

Supplementary Figures:

Relationship of tongue coating microbiome on volatile sulfur compounds in healthy and halitosis adults

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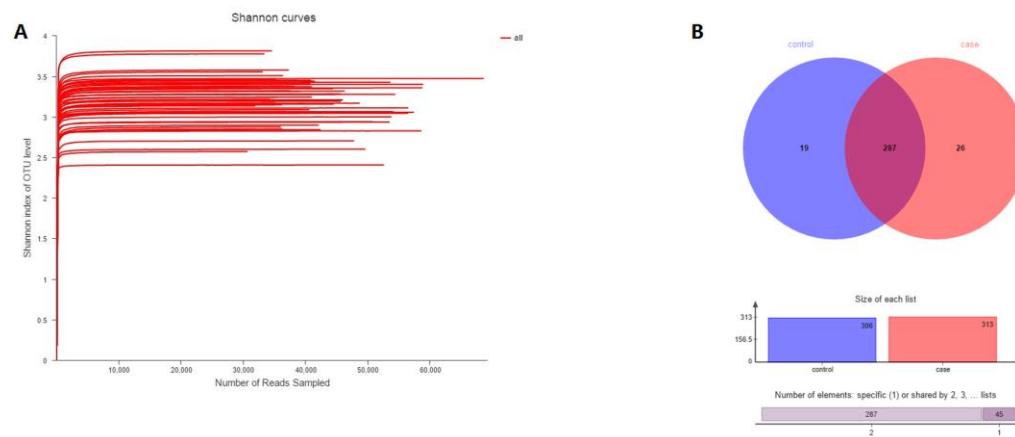


Figure S1: Overall sequencing data. (A) The rarefaction curves determined by the Shannon index were generated to evaluate sequencing depth. (B) Venn diagram-depicted OTU distributions.

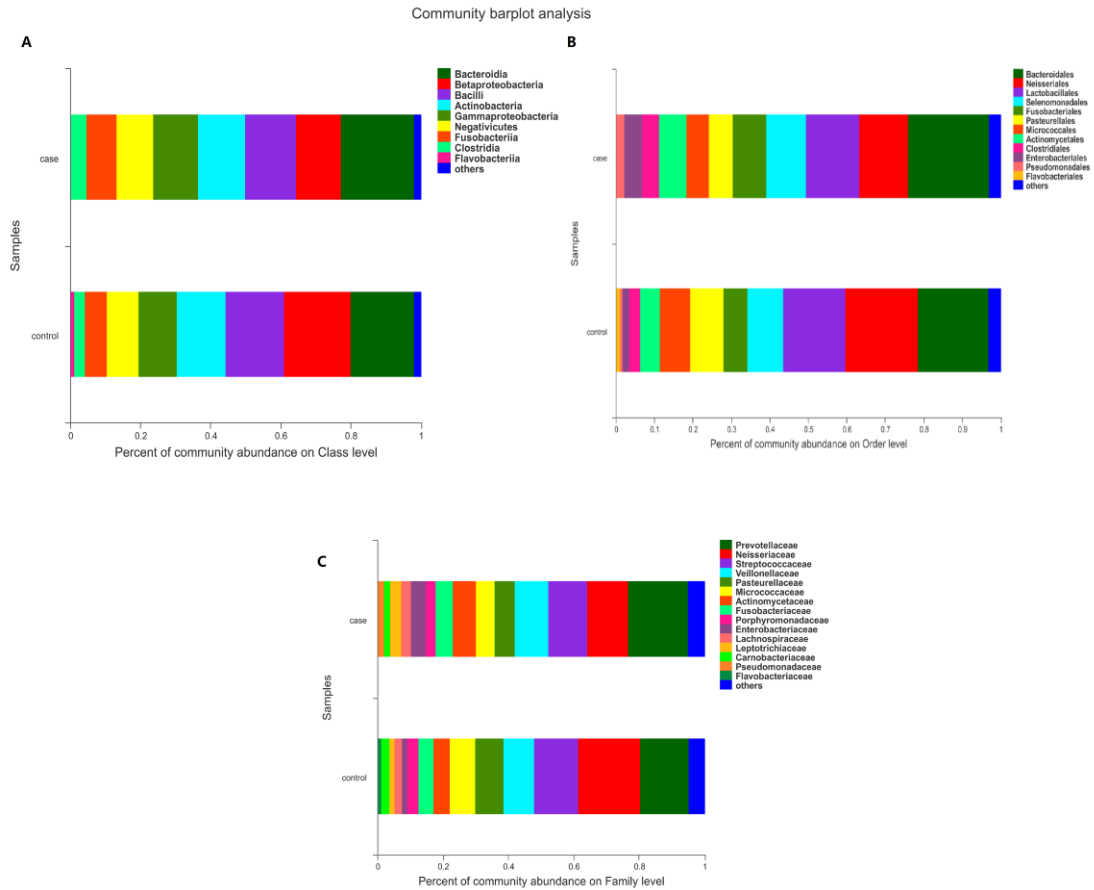


Figure S2: Composition of tongue coating bacterial communities across samples at different levels. (A) Community bar-plot analysis showed a relative abundance of bacteria at the class level in the halitosis and control groups. (B) Community bar-plot analysis showed the relative abundance of bacteria at the order level in the halitosis and control groups. (C) Community bar-plot analysis showed the relative abundance of bacteria at the family level in the halitosis and control groups.