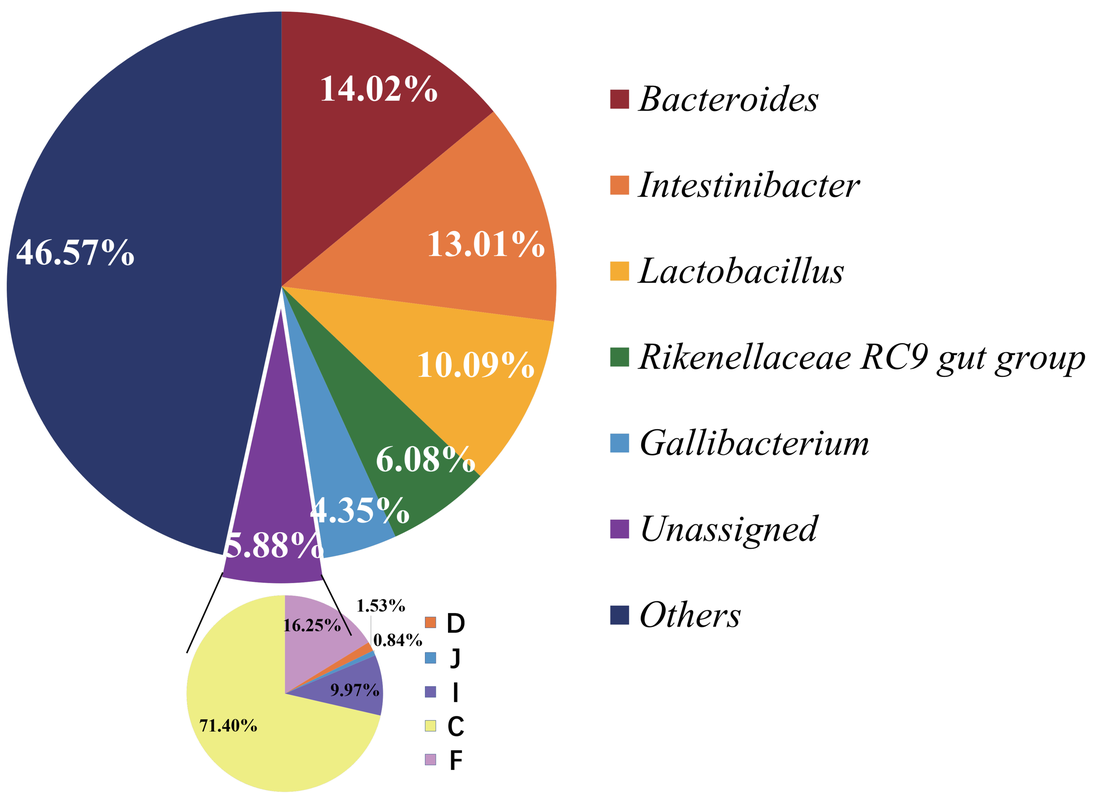
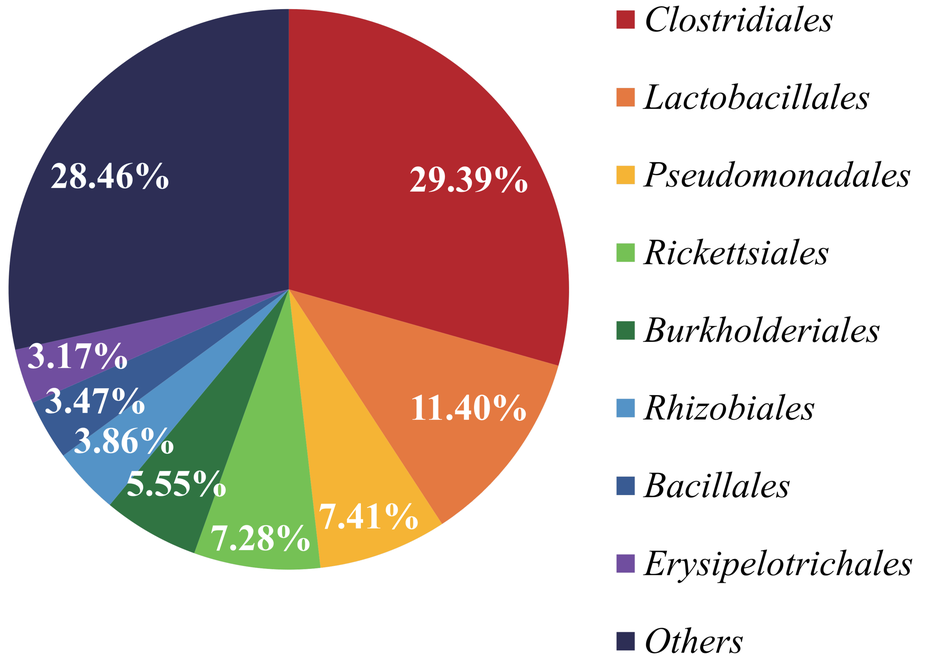
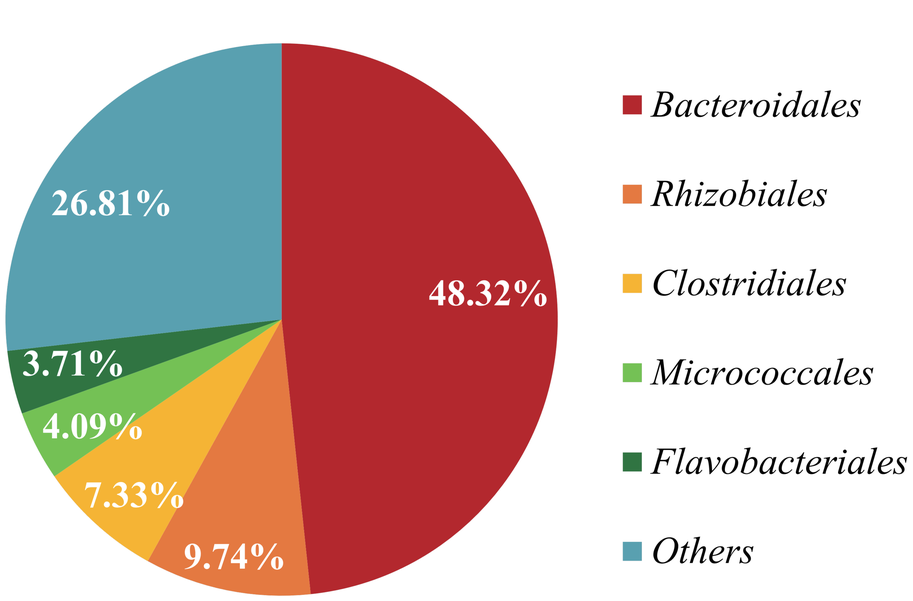
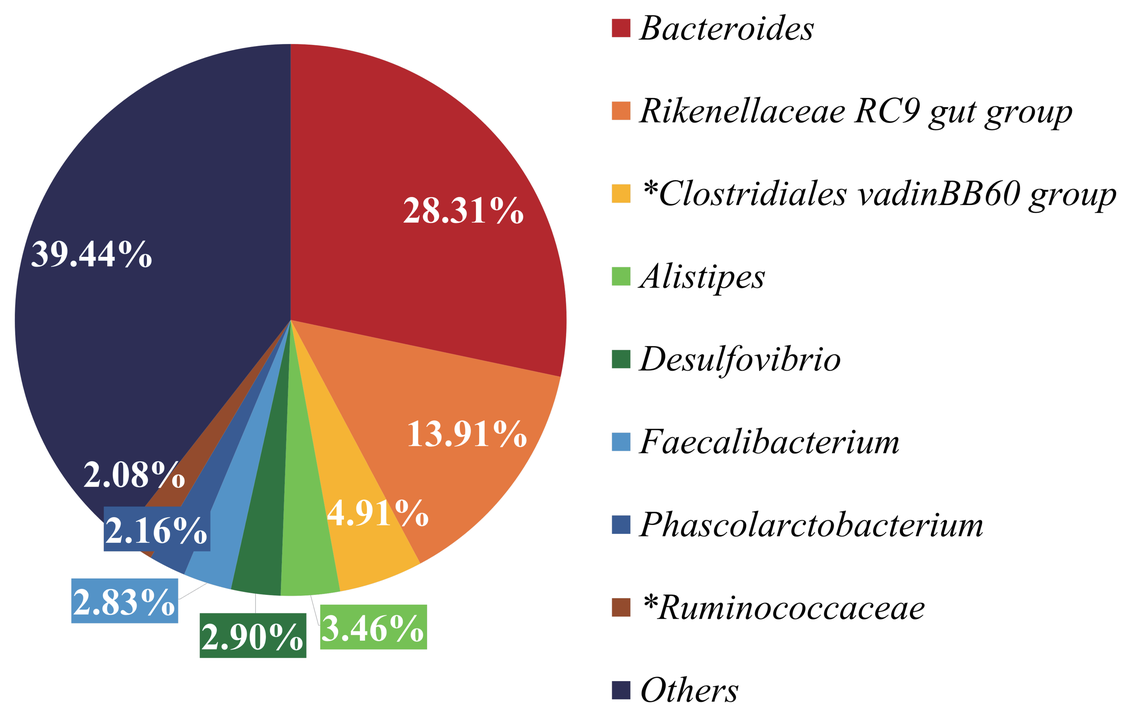
SUPPLEMENTARY FIGURES









Supplementary Figure S1 The predominant taxa. (A) The dominant taxa of the core microbiota. The sequences of unassigned taxa are partitioned to different sampling sites (duodenum, jejunum, ileum, ceca and feces) shown in the small pie chart. (B) The predominant taxa exclusively shared between feces and small intestine. (C) The predominant taxa exclusively shared between feces and cecum. (D) The dominated taxa in cecum samples.