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Supplemental Figure 7: Comparing Beta Diversity metrics between Deblur with and without positive filtering reveals little difference between the two based on beta diversity metrics

A) The weighted UniFrac intra-sample distances between the same biological samples based on ASVs/OTUs outputted by each sequence processing method and either Deblur or Deblur without positive filtering. B) The unweighted UniFrac intra-sample distances between the same biological samples based on ASVs/OTUs outputted by each sequence processing method and either Deblur or Deblur without positive filtering. C) The Bray-Curtis dissimilarity intra-sample distances between the same biological samples based on ASVs/OTUs outputted by each sequence processing method and either Deblur or Deblur without positive filtering. C) The Bray-Curtis dissimilarity intra-sample distances between the same biological samples based on ASVs/OTUs outputted by each sequence processing method and either Deblur or Deblur without positive filtering.