

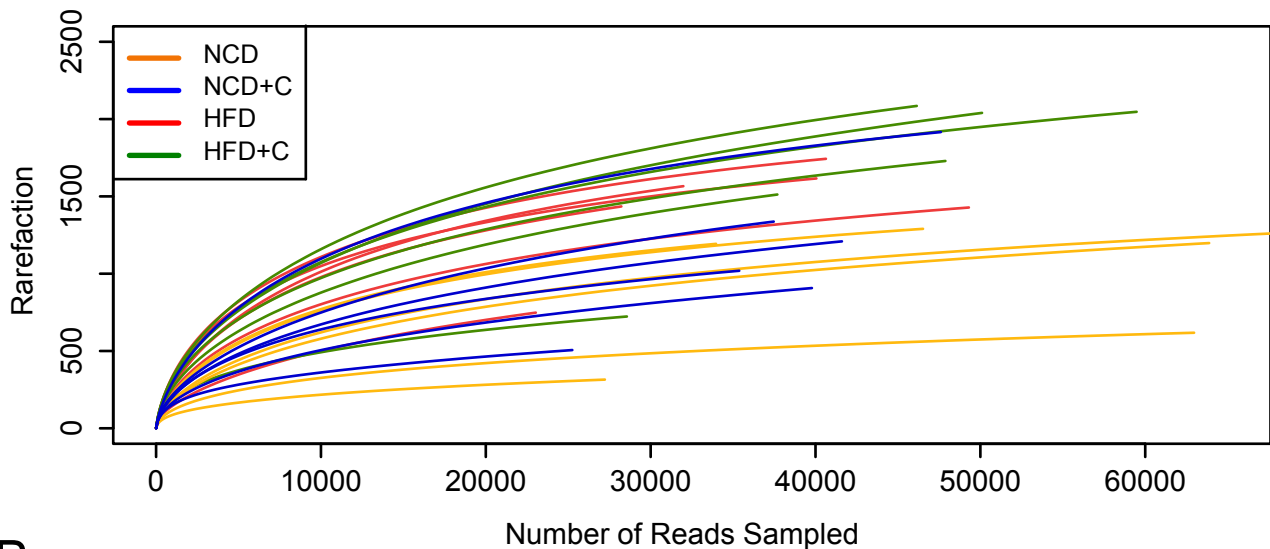
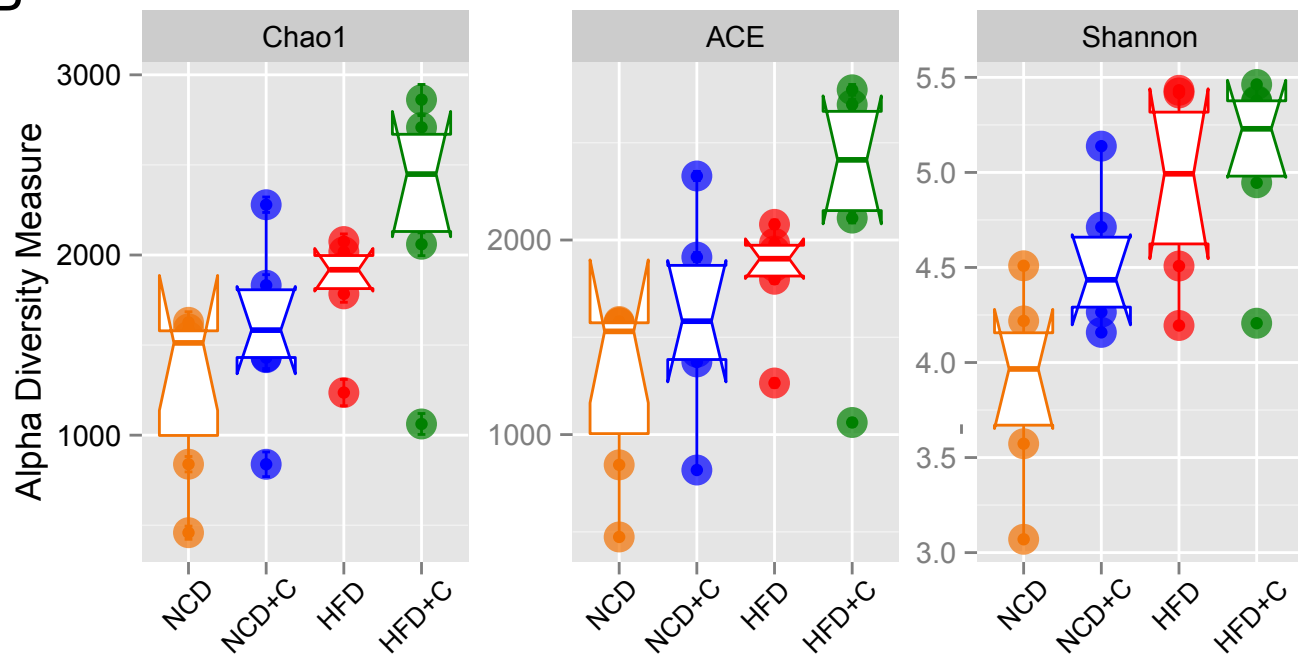
A**B**

FIG S4. CAP beneficially alters gut microbiota. Rarefaction curves of sequencing samples (A) and α -diversity (B) of gut microbial communities assessed by Chao1, ACE and Shannon α -diversity index, respectively (data are presented as median with inter-quartile range).