

Supplemental Figure 1: Gut microbiota in T2D patients and healthy controls. [A] Phylogenetic structure of the gut microbiota ecosystem of T2D patients (red) and healthy controls (blue); colors were assigned for all families detected. [B] Rarefaction curves with different alpha-diversity metrics: the Chao 1 measure of microbial richness, observed species, the Shannon index of biodiversity and the Faith's phylogenetic diversity (PD w.t.).