Figure S6. RDA plot of microbiota composition at genus level in the non-secretors (white) and the secretors (black) (A) and among the FUT2 genotypes, AA (white), AG (grey) and GG (black) (B). Plots are based on random subsample of the pyrosequencing data set (2500 sequences per sample, total 60,000 sequences). The centroids of each group are indicated by triangles. P-values show statistical significance in ANOVA test.