

Table S6. Bacterial families in stool samples with statistically significant changes in the abundance between baseline and follow up samples after fecal microbiota transplantation (assessed by Metastats).

Family	Day	Abundance (%)	P-value
<i>Bacteroidaceae</i> ↑	Baseline	0.9 ±0.6	
	Day1	12.2 ± 5.7	0.03
	Day 7	13.5 ± 6.6	0.04
	Day 30	23.0 ± 11.2	0.03
	Day 90	19.6 ± 4.9	0.001
<i>Enterococcaceae</i> ↓	Baseline	5.7 ±2.4	
	Day 7	0.1 ±0.1	0.01
	Day 90	0.1 ±0.1	0.01
<i>Enterobacteriaceae</i> ↓	Baseline	25.8 ± 10.6	
	Day 30	3.3 ± 2.6	0.02
	Day 90	0.1 ± 0.1	0.02

↑relative increase in abundance

↓relative decrease in abundance