Supplemental Figure 6: Unweighted UniFrac intra-sample distances between sequence processing methods based on three real datasets reveal highly different biological profiles.

A-C) The unweighted UniFrac distances between the same biological samples based on ASVs/OTUs outputted by each of the different sequence processing methods on the soil, human associated, and Exercise datasets respectively. D-F) Principal coordinates analysis of the unweighted UniFrac distances of all the samples in the soil, human associated, and Exercise datasets respectively. The four different sample profiles generated for each biological sample are colour-coded and are joined by an interconnecting line.