Gut Microbiome Profiles and Associated Metabolic Pathways in HIV-Infected Treatment-Naïve Patients

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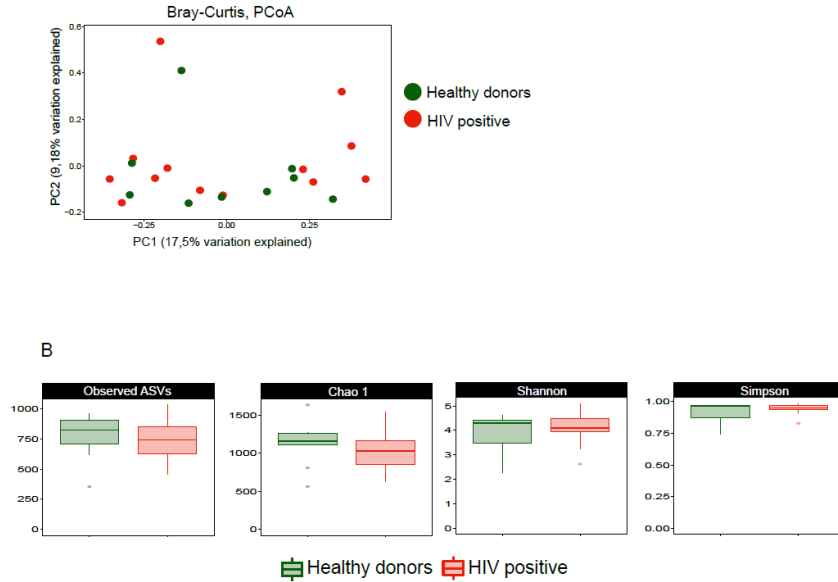
Supplementary Figures:

Figure 1. Comparison of gut microbiota of HIV-1 positive and healthy donors. PCoA of Bray Curtis distances among groups (a). Total number of ASVs observed, Richness (Chao1) and diversity (Shannon and Simpson) indices (b).
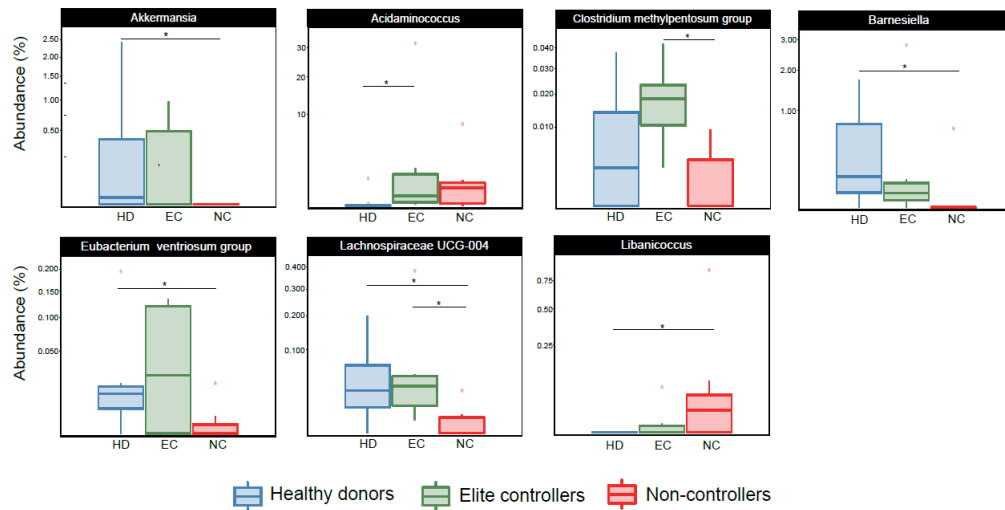

Figure S2. Comparison of gut microbiota of healthy donors (HD), elite controllers (EC) and Non-controllers (NC). The relative abundance of genera identified by LEfSe as being differentially abundant between groups were compared using the Kruskal-Wallis test. *P values <0.05.