



Supplementary Figure 1. Bacterial community and diversity comparison in the control and ESKD groups. Beta diversity did not show different microbial compositions between the control group and the ESKD group in the (A) antecubital fossa and (B) shin. Permutational multivariate analysis of variance was performed for statistical comparisons of samples in the two groups. Beta diversity analysis was performed using Bray-Curtis, Jensen-Shannon, generalized UniFrac, and UniFrac metrics. ESKD, end-stage kidney disease.