

**Effects of konjac glucomannan intake patterns on glucose and lipid metabolism
of obese mice induced by high fat diet**

Sijia Zhu ^{a, b}, Jiyu Yang ^{a, b}, Pengkui Xia ^{a, b}, Sha Li ^{a, b}, Qi Wang ^{a, b}, Kaikai Li ^{a, b}, Bin Li ^{a, b}, Jing
Li ^{a, b, *}

^a College of Food Science and Technology, Huazhong Agricultural University, Wuhan 430070,
China

^b Key Laboratory of Environment Correlative Dietology, Huazhong Agricultural University,
Ministry of Education, China

* Corresponding author: Jing Li

E-mail: lijingfood@mail.hzau.edu.cn

Table S1 Diet formula and energy

Material	D12450J (10 kcal% Fat)	D12492 (60 kcal% Fat)		
	gm	kcal	gm	kcal
Casein	200	800	200	800
L-Cystine	3	12	3	12
Corn Starch	506.2	2024.8	0	0
Maltodextrin 10	125	500	125	500
Sucrose	68.8	275.2	68.8	275.2
Cellulose,BW200	50	0	50	0
Soybean Oil	25	225	25	225
Lard	20	180	245	2205
Mnereal Mx S10026	10	0	10	0
DiCalcium Phosphate	13	0	13	0
Calcium Carbonate	5.5	0	5.5	0
Potassium Citrate,1 H2O	16.5	0	16.5	0
Vitamin Mx V10001	10	40	10	40
Choline Bitartrate	2	0	2	0
FD&C Yellow Dye#5	0.04	0	0	0
FD&C Blue Dye#1	0.01	0	0.05	0
	gm %	kcal%	gm%	kcal%
Protein	19.2	20	26.2	20
Carbohydrate	67.3	70.0	26.3	20.1
Fat	4.3	10	34.9	59.9
Total		100		100
Total	1055.05	4057	773.85	4057

Table S2 Primer information of genes to be tested

Item		Base sequence (5'-3')
<i>GCK</i>	sense	CCCAGTCGTTGACTCTGGTAG
	antisense	CCACGGTCCATCTCCTTCTG
<i>GSK-3β</i>	sense	GCTGTGTGTTGGCTGAATTGT
	antisense	TGCTCCTGGTGAGTCCTTGT
<i>Glut4</i>	sense	GGCTCTGACGTAAGGATGGG
	antisense	GCCACGTTGCATTGTAGCTC
<i>G-6-pase</i>	sense	TGAGACC GGACCAGGAAGTC
	antisense	GCAAGGTAGATCCGGGACAG
<i>PEPCK</i>	sense	GGGTGGAAGGT CGAATGTGT
	antisense	AGCCCTTAAGTTGCCTTGGG
	sense	ACAGGTGTGAAGATATTGACGAGT
<i>LDLR</i>	G	
	antisense	CGGTTGGTGAAGAGCAGATAGC
<i>SR-B1</i>	sense	CCTCTGTCTCCGTCTCCTTC
	antisense	TCTATGCGGACATTCTTGAGCA
<i>CYP7A1</i>	sense	ACACTACTTCTGCGAAGGCAT
	antisense	CCCGGAGGTCTCATGACAGA
	sense	TGAGGGCTGCAAAGGTTCT
<i>FXR</i>	antisense	ACCTGTATA CATACATT CAGCAA
	C	
<i>PPARα</i>	sense	TGCCTTCCCTGTGAAGTGAC
	antisense	CACAGAGCGCTAACGCTGTGA
<i>β-actin</i>	sense	CACTGTCGAGTCGCGTCC
	antisense	TCATCCATGGCGAACTGGTG

Table S3 Linear regression equation and precision of SCFA standards

SCFA	Retention Time (min)	Regression equation	Linear range ($\mu\text{moL/mL}$)	R	Repeatability RSD (%)
Acetic	5.45	$y=9.8424x-520.62$	78.125~5000	0.9975	5.44
Propionic	6.55	$y=21.808x-1072$	78.125~5000	0.9992	4.72
Isobutyric	6.91	$y=29.927x-1217$	78.125~5000	0.9975	5.65
Butyric	7.73	$y=33.669x-1534.5$	78.125~5000	0.9975	5.51
Isovaleric	8.23	$y=46.841x-1962.9$	78.125~5000	0.9980	7.59
Valeric	9.13	$y=47.235x-2236.8$	78.125~5000	0.9980	7.62

Table S4 Effects of different treatments on body weight of mice.

Wee k	C	HF	KS	DK	KG	FKG
0	19.67±1.07 ^a	19.63±0.99 ^a	19.63±0.90 ^a	19.63±0.78 ^a	19.64±0.72 ^a	19.63±0.64 ^a
1	20.88±0.77 ^a	21.81±1.13 ^a	21.13±0.63 ^a	21.25±1.26 ^a	20.68±0.48 ^a	21.35±0.98 ^a
2	22.02±0.83 ^a	22.89±1.39 ^a	21.76±0.70 ^a	22.17±1.26 ^a	21.78±0.40 ^a	22.16±1.45 ^a
3	22.66±1.04 ^a	23.91±1.53 ^a	22.69±0.93 ^a	23.22±1.27 ^a	23.19±0.79 ^a	23.11±1.49 ^a
4	23.69±1.17 ^a	25.06±1.63 ^a	23.68±0.98 ^a	23.95±1.18 ^a	24.22±0.87 ^a	24.12±1.69 ^a
5	25.12±1.40 ^a	26.46±1.81 ^b	24.80±1.11 ^a	25.27±1.42 ^{ab}	25.30±1.11 ^{ab}	25.05±1.69 ^{ab}
6	25.53±1.45 ^a	27.68±1.80 ^b	25.70±1.12 ^a	25.82±1.23 ^{ab}	25.85±1.15 ^{ab}	25.54±1.90 ^{ab}
7	25.44±1.30 ^a	28.75±2.08 ^b	26.43±1.33 ^{ab}	26.97±1.34 ^{ab}	26.8±1.45 ^{ab}	26.22±2.06 ^{ab}
8	25.77±1.30 ^a	29.68±2.10 ^b	26.78±1.40 ^a	27.53±1.12 ^{ab}	27.22±1.82 ^{ab}	26.63±2.40 ^{ab}
9	26.17±1.32 ^a	30.66±1.87 ^b	27.23±1.40 ^a	28.26±1.24 ^a	28.16±1.85 ^a	27.25±2.36 ^a
10	26.26±1.36 ^a	31.53±2.24 ^b	27.72±1.36 ^a	28.79±1.36 ^a	28.84±1.81 ^a	27.66±2.29 ^a
11	26.00±1.21 ^a	32.00±2.31 ^c	28.29±1.36 ^{ab}	29.64±1.44 ^{ab}	29.50±2.11 ^{ab}	28.08±2.79 ^{ab}
12	26.25±1.15 ^a	32.84±1.63 ^c	28.96±1.00 ^b	29.86±1.89 ^b	29.86±1.39 ^b	28.30±1.73 ^{ab}

Note: Different lowercase letters in the same line indicated a statistically significant difference between groups ($P<0.05$).

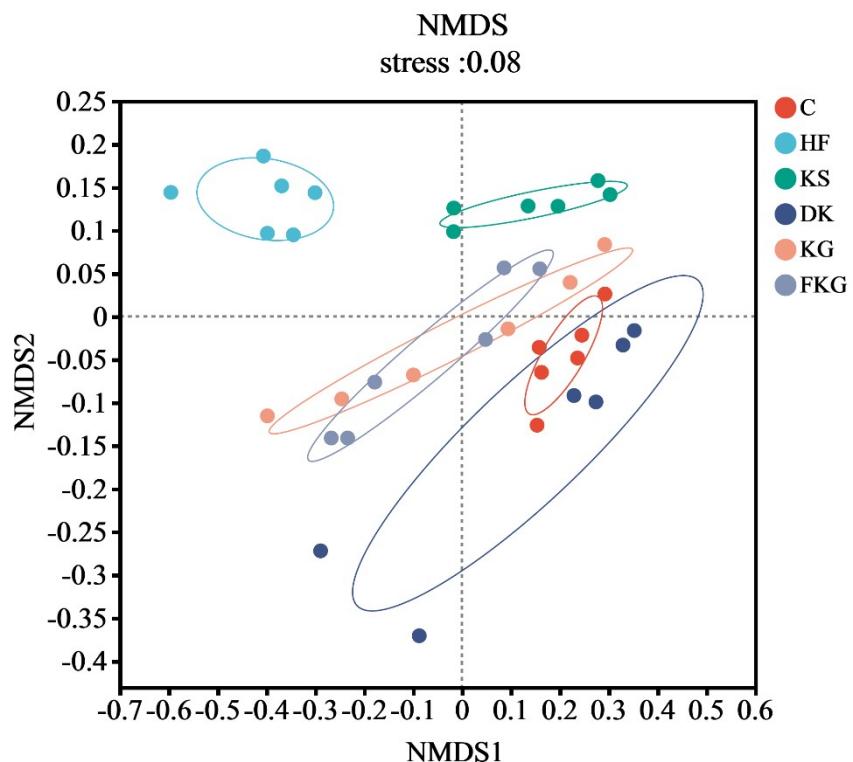


Fig.S1 NMDS analysis of β diversity index of mouse intestinal flora