



- CON
- FMT1
- FMT2

Adjusted R-squared

Variable	Adjusted R-squared
FMT	0.154
SOW	0.064
Intersect	0
Residuals	0.785

db-RDA2: 22.29 %

db-RDA1: 66.14 %

Enterococcus hirae

Enterococcus faecium

Bacillus cereus

Streptococcus suis

Enterobacter cloacae

Enterococcus faecalis

Streptococcus viridans

Streptococcus gallolyticus

Escherichia coli

Faecalicoccus pleomorphus

Rothia nasimurium

Shigella sonnei

Clostridioides difficile

Lactobacillus crispatus

Lactobacillus amylovorus

Limosilactobacillus reuteri