Supplementary material to article by Y. Deng et al. “Patients with Acne Vulgaris Have a Distinct Gut Microbiota in Comparison with Healthy Controls”

Fig. S1. Difference in microbial structure among acne patient subgroups of different severity. (a) Principal coordinates analysis (PCoA) plot with different relative abundances of operational taxonomic units (OTUs) among patient subgroups. (b) Heatmap with weighted UniFrac phylogenetic distances of the microbial taxa among patients divided by disease severity.