



Figure 3. PCoA based on Bray-Curtis dissimilarity matrix computed on Hellinger-transformed ASVs relative abundances per sampling (PCoA1/PCoA2 panel on the left) and Ward.D2 clustering on Euclidean distances between samplings' coordinates along the first seven axes (on the right). Colours refer to clusters in both panels. The vertical dashed line represents the threshold used to define the three clusters.