

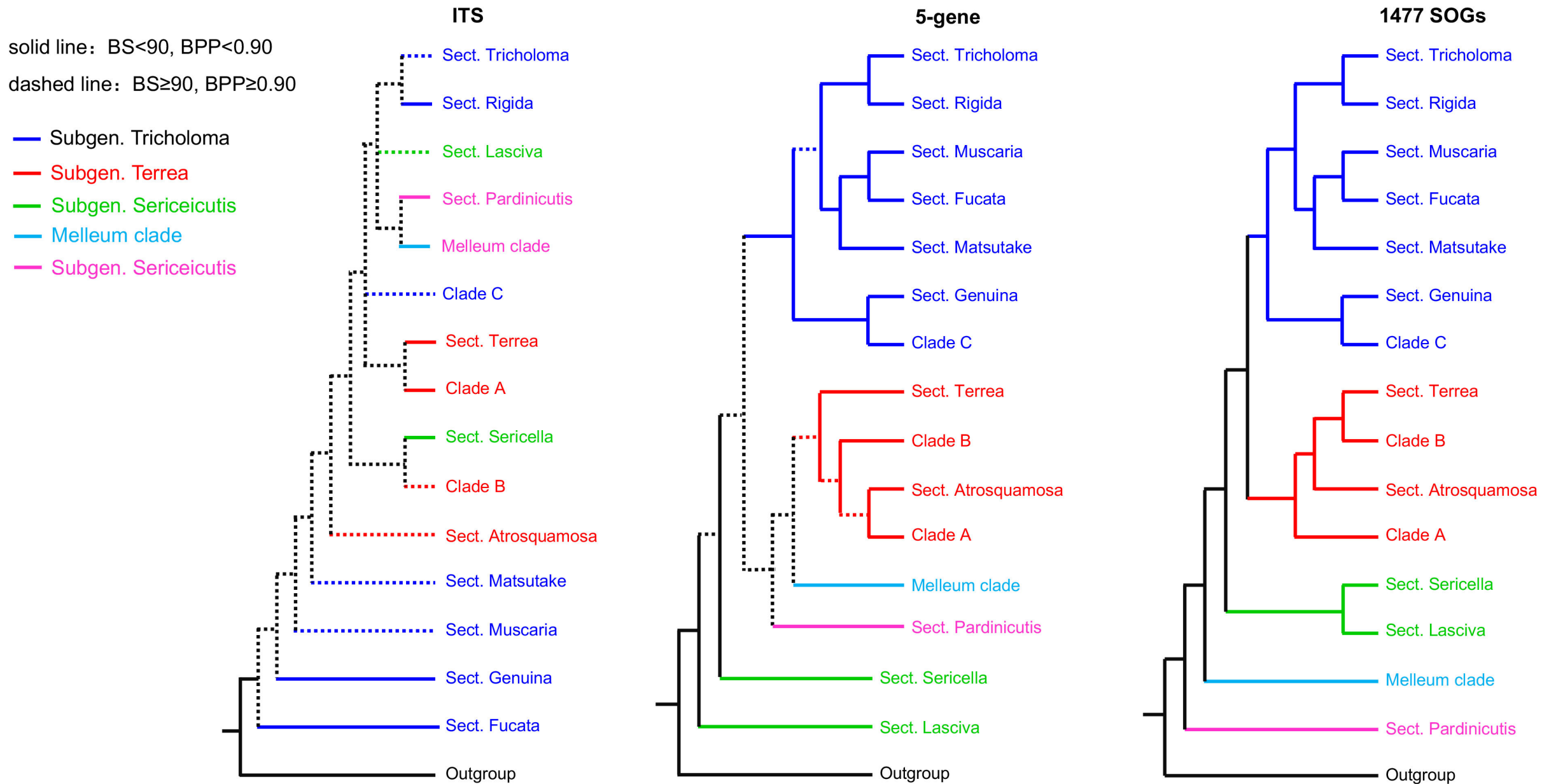


Fig. S1 A comparison of the three *Tricholoma* phylogenetic trees inferred from Dataset II (1 477 SOGs), Dataset III (ITS/RPB2/MCM7/EF1- α /mtSSU) and Dataset IV (ITS)