Supplementary Figure S1. (A, B) The gut microbiota between the HPC group (n = 36) and PE patients without comorbidities of chronic hypertension or gestational diabetes (n = 14) differed significantly at the phylum and genus levels. *p < 0.05, **p < 0.01, ***p < 0.001.
Supplementary Figure S2. (A, B) When compared between the HPC group (n = 36) and PE patients without comorbidities of chronic hypertension or gestational diabetes (n = 14), the faecal butyric acid levels did not markedly differ, whereas the valeric acid level decreased significantly in patients with PE.