

Table S1: Effects of COST on body weight and various physiological and biochemical parameters in mice with disorders of lipid metabolism, correlated with Figure 1 ($n = 6$). Data are expressed as mean \pm SEM and statistically analyzed via one-way ANOVA; * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$; **** $p < 0.0001$.

	CTRL	MOD	COST-H	COST-M	COST-L
Body weight(g)	26.76 \pm 0.083***	33.62 \pm 0.48	32.28 \pm 0.56	33.10 \pm 0.60	33.64 \pm 0.89
Serum TC(mmol/L)	3.90 \pm 0.13****	5.66 \pm 0.20	4.35 \pm 0.17****	4.59 \pm 0.12**	4.98 \pm 0.26*
Serum TG(mmol/L)	0.33 \pm 0.04****	0.67 \pm 0.03	0.48 \pm 0.02**	0.47 \pm 0.03**	0.52 \pm 0.04*
Serum HDL-C(mmol/L)	7.84 \pm 0.31****	4.70 \pm 0.06	6.84 \pm 0.26****	6.42 \pm 0.25***	5.52 \pm 0.10
Serum LDL-C(mmol/L)	1.83 \pm 0.07****	3.60 \pm 0.21	2.45 \pm 0.15***	2.67 \pm 0.15**	3.34 \pm 0.26
Serum NEFA (mmol/L)	0.53 \pm 0.01****	1.04 \pm 0.02	0.56 \pm 0.02****	0.61 \pm 0.023****	0.64 \pm 0.026****
Serum GLU (mmol/L)	5.51 \pm 0.11****	12.23 \pm 0.34	9.18 \pm 0.47****	10.85 \pm 0.053	11.62 \pm 0.55

Table S2: Mice body weight, serum and liver biochemical markers, and liver and adipose tissue weight during FMT, correlated with Fiure 3, Figure 4 and Figure 5 ($n = 6$).Data are expressed as mean \pm SEM and statistically analyzed via one-way ANOVA; * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$; **** $p < 0.0001$.

	STD	Control	Model	COST-H-FMT	COST-M-FMT	COST-L-FMT
Body weight(g)	28.84 \pm 0.55***	29.85 \pm 0.52*	32.49 \pm 0.50	26.19 \pm 0.65***	27.38 \pm 0.71***	27.88 \pm 0.65***
LiverWeight(g)	0.9807 \pm 0.05**	1.0156 \pm 0.04*	1.1457 \pm 0.03	0.9214 \pm 0.02***	0.9391 \pm 0.03**	0.8758 \pm 0.03***
Brown adipose tissue (g)	0.132 \pm 0.01	0.125 \pm 0.01	0.098 \pm 0.00	0.118 \pm 0.00	0.109 \pm 0.01	0.108 \pm 0.01
Subcutaneous adipose	0.199 \pm 0.01***	0.400 \pm 0.04*	0.557 \pm 0.04	0.207 \pm 0.03***	0.326 \pm 0.04***	0.352 \pm 0.05**
Perirenal adipose	0.071 \pm 0.01***	0.210 \pm 0.03***	0.422 \pm 0.04	0.150 \pm 0.02***	0.175 \pm 0.03***	0.194 \pm 0.03***
Epididymal adipose	0.344 \pm 0.02***	0.789 \pm 0.10***	1.325 \pm 0.09	0.540 \pm 0.10***	0.601 \pm 0.07***	0.652 \pm 0.08***
Fat/body ratio (%)	2.58%***	5.08%***	7.41%	3.88%***	4.43%***	4.68%***
Serum TC(mmol/L)	7.84 \pm 0.28***	6.89 \pm 0.52***	10.30 \pm 0.42	7.34 \pm 0.19***	7.71 \pm 0.37***	7.96 \pm 0.42**
Serum TG(mmol/L)	0.95 \pm 0.03*	0.83 \pm 0.13**	1.45 \pm 0.16	0.50 \pm 0.16***	0.77 \pm 0.07**	1.04 \pm 0.08
Serum HDL-C(mmol/L)	3.76 \pm 0.01***	3.14 \pm 0.16	2.56 \pm 0.16	4.26 \pm 0.15***	4.22 \pm 0.21***	3.61 \pm 0.21***
Serum LDL-C(mmol/L)	0.28 \pm 0.04***	0.95 \pm 0.15***	2.40 \pm 0.38	0.90 \pm 0.19***	0.87 \pm 0.08***	1.22 \pm 0.15**
Serum ALT(mmol/L)	82.72 \pm 3.88***	45.38 \pm 5.85***	131.16 \pm 10.71	50.16 \pm 2.39***	50.16 \pm 5.85***	46.99 \pm 0.48***
Serum AST(mmol/L)	84.30 \pm 3.80**	79.42 \pm 2.16	106.40 \pm 1.52	67.73 \pm 2.46***	76.90 \pm 2.98	77.10 \pm 2.78
Serum Glucose(mmol/L)	8.06 \pm 0.37**	7.23 \pm 0.44***	11.02 \pm 0.42	6.35 \pm 0.47***	8.02 \pm 0.22***	7.99 \pm 0.36***
Serum NEFA (mmol/L)	0.94 \pm 0.06***	1.07 \pm 0.02*	1.25 \pm 0.04	0.75 \pm 0.04***	0.86 \pm 0.04***	0.99 \pm 0.03***
Serum TBA (mmol/L)	2.31 \pm 0.16***	4.37 \pm 0.16	5.28 \pm 0.46	3.04 \pm 0.18***	3.30 \pm 0.14***	4.16 \pm 0.25*
Liver TG level(mmol/L)	3.47 \pm 0.51***	5.80 \pm 1.10**	9.04 \pm 2.42	2.96 \pm 0.31***	4.99 \pm 0.96**	5.04 \pm 1.46**
Liver TC level(mmol/L)	1.47 \pm 0.39**	1.98 \pm 0.15*	2.69 \pm 0.40	1.68 \pm 0.411**	1.70 \pm 0.23*	2.16 \pm 0.28
Liver LDL- C level	0.38 \pm 0.076**	0.44 \pm 0.097	0.67 \pm 0.060	0.38 \pm 0.087*	0.39 \pm 0.092*	0.47 \pm 0.11
Liver TBA level	59.13 \pm 8.89***	86.13 \pm 11.77*	123.33 \pm 18.36	66.90 \pm 13.43***	78.74 \pm 19.06**	83.98 \pm 22.45**

Table S3 Animal Feed Information

Feed	Protein (%)	Carbohydrate (%)	FAT (%)	Fiber (%)	Calcium/Phosphorus (%)
Standard formula feed	20	0	4.3	4.8	1.19
D12327 High- fat and high- sugar feed	23	46.1	20.4	5.5	5

Table S4 Primer sequence

Gene	Forward primer	Reverse primer
SREBP	5'-CAGCTATTGGCCTTCCTCAG-3'	5'-GGTTACTGGCGGTCACTGTC-3'
FAS	5'-ATGCACACTCTGCGATGAAG-3'	5'-TTCAGGGTCATCCTGTCTCC-3'
PPAR γ	5'-AGACCACTCGCATTCCCTTTG-3'	5'-CATTGGGTCACTCTTGTGA-3'
PPAR α	5'-GCAGCTCGTACAGGTCATCA-3'	5'-ACTGCCGTTGTCTGTCACTG-3'
CYP7A1	5'-GGGATTGCTGTGGTAGTGAGC-3'	5'-GGTATGGAATCAACCCGTTGTC-3'
FXR	5'-GCTTGATGTGCTACAAAAGCTG-3'	5'-GTGGTGATGGTTGAATGTCC-3'
Occludin	5'-ATGTCCGGCCGATGCTCTC-3'	5'-TTTGGCTGCTCTTGGGTCTGTAT-3'
Claudin	5'-TCTACGAGGGACTGTGGATG-3'	5'-TCAGATTCAGCTAGGAGTCG-3'
ZO1	5'-ACCCGAAACTGATGCTGTGGATAG-3'	5'-AAATGGCCGGGCAGAACTTGTGTA-3'
β -Actin	5'-CGTGAAAAGATGACCCAGA-3'	5'-GTCCATCACAATGCCTGT-3'

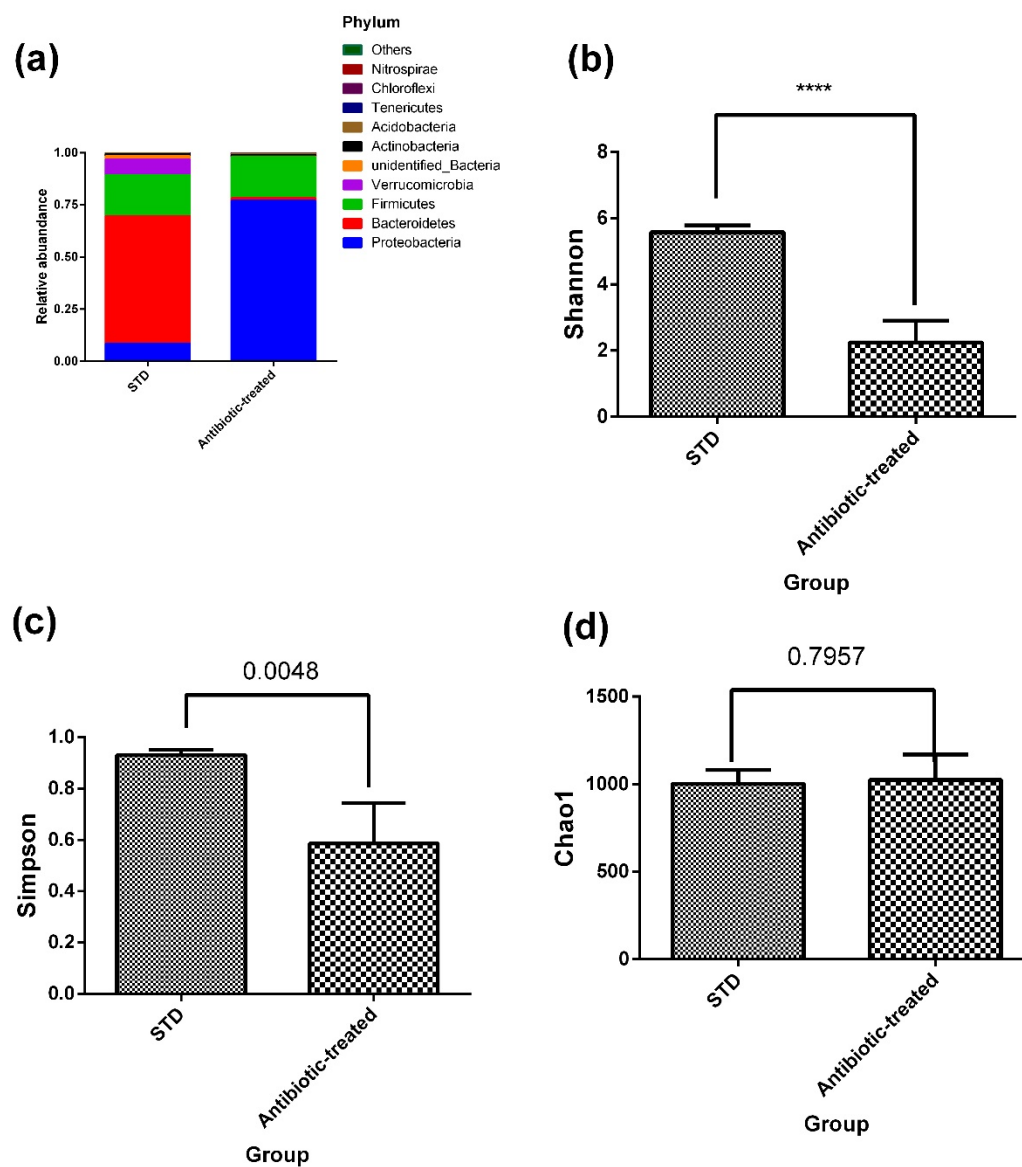


Figure S1: Species levels of phylum differences in mice following antibiotic intervention (a), and changes in Alpha index levels (b-d)($n = 6$).