

*Supporting Information*

In Vitro Fermentation of Different Indigestible Glucans with Varying Physico-Chemical Properties by Human Fecal Microbiota

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Method supplement

Changes of molecular weight of polysaccharides from indigestible glucans during fermentation.

Molecular weight of polysaccharide from indigestible glucans during fermentation was determined by high performance liquid chromatography (HPLC 1260, Agilent, USA) as described by Wang et al. [1], with modification. 100 uL of supernatant (4 °C, 15 min, 13000 rpm) was mixed with 400 μL of anhydrous ethanol and rested for 2 h to obtain sediment, the residual ethanol was removed by vacuum centrifugation, then re-dissolved with 1 mL of 0.02% NaN3 mobile phase and filtered (0.22 μm) for determination.

Standard curves were made using glucose and dextran (Dextran T-10, T-40, T-70, T-500, T-2000) as standards. Chromatographic conditions: Ultrahydrogel™ Linear (7.8 mm × 300 mm) column, 0.02% NaN₃ solution was used as mobile phase at a flow rate of 0.6 mL/min, column temperature of 35 °C, and data acquisition time was 30 min.

Result supplement

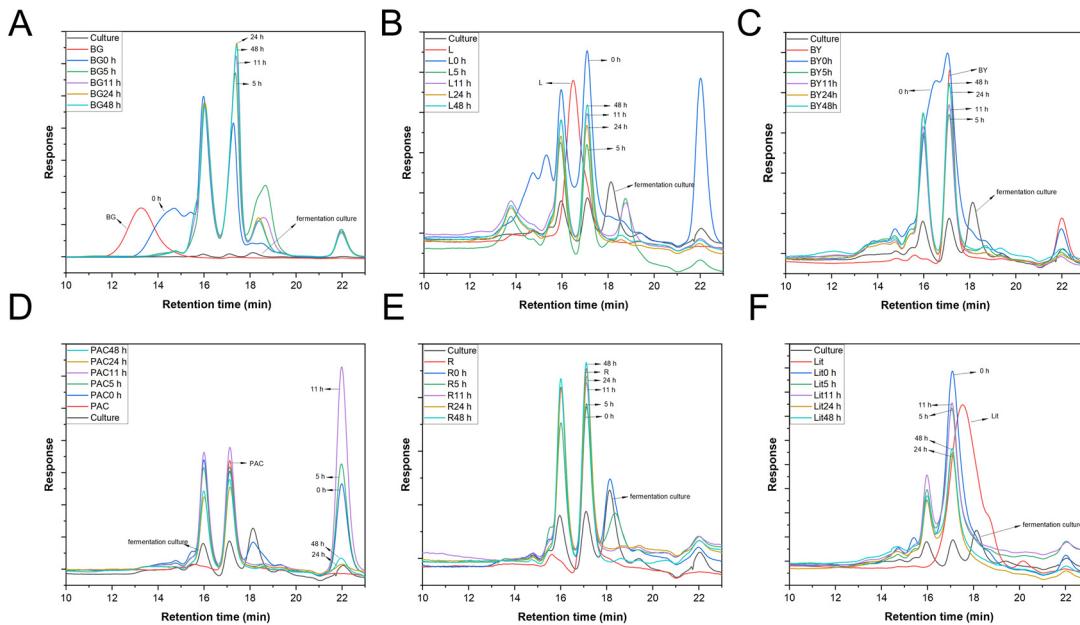


Figure S1. Changes of molecular weight of polysaccharides from indigestible glucans during fermentation. (A) Barley beta-glucan; (B) Laminarin; (C) Yeast beta-glucan; (D) Pachyman; (E) Resistant starch; (F) Litesse.

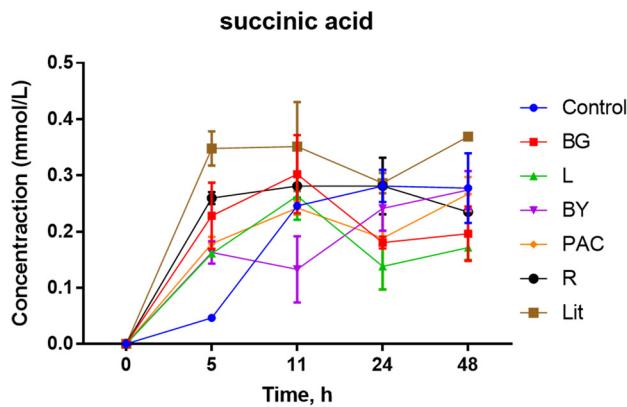


Figure S2. Changes in the content of succinic acid during fermentation.

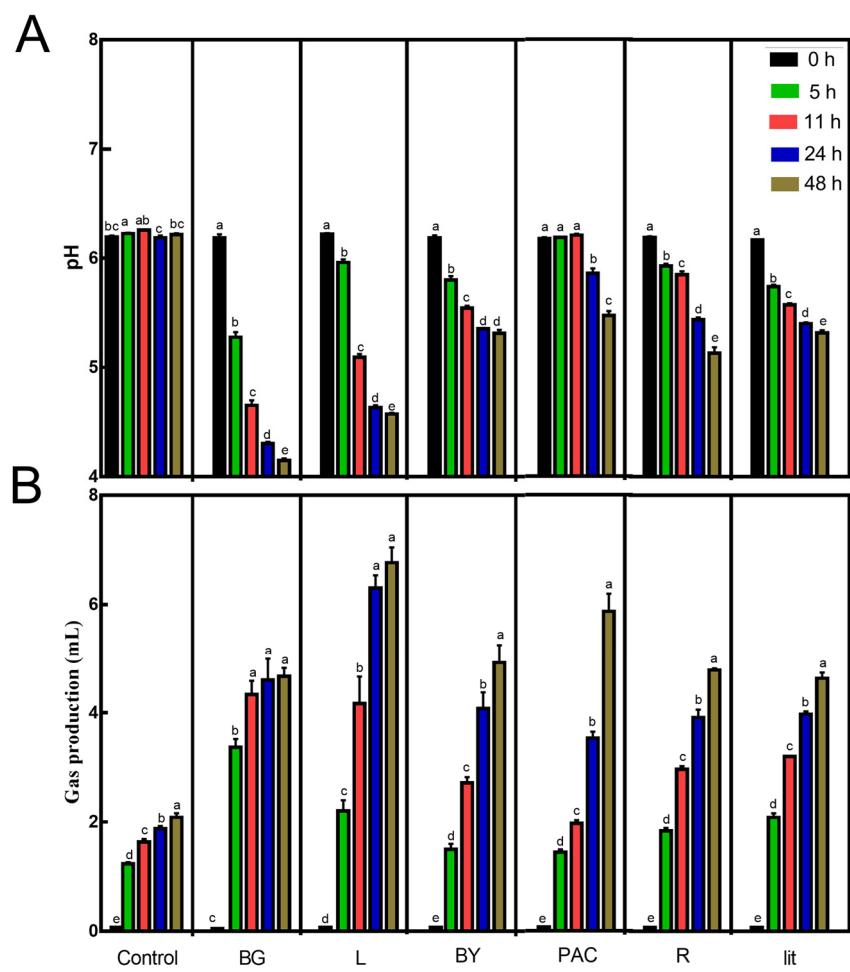


Figure S3. Bar chart for statistical analysis of pH and gas production. (A) pH change. (B) Gas production. BG, barley β -glucan; L, laminarin; BY, yeast β -glucan; PAC, pachymannan; R, resistant starch; Lit, litesse. Different letters indicate significant difference between groups, $p < 0.05$.

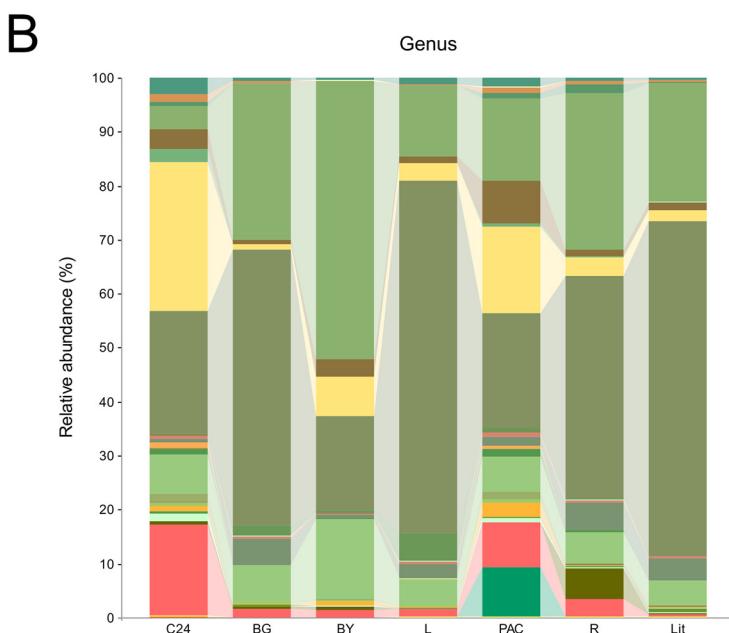
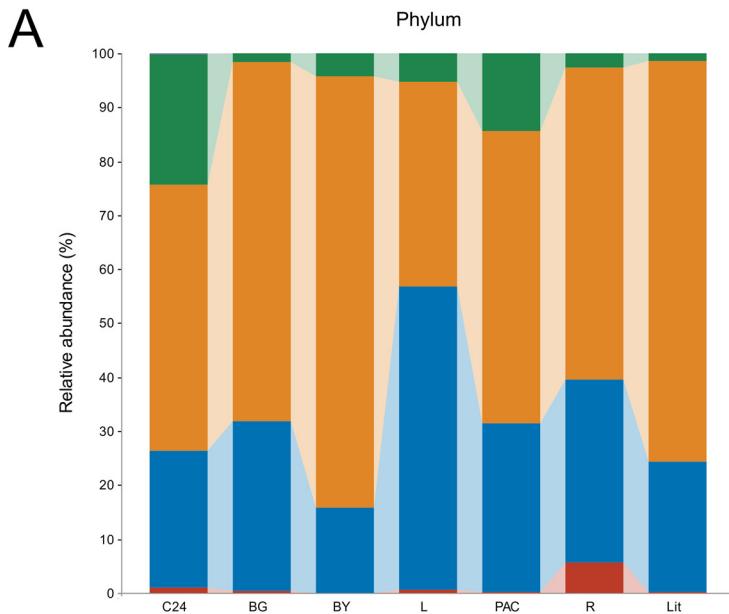


Figure S4. Bar chart of gut microbiota composition. (A) Phylum level;(B) Genus level. C24, control 24 h; BG, barley β -glucan; L, laminarin; BY, yeast β -glucan; PAC, pachyman; R, resistant starch; Lit, litesse.

Table S1. Changes in the content of lactic acid during fermentation.

	Control	BG	L	BY	PAC	R	Lit
0 h	0.46 \pm 0 ^d	1.26 \pm 0.07 ^c	0.77 \pm 0.04 ^c	1.26 \pm 0.13 ^b	1.15 \pm 0.03 ^{ab}	1.17 \pm 0.01 ^{cd}	1.22 \pm 0 ^b
5 h	1.16 \pm 0.03 ^c	1.23 \pm 0.05 ^c	1.23 \pm 0.11 ^c	1.37 \pm 0.1 ^b	0.92 \pm 0.08 ^b	1.1 \pm 0.01 ^d	1.22 \pm 0.03 ^b
11 h	1.68 \pm 0.12 ^{ab}	14.46 \pm 0.62 ^b	11.91 \pm 0.18 ^a	0.74 \pm 0.23 ^b	1.29 \pm 0.09 ^{ab}	1.77 \pm 0.04 ^{ab}	1.30 \pm 0.53 ^b
24 h	1.81 \pm 0.20 ^a	24.17 \pm 0.33 ^a	8.85 \pm 1.88 ^b	2.25 \pm 0.19 ^a	1.87 \pm 0.48 ^a	1.49 \pm 0.07 ^{bc}	1.56 \pm 0.06 ^b
48 h	1.37 \pm 0.17 ^{bc}	25.5 \pm 0.72 ^a	10.89 \pm 0.84 ^{ab}	2.5 \pm 0.47 ^a	1.46 \pm 0.34 ^{ab}	2.1 \pm 0.24 ^a	2.42 \pm 0.20 ^a

BG, barley β -glucan; L, laminarin; BY, yeast β -glucan; PAC, pachyman; R, resistant starch; Lit, litesse. Different letters indicate significant difference between groups, $p < 0.05$.

Table S2. Released free monosaccharides and oligosaccharides during in vitro fermentation (mg/mL).

	Control	BG	L	BY	PAC	R	Lit
Monosaccharid							
e							
0 h	ND	¹ 0.16 ± 0.03 ^c	0.4 ± 0.05 ^c	0.25 ± 0.05 ^b	0.11 ± 0.01 ^a	0.29 ± 0.02 ^a	0.36 ± 0.04 ^a
5 h	ND	3.10 ± 0.08 ^a	2.19 ± 0.16 ^a	0.5 ± 0.08 ^a	0.01 ± 0 ^c	0.06 ± 0.01 ^b	0.56 ± 0.05 ^b
11 h	ND	1.20 ± 0.05 ^b	1.24 ± 0.06 ^b	0.24 ± 0.01 ^b	0.02 ± 0.01 ^c	0.11 ± 0.03 ^b	0.32 ± 0.14 ^a
24 h	ND	0.08 ± 0.05 ^c	0.05 ± 0.01 ^d	0.04 ± 0.01 ^c	0.07 ± 0.01 ^b	0.06 ± 0.02 ^b	0.14 ± 0.02 ^b
48 h	0.02 ± 0.01	0.10 ± 0.03 ^c	0.01 ± 0 ^d	0.01 ± 0.01 ^c	0.05 ± 0.01 ^b	0.06 ± 0.01 ^b	0.09 ± 0.01 ^b
Disaccharide							
0 h	ND	0.02 ± 0 ^b	ND	0.04 ± 0.02 ^a	ND	ND	*0±0
5 h	ND	1.55 ± 0.05 ^a	0.02 ± 0.01 ^a	ND	ND	ND	ND
11 h	ND	1.33 ± 0.05 ^a	*0 ± 0 ^b	ND	ND	ND	ND
24 h	ND	1.17 ± 0.41 ^a	ND	ND	ND	ND	ND
48 h	ND	1.45 ± 0.07 ^a	ND	ND	ND	ND	ND
Trisaccharide							
0 h	ND	0.02 ± 0 ^a	ND	0.02 ± 0 ^a	ND	ND	*0 ± 0 ^b
5 h	ND	0.01 ± 0 ^b	0.18 ± 0.06 ^a	*0 ± 0 ^b	ND	ND	0.05 ± 0 ^a
11 h	ND	*0 ± 0 ^b	0.16 ± 0.01 ^a	ND	ND	ND	0.02 ± 0.02 ^{ab}
24 h	ND	*0 ± 0 ^b	*0 ± 0 ^b	ND	ND	ND	0.01 ± 0.01 ^b
48 h	ND	*0 ± 0 ^b	*0 ± 0 ^b	ND	ND	ND	0.01 ± 0 ^b
Tetrasaccharide							
0 h	ND	ND	ND	ND	ND	ND	ND
5 h	ND	ND	ND	ND	ND	ND	ND
11 h	ND	ND	ND	ND	ND	ND	ND
24 h	ND	ND	ND	ND	ND	ND	0.05±0 ^a
48 h	ND	ND	ND	ND	ND	ND	0.04±0 ^a

¹Results are expressed as mean ± standard deviation ($n = 3$), and significant comparisons were made between groups of the same polysaccharide in the same column. Different letters indicate significant difference between groups, $p < 0.05$. ND indicates not detected by the equipment. * Indicates lower value, not shown (<0.01 mg/mL). BG, barley β -glucan; L, laminarin; BY, yeast β -glucan; PAC, pachymann; R, resistant starch; Lit, litesse.

Table S3. Primer sequences used in real-time quantification PCR.

Target bacterium	Primer name	Sequence (5'-3')	Amplicon size (bp)	Tm (°C)	Reference
Total bacteria	338F	ACTCCTACGGGAGGCAGCAGT	192	60	Castillo et al [2]
	518R	ATTACCGCGGCTGCTGGC			
<i>Lactobacillus</i>	Lacto-F	GGAATCTTCCACAATGGACG	217	56	Bakar et al [3]
	Lacto-R	CGCTTTACGCCAATAATCCGG			
<i>Bifidobacterium</i>	Bifid-F	TCGCGTCYGGTGTGAAAG	243	58	Rinttilä et al [4]
	Bifid-R	CCACATCCAGCRTCCAC			
<i>Faecalibacterium prausnitzii</i>	FPR-2F	GGAGGAAGAACGGTCTCGG	245	60	Carlett et al [5]
	Fprau645R	AATTCCGCCTACCTCTGCACT			
<i>Bacteroides</i>	g-Bfra-F	ATAGCCTTCGAAAGRAAGAT	493	60	Matsuki et al [6]
	g-Bfra-R	CCAGTATCAACTGCAATTAA			
<i>Prevotella</i>	g-Prevo-F	CACRGTAACGATGGATGCC	515	56	Matsuki et al [7]
	g-Prevo-R	GGTCGGGTTGCAGACC			

Table S4. Permutational MANOVA analysis of β -diversity among groups.

Group 1	Group 2	Sample size	Permutations	pseudo-F	p-value	q-value
BG24	BY24	8	0.999	49.48552	0.026	0.040385
BG24	C0	8	0.999	68.27435	0.031	0.040385
BG24	C24	8	0.999	111.5937	0.028	0.040385
BG24	L24	8	0.999	9.321454	0.032	0.040385
BG24	Lit24	8	0.999	7.989927	0.034	0.040385
BG24	PAC24	8	0.999	64.91326	0.028	0.040385
BG24	R24	8	0.999	10.9191	0.036	0.0405
BY24	C0	8	0.999	24.86024	0.023	0.040385
BY24	C24	8	0.999	61.42028	0.035	0.040385
BY24	L24	8	0.999	51.91417	0.035	0.040385
BY24	Lit24	8	0.999	40.44521	0.028	0.040385
BY24	PAC24	8	0.999	39.01315	0.026	0.040385
BY24	R24	8	0.999	34.30895	0.034	0.040385
C0	C24	8	0.999	63.9431	0.017	0.040385
C0	L24	8	0.999	65.44541	0.03	0.040385
C0	Lit24	8	0.999	47.23702	0.028	0.040385
C0	PAC24	8	0.999	47.20966	0.017	0.040385
C0	R24	8	0.999	40.21948	0.03	0.040385
C24	L24	8	0.999	78.13104	0.034	0.040385
C24	Lit24	8	0.999	59.77891	0.032	0.040385
C24	PAC24	8	0.999	25.2231	0.029	0.040385
C24	R24	8	0.999	65.11511	0.029	0.040385
L24	Lit24	8	0.999	3.021266	0.053	0.056786
L24	PAC24	8	0.999	60.33565	0.026	0.040385
L24	R24	8	0.999	17.07078	0.034	0.040385
Lit24	PAC24	8	0.999	50.23807	0.027	0.040385
Lit24	R24	8	0.999	9.727374	0.028	0.040385
PAC24	R24	8	0.999	41.38957	0.028	0.040385

C0, control 0 h; C24, control 24 h; BG24, barley β -glucan 24 h; L24, laminarin 24 h; BY24, yeast β -glucan 24 h; PAC24, pachymann 24 h; R24, resistant starch 24 h; Lit24, litesse 24 h.

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