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A circular phylogenetic tree showing the relationships between 100 bacterial strains. The branches are colored blue and orange, representing two distinct groups. The strains are labeled with their accession numbers around the perimeter of the tree.

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A circular phylogenetic tree representing the genetic relationships between 1000 human genomes. The tree is color-coded by population: African (red), European (blue), and Asian (grey). The branches radiate from a central point, with labels for various populations and individuals around the perimeter, such as 'AFR1', 'EUR1', 'AS1', 'AFR2', 'EUR2', 'AS2', etc. The tree shows a clear clustering of individuals from the same population, with African individuals forming a distinct cluster, followed by European and then Asian individuals.

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t p = 0.051, Wilcoxon p = 0.139

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