

Table S5. Bacterial families in mucosal 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	1.5±1.2	
	Day1	16.0± 4.9	0.005
	Day 7	11.9 ± 4.0	0.02
	Day 30	18.2 ± 6.2	0.007
	Day 90	15.0 ± 6.1	0.02
<i>Enterococcaceae</i> ↓	Baseline	6.5 ±2.4	
	Day 7	0.2 ±0.2	0.01
	Day 90	0.1 ±0.1	0.009
<i>Turicibacteraceae</i> ↑	Baseline	0	
	Day 7	1.2 ± 1.1	0.01
<i>Clostridiales Family XIII</i> ↑	Baseline	0	
	Day 7	1.0 ± 1.0	0.01
<i>Enterobacteriaceae</i> ↓	Baseline	20.7 ± 10.8	
	Day 90	0.8 ± 0.4	0.047

↑relative increase in abundance

↓relative decrease in abundance