Supplementary Figure 3. Abundance of bacterial OGAs-encoding genes in 1130 metagenomic samples from healthy people. (**A**) The overall abundance of OGAs-encoding genes. (**B**) The abundance of OGAs-encoding genes according to different phylum. (**C**) The abundance of genes that encode OGAs with or without NAGidase domain. (**D**) The abundance of OGAs-encoding genes in different genders. (**E**) The abundance of OGAs-encoding genes in samples from different countries. (**F**) Correlation between the abundance of OGAs-encoding genes and BMI (upper) or age (lower) calculated by Spearman correlation analysis.