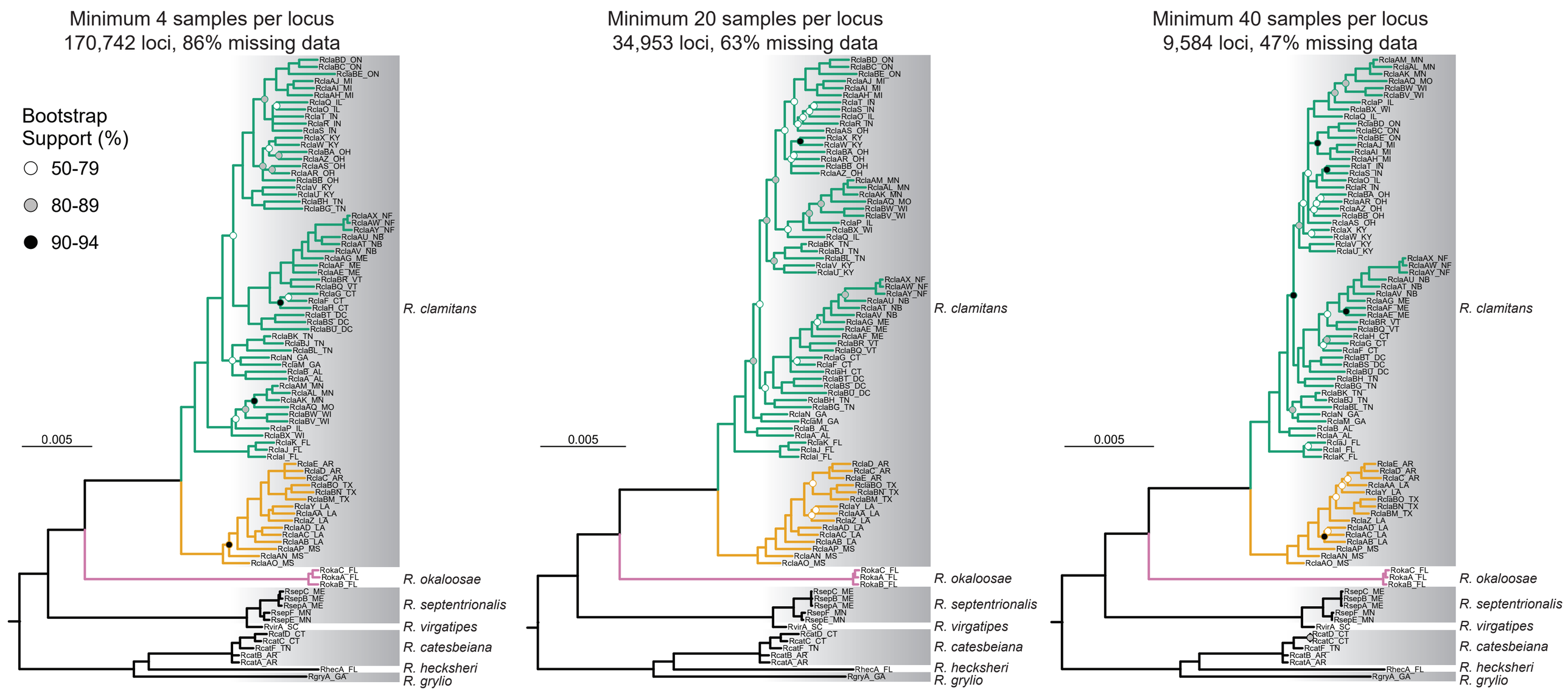
Minimum 4 samples per locus
170,742 loci, 86% missing data
Minimum 20 samples per locus
34,953 loci, 63% missing data
Minimum 40 samples per locus
9,584 loci, 47% missing data
Bootstrap Support (%)
50-79
80-89
90-94
0.005
R. clamitans
R. okaloosae
R. septentrionalis
R. virgatipes
R. catesbeiana
R. hecksheri
R. grylio