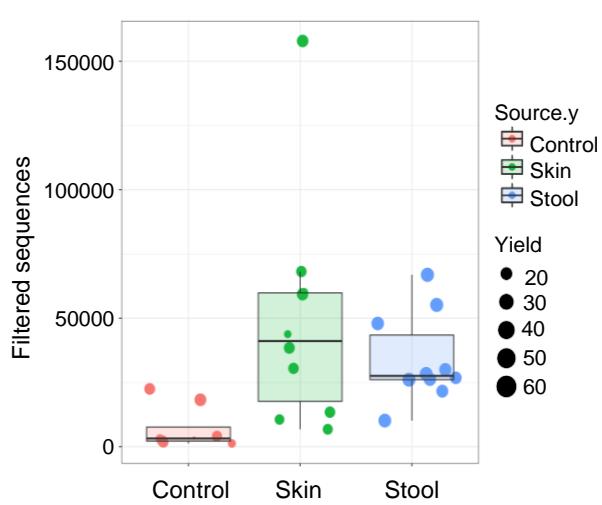
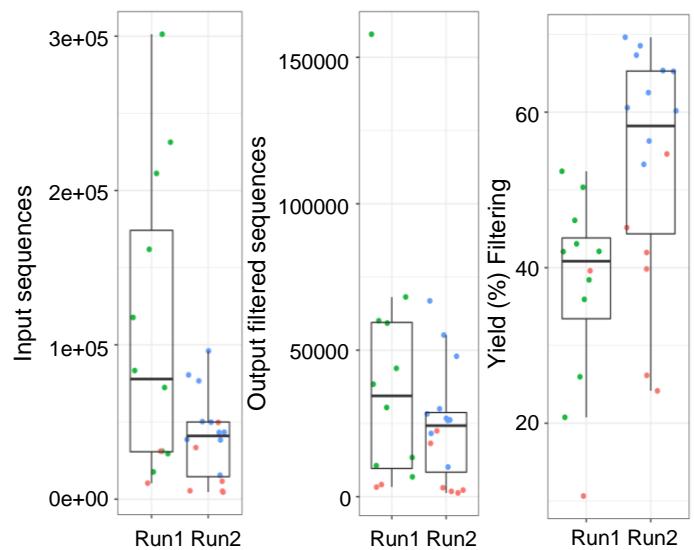
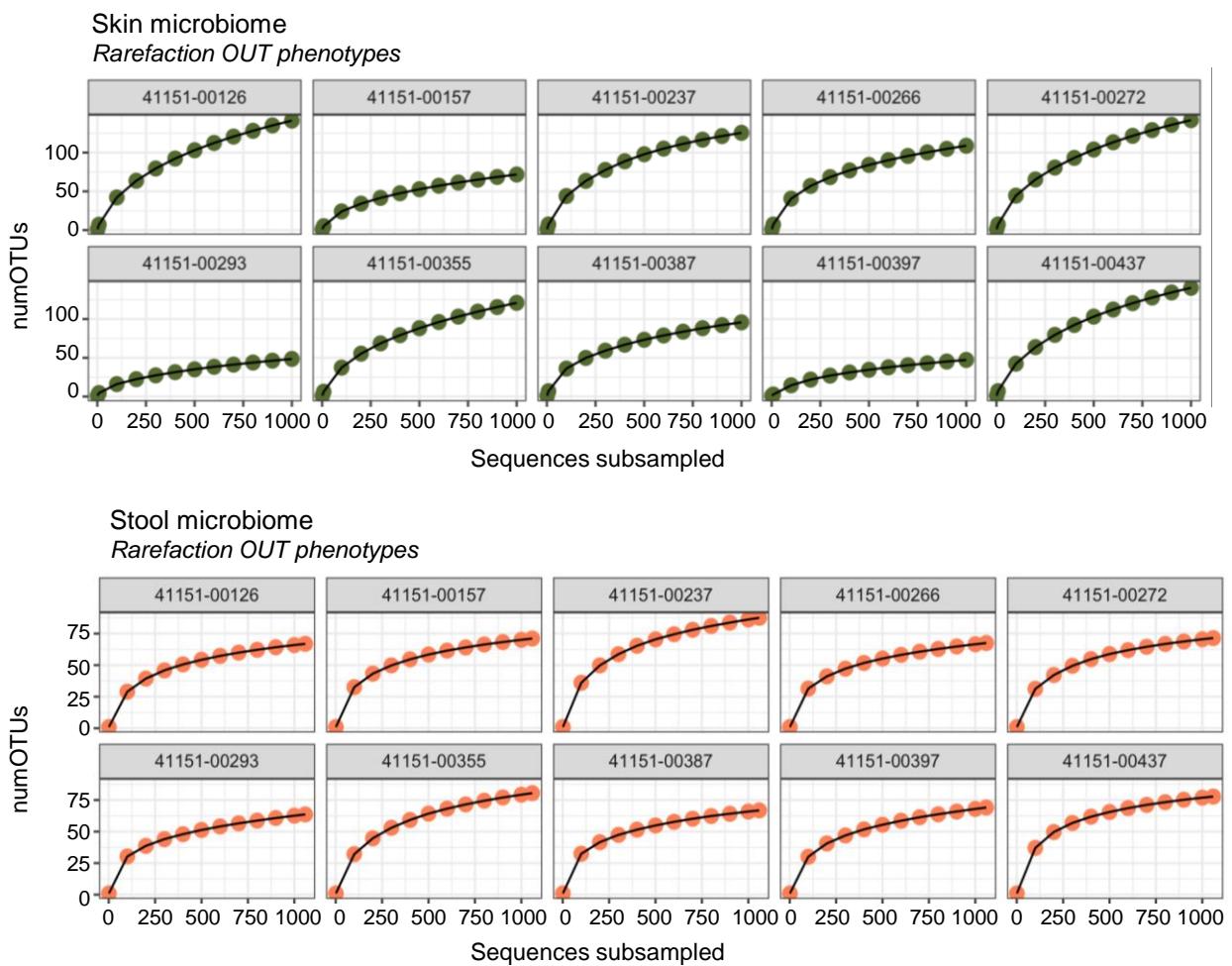


A**B****C**

Supplemental Figure 1. Sequences quality control for microbiota analyses **(A)** Total number of filtered sequences per sample in control, skin and stool samples. **(B)** From left to right: Distribution of the number of total raw sequences (Input sequences), filtered sequences (Output filtered sequences), and filtering yield values per each type of sample, grouped by the two independent sequencing batches (Run1 and Run2). **(C)** Rarefaction curves for skin and stool samples. Panel identifiers correspond to different individuals.