Supplementary Material

# Supplementary Figures

Fig S1

**Supplementary Figure 1.** Rarefaction curves of the 192 oral samples.

Fig S2

**Supplementary Figure 2.** Oral microbiota composition. (A) Comparison of the oral microbiota composition between the reflux esophagitis (RE) and the healthy groups at the genus level. (B-C) The influence of *Hp* infection on the oral microbiota of reflux esophagitis patients. The alpha diversity of the RE group was not different from that of the C, REHpn, and REHpp groups, *P* > 0.05. (D) Comparison of oral microbiota composition between the REHpn and REHpp groups at the phylum level.