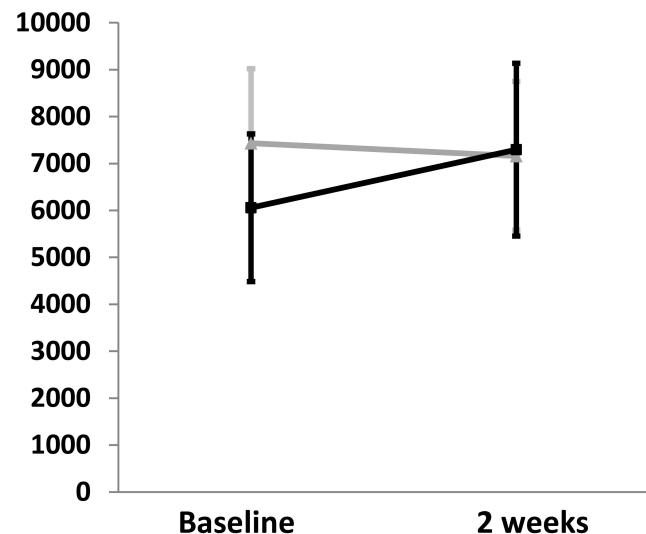


Supplementary figure 4

—▲— SIT
—■— MICT

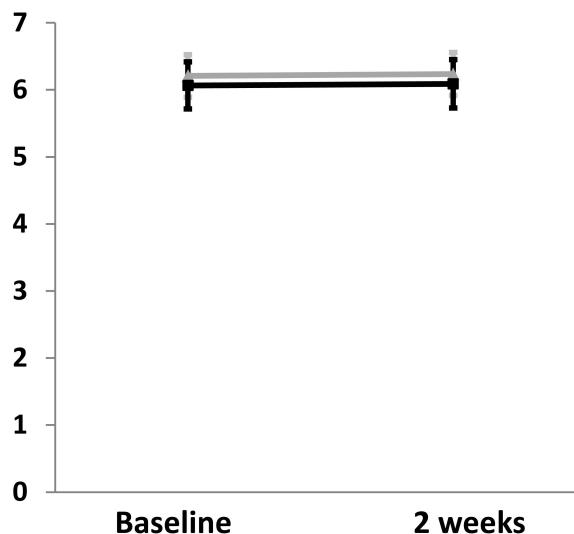
a)

CHOA 1



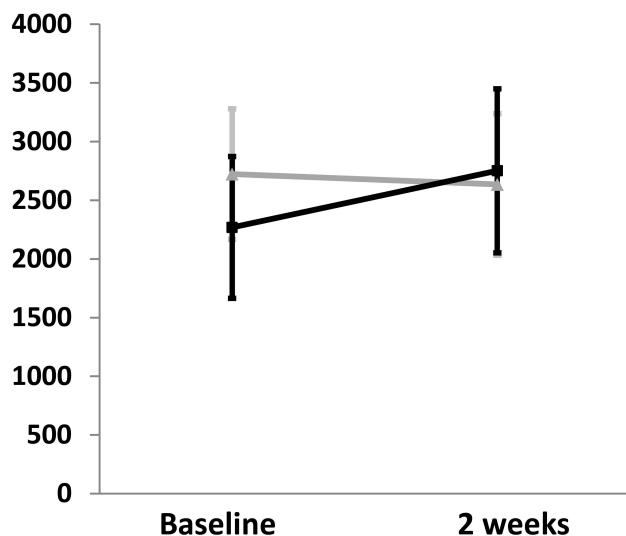
b)

SHANNON



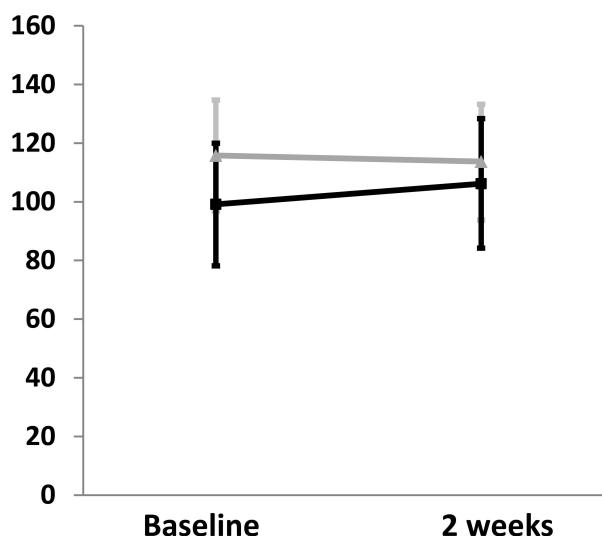
c)

OBSERVED OTUS



d)

PD WHOLE TREE



Supplementary figure 4: Changes in microbiome richness and diversity. a) Chao1; b) Shannon; c) observed otus and d) PD whole tree). All values are expressed as model-based means and bars are confidence intervals [95 % CI]