

Supplementary Table 1. Experimental diets and fatty acid composition

Study Diets	Low-fat (AIN-93M)	Saturated- Milk Fat	Saturated- Lard Fat	PUFA (Safflower oil)
Fat (%kcal)	5	38	38	38
-Saturated (%total)	0.6	65	39	9
-PUFA	3.9	3.5	11	78
Protein	19	16	16	16
Carbohydrate	66	47	47	47
Micronutrients/Fiber	*ND	*ND	*ND	*ND

		Milk		Safflower
		Fat	Lard	Oil
4:0	Butyric	3.8		
6:0	Caproic	2.3		
8:0	Caprylic	1.1		
10:0	Capric	2		
11:0		0.1		
12:0	Lauric	3.1		
14:0	Myristic	11.7	1.5	
14:1		0.8		
15:0		1.6		
15:1		0		
16:0	Palmitic	26.2	24.8	6.5
16:1	Palmitoleic	1.9	3.1	
17:0	Margaric	0.7		
17:1		0.2		
18:0	Stearic	12.5	12.3	2.4
18:1	Oleic	28.2	45.1	13.1
18:2	Linoleic	2.9	10	78
18:3	Linolenic	0.5	0.1	
18:4		0		
20:0	Arachidic	0	0.2	
20:1		0.2	1.3	
20:2		0	0.1	
20:3				
20:4	Arachidonic	0.1	0.4	
20:5	EPA			
21:5		0		
22:1		0		
22:4				
22:5	DPA	0		
22:6	DHA	0		

**Supplementary Table 2. Assignment to the genus level of 324 near full-length 16s rRNA sequences**

	Low-fat	PUFA	MF
Genus			
<i>Bacteroides</i>	8	31	34
<i>Tannerella</i>	1	1	2
<i>Parabacteroides</i>	5	48	39
<i>Allobaculum</i>	20	20	7
<i>Anaerotruncus</i>	10	2	2
<i>Clostridium</i>	25	3	
<i>Peptostreptococcaceae Incertae Sedis</i>	2	1	4
<i>Lachnospiraceae Incertae Sedis</i>	10	1	10
<i>Erysipelotrichaceae Incertae Sedis</i>	24	1	
<i>Bilophila</i>			6
<i>Lactobacillus</i>			5
<i>Alistipes</i>			3
<i>Papillobacter</i>	2	1	1
<b>Sum</b>	<b>107</b>	<b>109</b>	<b>111</b>

**Supplementary Table 3. Sample metadata**

	<b>Clone Library</b>	<b>454-Pyrosequencing</b>
<b>Sequencing depths and # reads</b>	<ul style="list-style-type: none"><li>• 3525 average total sequences</li><li>• 110 sequences/sample</li></ul>	<ul style="list-style-type: none"><li>• 3000 reads/sample</li></ul>
<b>Total Mice</b>	32	38
<i>C57BL/6</i>	10	23
<i>IL10<sup>-/-</sup></i>	22	15

**Supplementary Table 4. Correlation between abundance of *B. wadsworthia* and colitis score calculated by Pearsons correlation coefficient**

	<b>Correlation of Bw abundance and colitis score (R-value)</b>
<b>Diet treatment groups</b>	
SPF IL10 <sup>-/-</sup> + LF	0.36
SPF IL10 <sup>-/-</sup> + PUFA	0.26
SPF IL10 <sup>-/-</sup> + MF	0.92
GF IL10 <sup>-/-</sup> + LF	-
GF IL10 <sup>-/-</sup> + PUFA	-
GF IL10 <sup>-/-</sup> + MF	0.94
<b>Bile treatment groups</b>	
SPF IL10 <sup>-/-</sup> + PBS	0.27
SPF IL10 <sup>-/-</sup> + GC	0.54
SPF IL10 <sup>-/-</sup> + TC	0.96
GF IL10 <sup>-/-</sup> + PBS	-
GF IL10 <sup>-/-</sup> + GC	-
GF IL10 <sup>-/-</sup> + TC	0.95