



Figure S5. RDA plot based on the OTUs detected in the non-secretors (white) and the secretors (black) (A) and among the *FUT2* genotypes, AA (white), AG (grey) and GG (black) (B). Threshold of 97% similarity was used for clustering of the sequences into OTUs. The centroids of each group are indicated by triangles. P-values show statistical significance in ANOVA test.