



S9 Fig. Stacked barchat (a) showing the seven α -Proteobacteria orders, relative abundances and their PCoA plots (b), revealing the nucleotide composition variation among the reactors based on the Euclidean model. The plots revealed close proximity of reactor 1 and 6 nucleotide composition and were positioned on the y-axis. However, the composition of reactor 3 and 11 clustered partially on the upper right quadrant while the nucleotide compositions of the other treatments were found to reveal dissimilarity.