

Table S1. Average number of reads and operational taxonomic units (OTUs) of 16S rDNA

<b>Samples*</b>	<b>No. of reads (standard deviation)</b>	<b>No. of screened reads (standard deviation)</b>	<b>No. of OTUs (standard deviation)</b>
<b>CTL</b>	42,992 (7,535)	35,421 (6,467)	282 (23)
<b>HFD</b>	48,882 (4,504)	38,212 (4,037)	188 (40)
<b>HFL</b>	53,485 (32,975)	43,857 (28,670)	154 (44)

*Notes:* (\*) Samples were named by sampling time (D0: Day 0, W1-6: Week 1-6), treatment (CTL: Normal diet, HFD: High-fat diet, HFL: HFD + Laminarin). 'A' and 'B' denote different cage.

Table S2. Number of differentially abundant OTUs and their read abundance during laminarin ingestion detected by comparing HFD and HFL using Metastats

Genera	HFD		HFL	
	No. of OTUs	No. of reads	No. of OTUs	No. of reads
<b>Bacteroides</b>	0	0	4	22558
<b>unclassified</b>	154	31403	60	12797
<b>Clostridium_XIVa</b>	0	0	2	1865
<b>Lactobacillus</b>	1	3	1	1775
<b>Parabacteroides</b>	0	0	1	1782
<b>Clostridium_XVIII</b>	0	0	1	2314
<b>Asaccharobacter</b>	0	0	1	181
<b>Proteus</b>	0	0	1	89
<b>Enterococcus</b>	0	0	1	103
<b>Allobaculum</b>	0	0	1	64
<b>Eubacterium</b>	0	0	1	106
<b>Blautia</b>	0	0	1	46
<b>Staphylococcus</b>	0	0	1	26
<b>Acinetobacter</b>	0	0	1	13
<b>Streptococcus</b>	0	0	1	17
<b>Johnsonella</b>	0	0	1	13
<b>Legionella</b>	0	0	1	5
<b>Exiguobacterium</b>	0	0	1	2
<b>Hydrogenophaga</b>	1	2	0	0
<b>Arcobacter</b>	1	2	0	0
<b>Emticicia</b>	1	2	0	0
<b>Clostridium_XIVb</b>	3	374	0	0
<b>Gemella</b>	1	5	0	0
<b>Oscillibacter</b>	5	871	0	0
<b>Butyricicoccus</b>	1	25	0	0
<b>Turicibacter</b>	1	20	0	0
<b>Clostridium_IV</b>	1	34	0	0
<b>Dorea</b>	1	59	0	0
<b>Pseudoflavonifractor</b>	1	50	0	0
<b>Enterorhabdus</b>	1	132	0	0
<b>Barnesiella</b>	1	183	0	0
<b>Lactococcus</b>	1	1366	0	0
<b>Alistipes</b>	1	266	0	0
<b>Bifidobacterium</b>	1	717	0	0
<b>Mucispirillum</b>	1	417	0	0
<b>Odoribacter</b>	1	451	0	0
<b>Clostridium_XI</b>	1	2234	0	0
<b>Total</b>	180	38616	81	43756

Table S3. Number of differentially abundant OTUs and their read abundance after termination of laminarin ingestion detected by comparing HFD and HFL using Metastats

<b>Genera</b>	<b>HFD</b>		<b>HFL</b>	
	<b>Number of OTUs</b>	<b>No. of reads</b>	<b>Number of OTUs</b>	<b>No. of reads</b>
<b>Parabacteroides</b>	0	0	1	684
<b>unclassified</b>	11	3653	8	1099
<b>Pseudoflavonifractor</b>	0	0	1	57
<b>Eubacterium</b>	0	0	1	28
<b>Blautia</b>	0	0	1	7
<b>Streptococcus</b>	0	0	1	5
<b>Oscillibacter</b>	1	36	0	0
<b>Clostridium_sensu_stricto</b>	1	117	0	0
<b>Barnesiella</b>	1	273	0	0
<b>Alistipes</b>	1	281	0	0
<b>Total</b>	15	4360	13	1880