

Supplementary Materials

Supplementary Figure S1

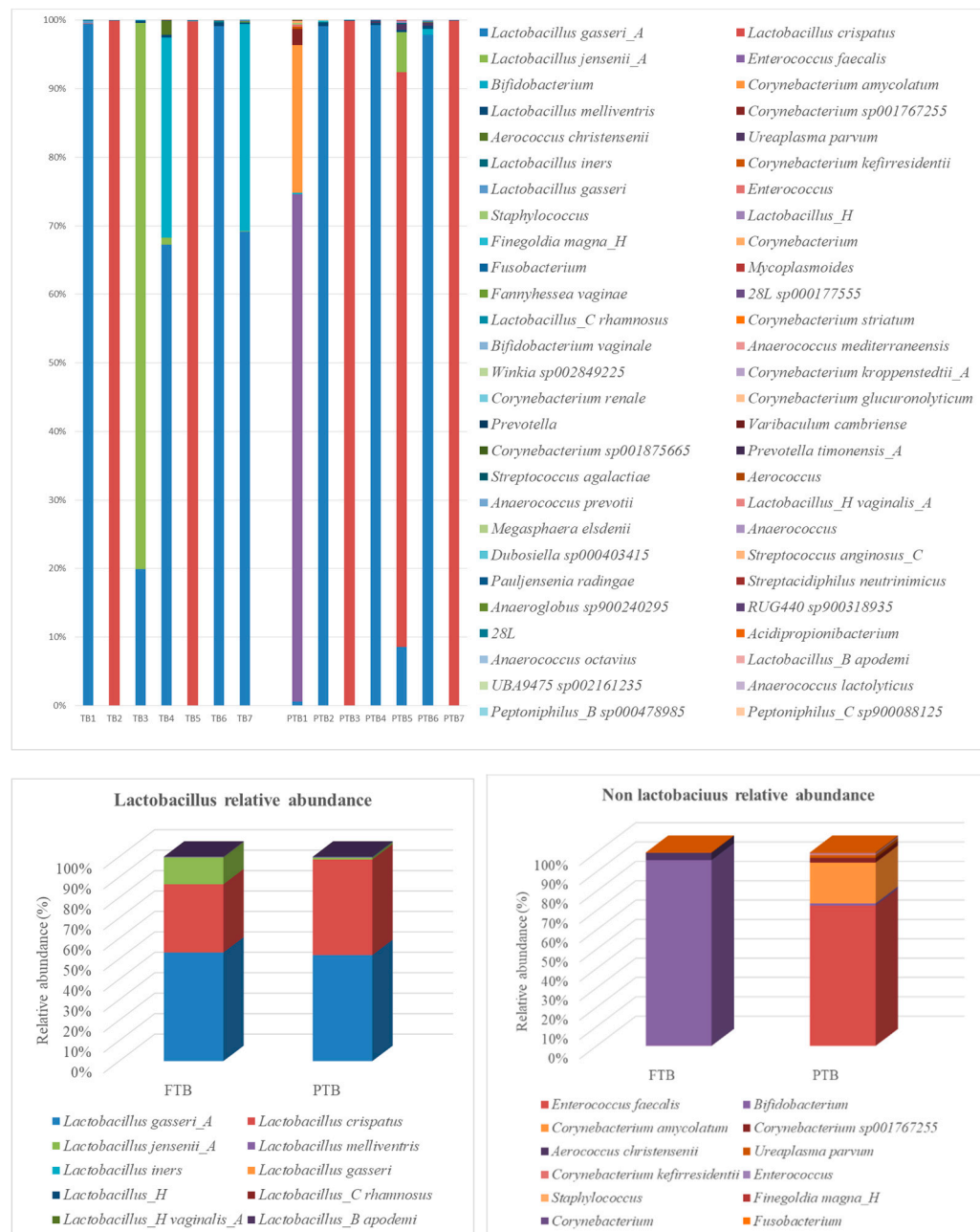


Figure S1. Microbiota profiling of cervicovaginal fluid (CVF) samples of full-term birth (FTB, n=7) and preterm birth (PTB, n=7) groups via next generation sequencing of the V1–V9 region.

Supplementary Figure S2

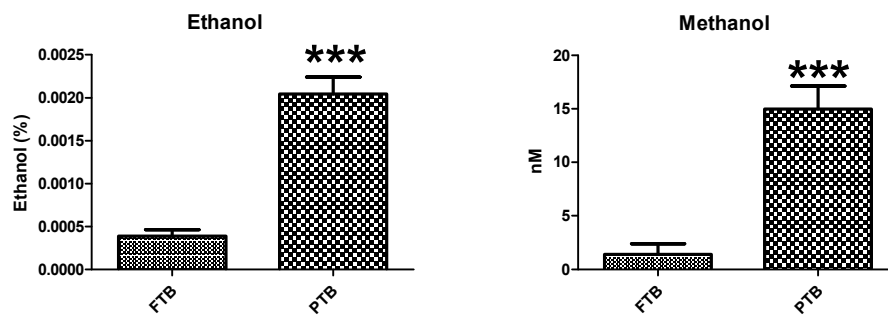


Figure S2. Metabolite analysis in cervicovaginal fluid (CVF) samples. Metabolites measured in CVF samples of full-term birth (FTB, $n = 20$) and preterm birth (PTB, $n = 20$) groups using the assay kits. Data are presented as the mean \pm standard deviation, and $p < 0.05$ was considered statistically significant; *** $p < 0.001$.