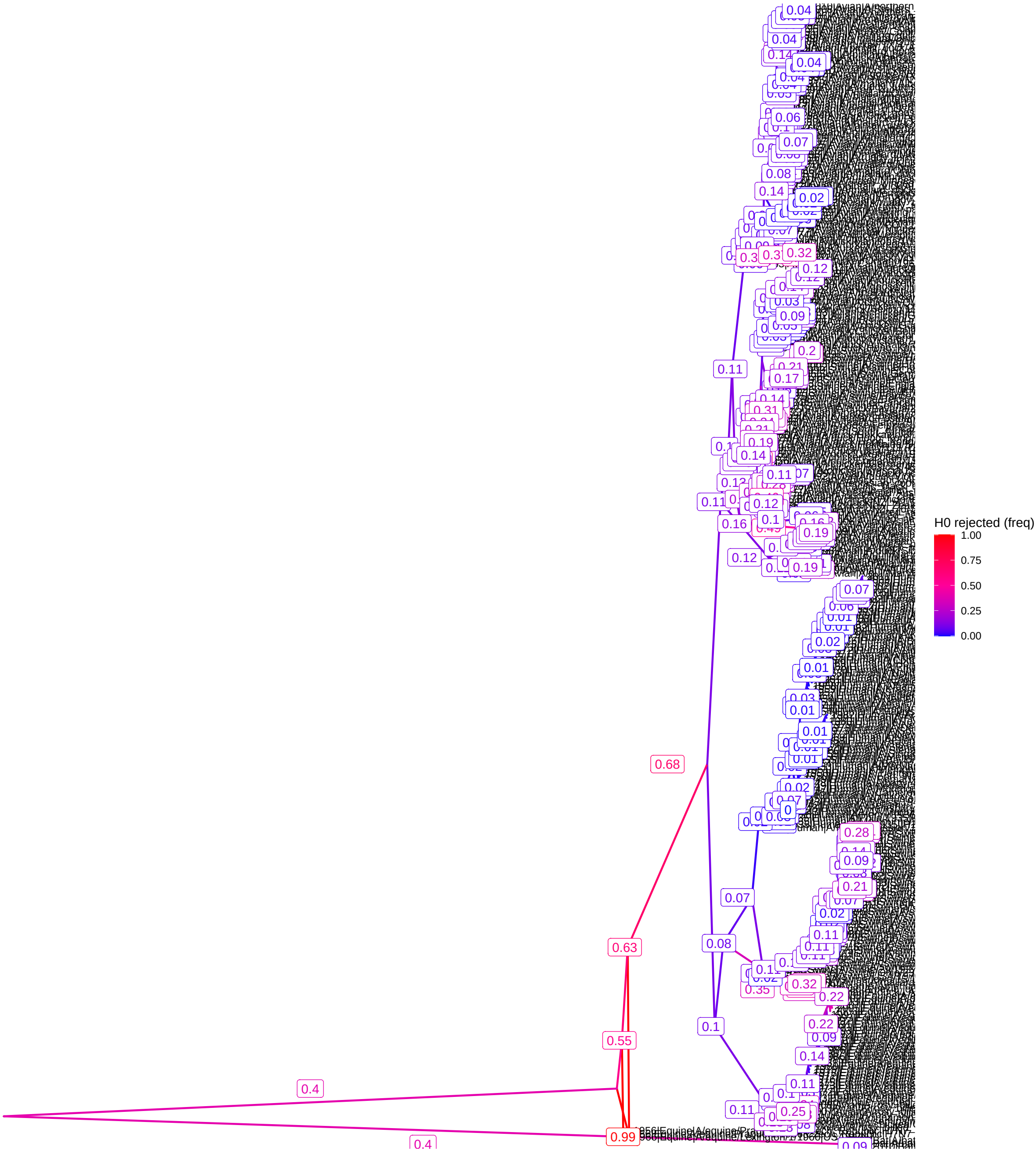
Stuart Test, omitted pairs: 11.53 %

Sequence

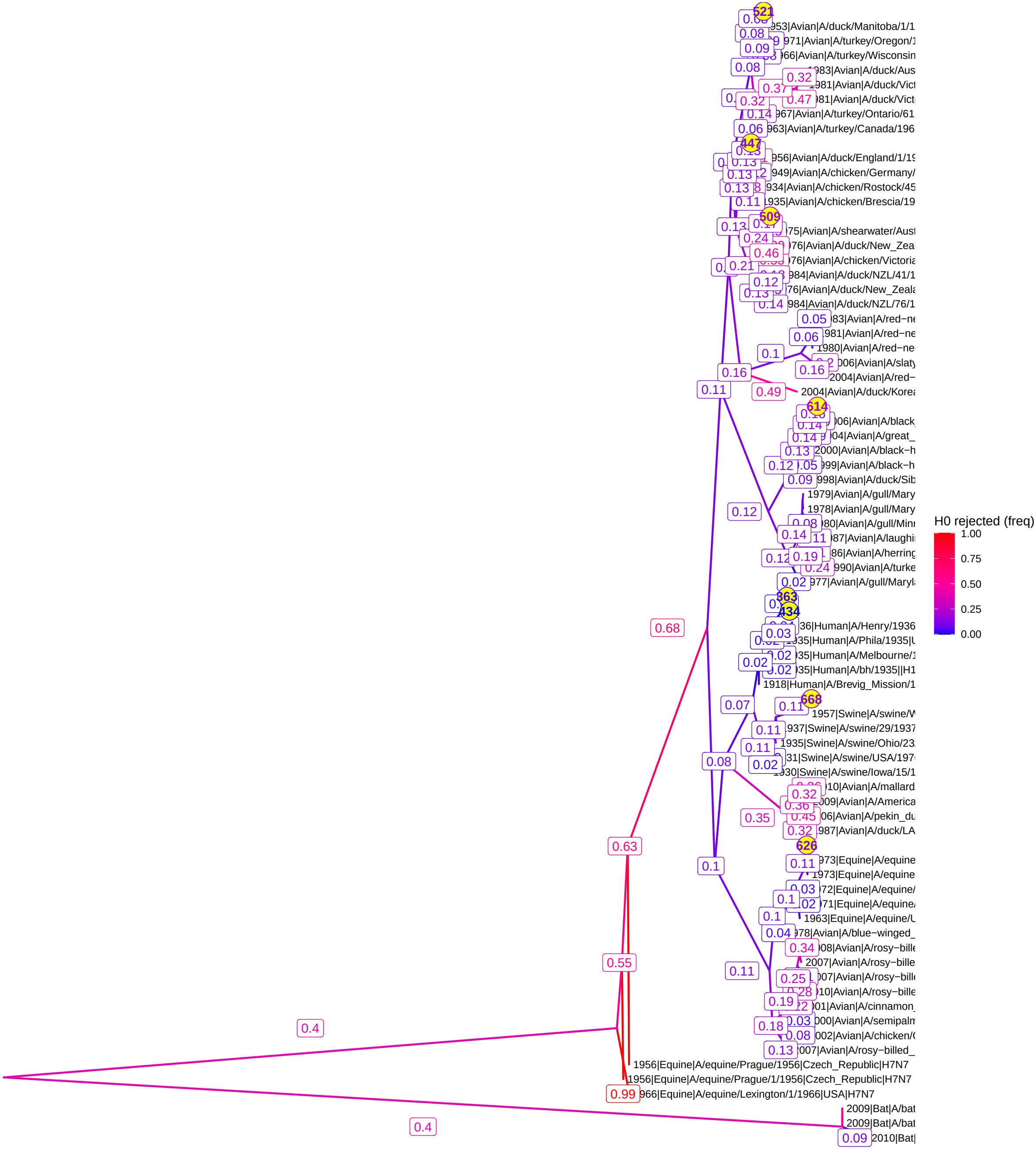
Sequence

p−value
1.00
0.75
0.50
0.25
0.00

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

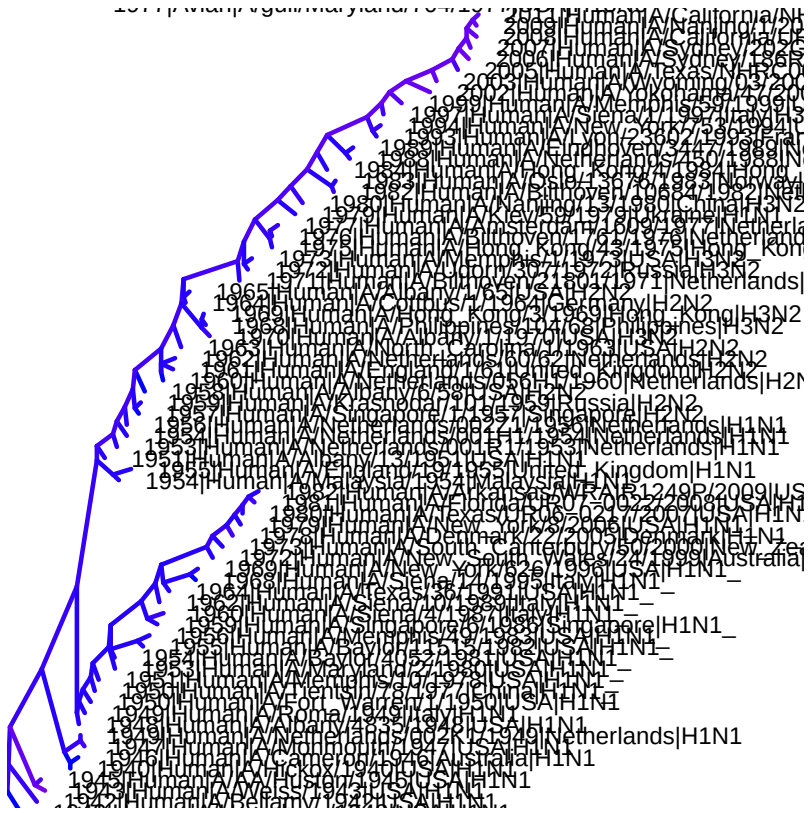


Stuart Test, Compressed Tree, Midpoint root

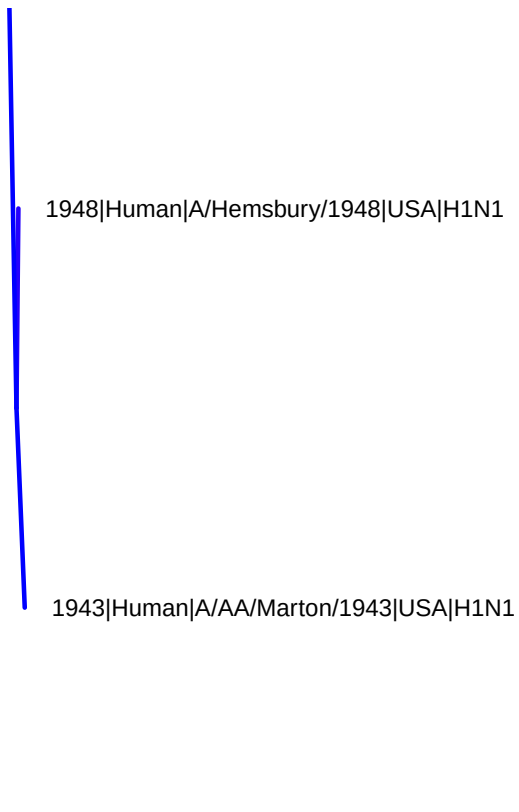


Collapsed Nodes

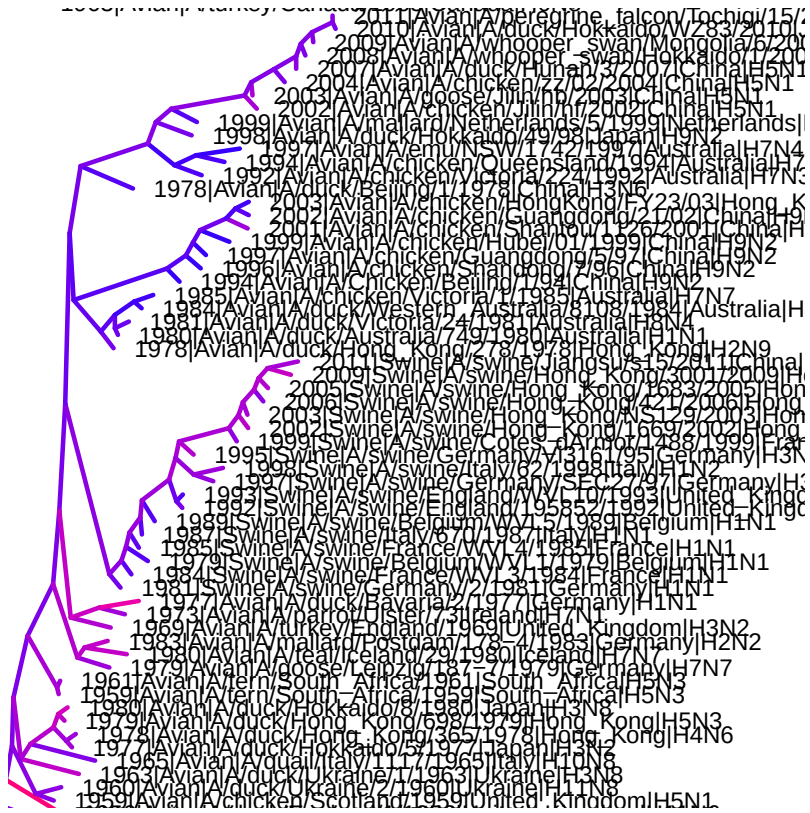
Node 363



Node 434

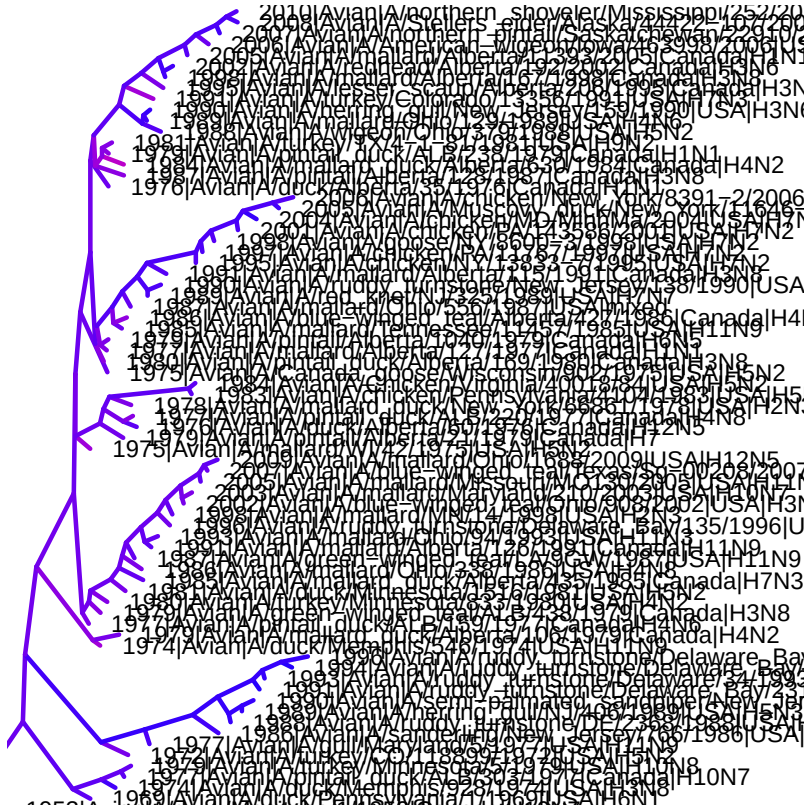


Node 447

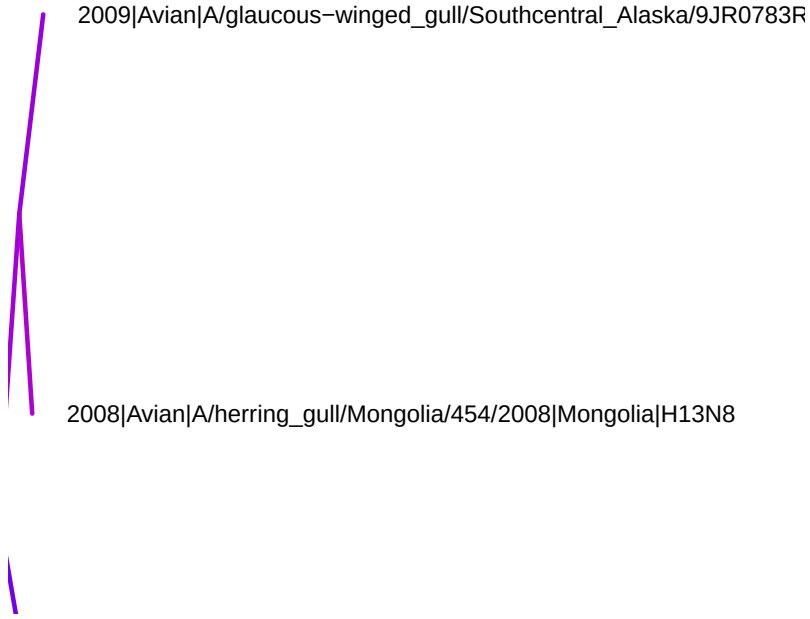


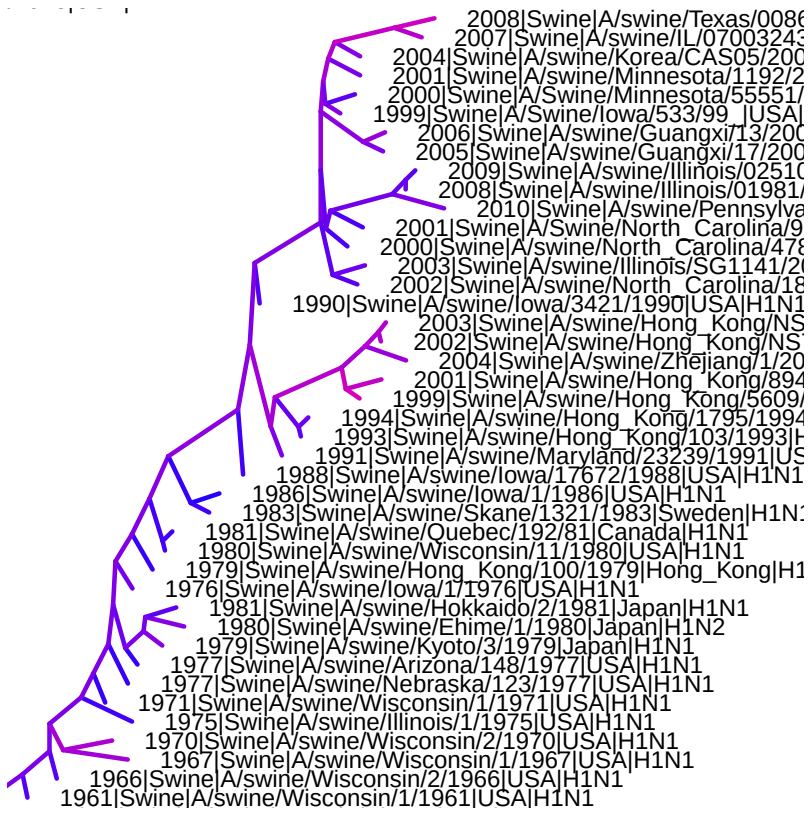
Node 509

Node 521



Node 614



Node 626

Node 668