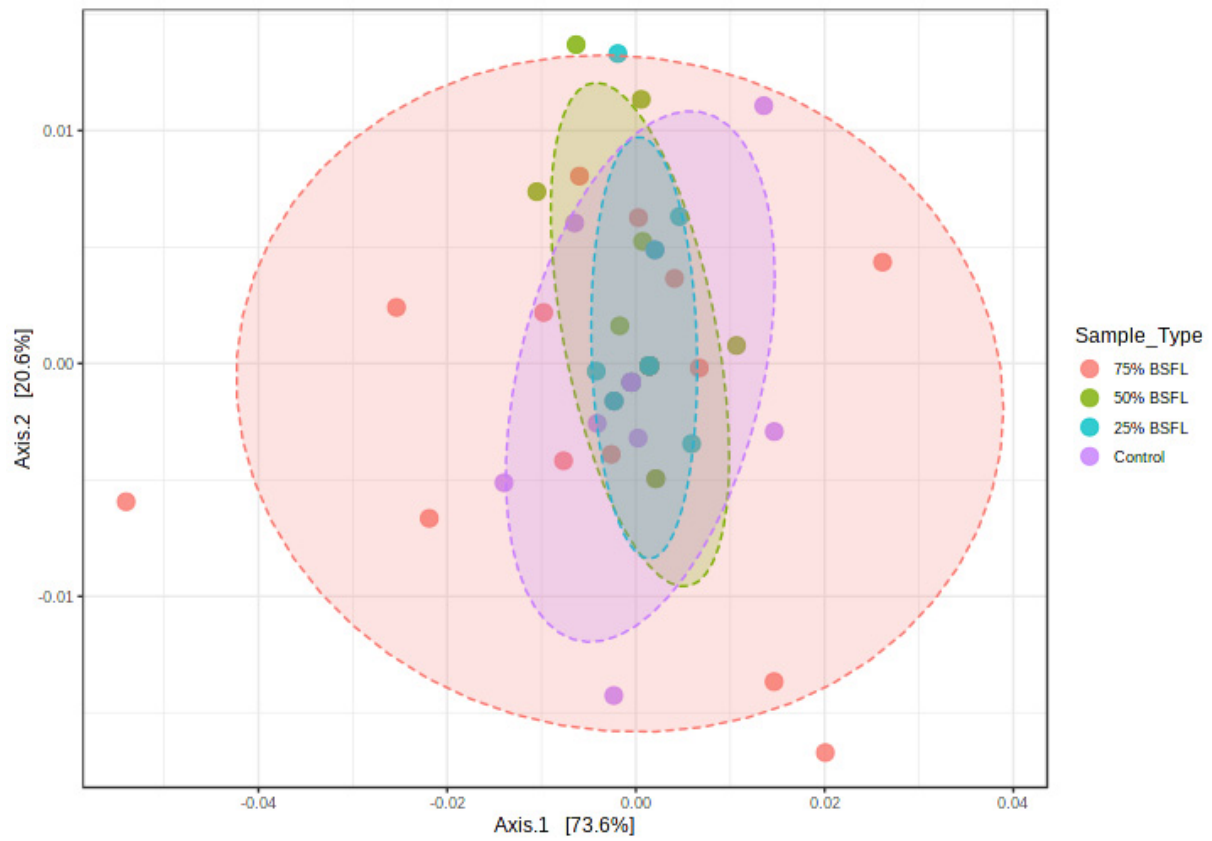


**Figure S1: Cumulative relative composition of bacteria Operational Taxonomic Units at family level observed across the different dietary treatments. The top 30 families with high abundance were selected.**



**Figure S2: Beta diversity PCoA plot based on Unifrac distance between the different dietary treatments. [PERMANOVA]  
R-Squared: 0.049; P-value > 0.001**