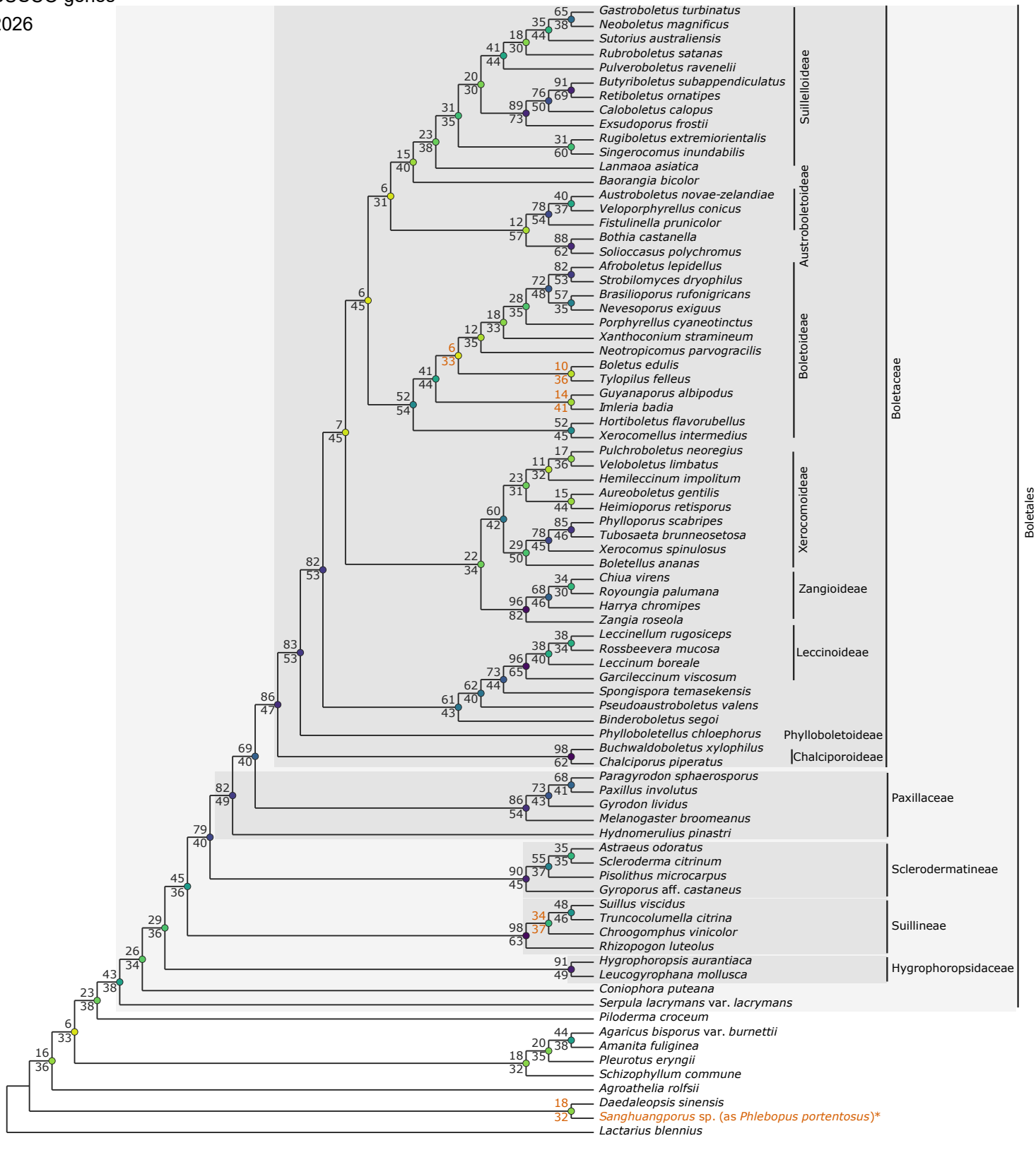
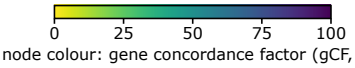
Boletales

71 genomes. 1440 BUSCO genes

Jerry Cooper, Nov. 2026

ML tree: suborders and Boletaceae concordant, backbone and Boletoideae not





node colour: gene concordance factor (gCF, %)

numbers: gCF above, sCF below the branch (%); cladogram, ML branch lengths not to scale
orange = UFBoot < 95 (52-87); all other nodes SH-aLRT / UFBoot >= 99 / 97
* genome deposited under a misidentified name (genomic ITS = Hymenochaetales)