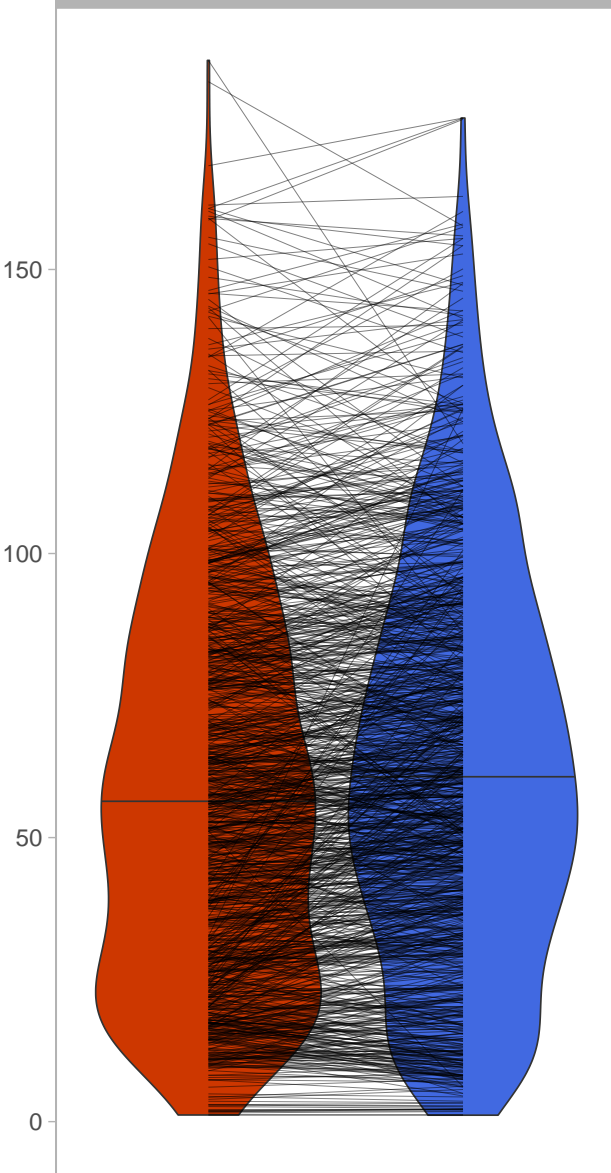
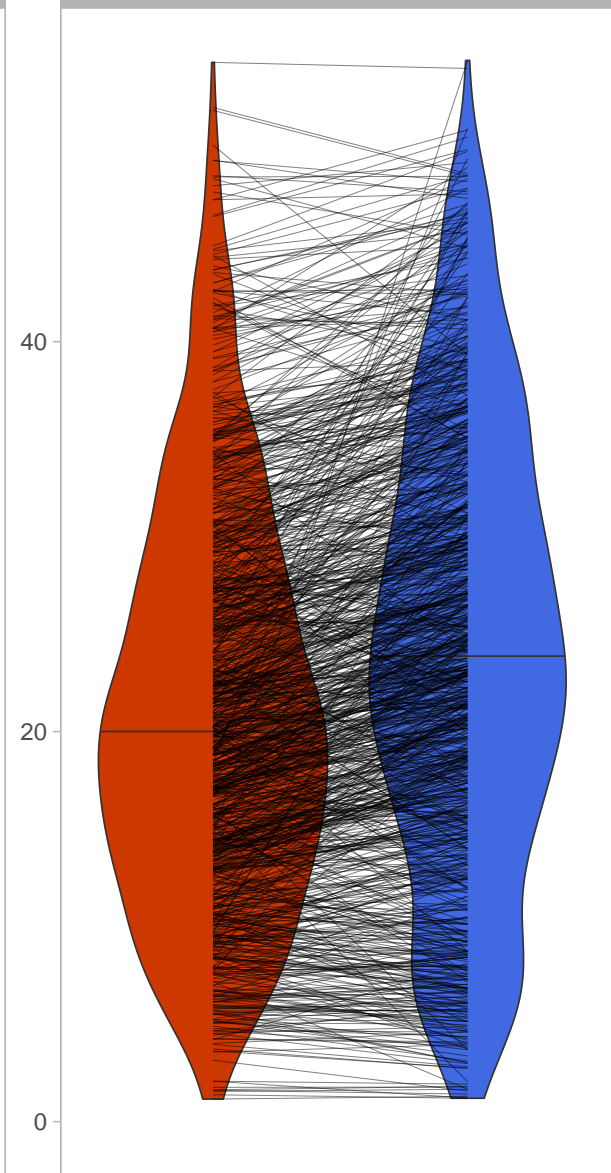


A. GTDB 220 (n = 1039)

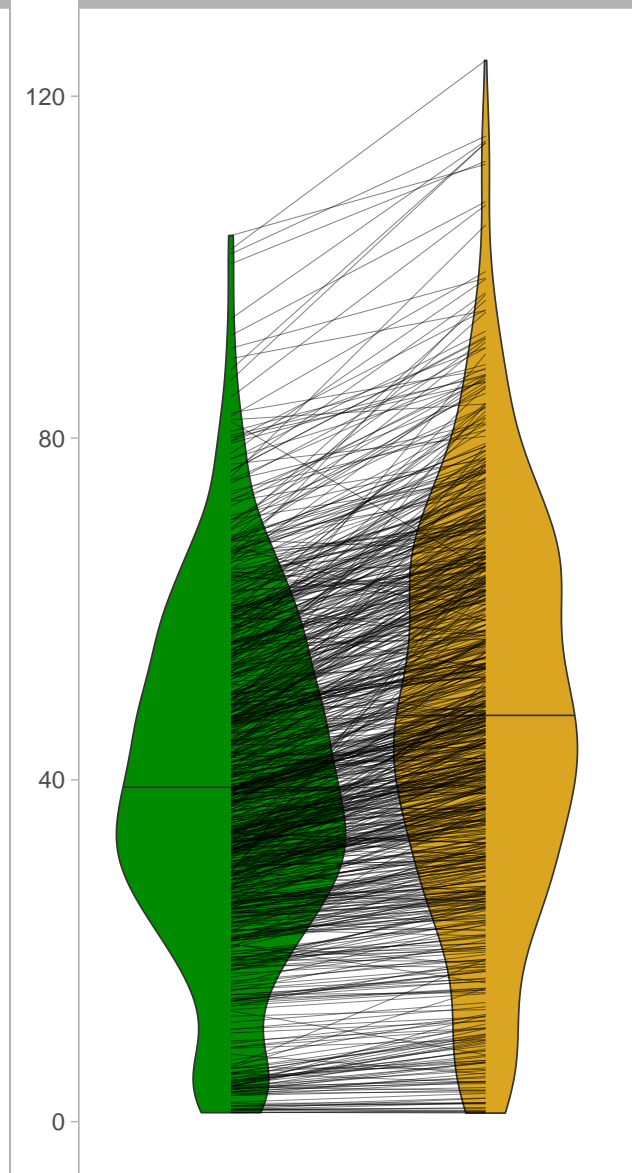
Hill number of order 1 (effective number of equally abundant species)



B. RefSeq 2024-12-28 (n = 1040)



C. DNA-to-marker methods (n = 792)



Methodology ■ Kraken2+Bracken ■ MetaPhlAn4 ■ mOTUs3 ■ Sourmash gather