

SUPPORTING INFORMATION

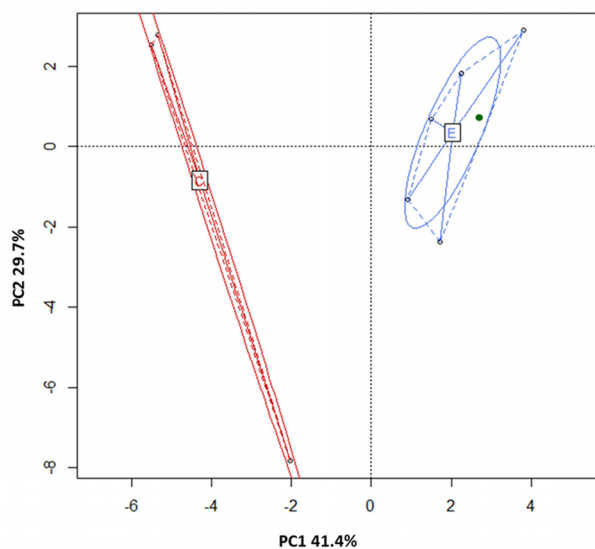


Figure S1. Centenarians and elderly subjects differ for gut microbiota composition. PCoA of Euclidean distances between genus-level taxonomic profiles. Subjects are colored as in Figure 1.

Please browse the full text version of this manuscript to see the Supplementary Table “KEGG Orthologous (KO) genes of core microbiome which correlated with ageing”.