Supplemental Figure 5: Bray-Curtis dissimilarity intra-sample distances between sequencing processing methods based on three real datasets reveals relatively similar biological profiles.

A-C) The Bray-Curtis dissimilarity 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) Non-metric multidimensional scaling plots 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.