

Taxa	REC6	REM6	p-value
Bacilli	3,781±1,309	0,897±0,228	0,023
Lactobacillales	3,265±1,088	0,771±0,19	0,015
Enterococcaceae	0,597±0,361	0,009±0,008	0,011
<i>Enterococcus</i>	0,597±0,361	0,009±0,008	0,011
Bacteroidaceae	13,145±4,893	29,586±5,346	0,04
<i>Bacteroides</i>	13,145±4,893	29,586±5,346	0,04
<i>Dorea</i>	2,445±0,726	5,996±0,943	0,008
<i>Ruminococcus</i>	0,213±0,105	2,348±0,983	0,036
<i>Dialister</i>	0±0	0,88±0,516	0,028

Supp. Table 3: Differentially represented taxa between samples from patients in remission or with endoscopic recurrence at 6 months after ICR

Shade less and grey shade lines indicate respectively increased taxa in patients with disease recurrence (REC) or remission (REM). Statistics was assessed by Wilcoxon test.