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Supplementary Material

Figure S1 Dietary intake during the entire experimental phase.

Figure S2 Changes in the abundance of selected genera.

Table S1 Changes in body weight during the entire experimental phase.

Table S2 Differentially expressed metabolites.

 Table S3 Enriched KEGG pathways in the GOS-fed group compared to the control.

Table S4 Prediction of differences in the metagenome functional content between GOS-fed and control mice using PICRUSt.