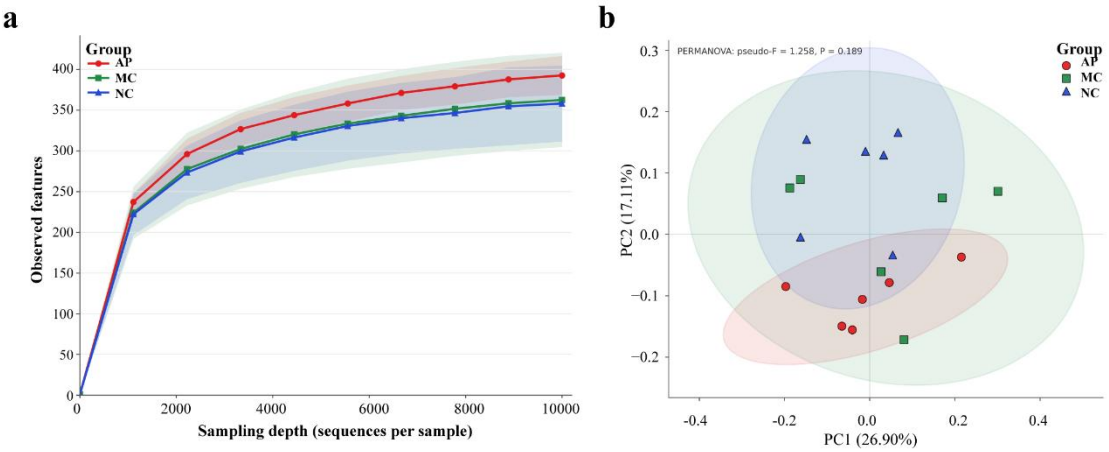


Fig. S1. Sequencing depth and beta-diversity analysis of gut microbiota among the NC, MC, and AP groups. (a) Rarefaction curves based on observed features. (b) Principal component analysis (PCA) of gut microbial community composition.