Supplemental figure 1. Hierarchical clustering of the HITChip microbiota profiles among the participants (n=51). Dash lines indicate participants in the WWB group and the solid lines participants in the RB group. Numbers denote participants, and the two branches of the tree represent microbiota composition of the pre- and post-intervention samples, respectively. HITChip, Human Intestinal Tract Chip; RB, rye bread; WWB, white wheat bread.