

**Modulation of gut microbiota and hypoglycemic/hypolipidemic activity of flavonoids from the fruits of *Lycium barbarum* on high-fat diet/streptozotocin-induced type 2 diabetic mice**

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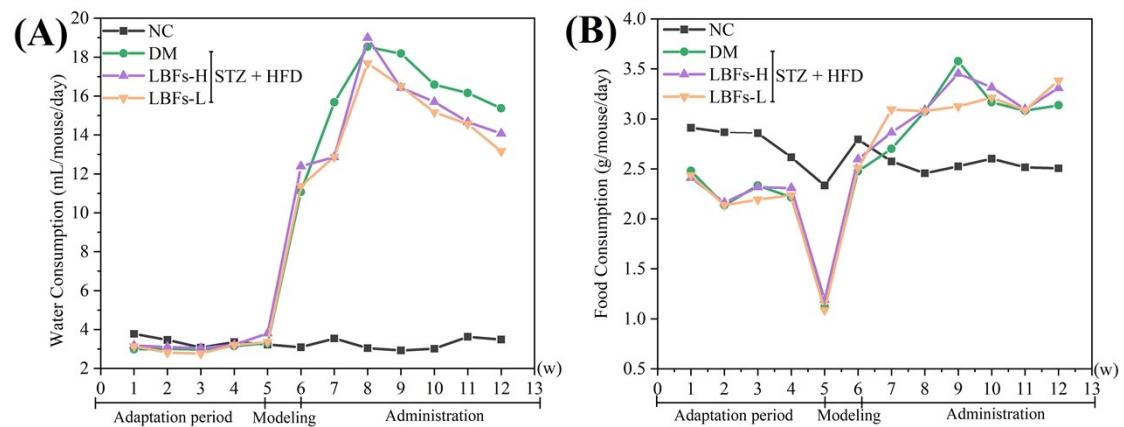
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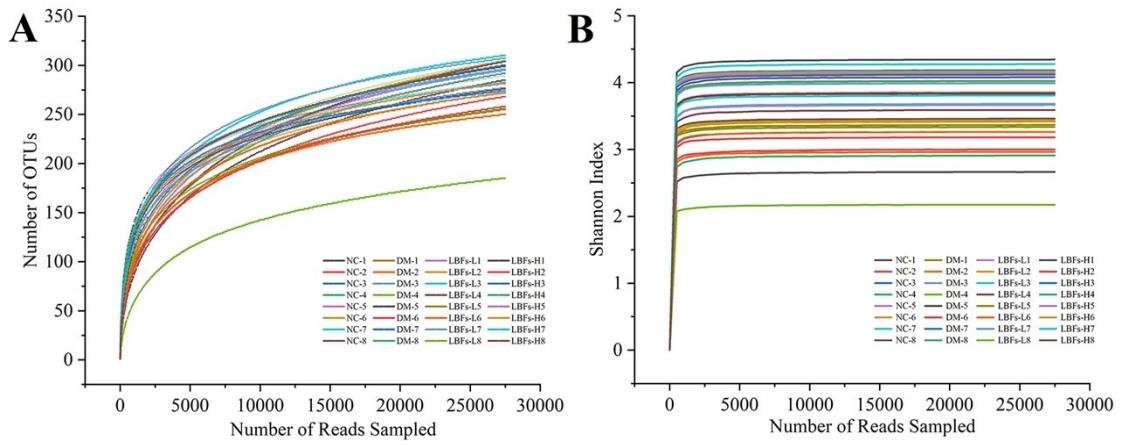
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<b>Target gene</b>	<b>Primer Sequence (5' - 3')</b>	<b>Annealing T (°C)</b>
GAPDH	FW: GGACTTACAGAGGTCCGCTT RV: CTATAGGGCCTGGGTCACTG	59
TLR-4	FW: GCTCTCAGCCATCCACAAAG RV: GAGTCGGGAAGAGGAAGAGG	60
TNF- $\alpha$	FW: CTCATGCACCACCATCAAGG RV: ACCTGACCACTCTCCCTTG	60
IL-6	FW: TAGTCCTCCTACCCCAATT RV: TTGGTCCTTAGCCACTCCTTC	60
IL-10	FW: AGTACAGCCGGGAAGACAAT RV: TCTAGGAGCATGTGGCTCTG	59
IFN- $\gamma$	FW: AGCTCTCCTCATGGCTGTT RV: GGTCAACCAACCACAAGCAT FW:	59
GK	AAAACACGTATGGAACAGGGTG RV: TTAGCCAGCGGATTACAGCA	60
G6Pase	FW: TCTACCTTGCAGGCTCACTTC RV: GAAAGTTTCAGCCACAGCAAT	59
PEPCK	FW: GCAGAGCATAAGGGCAAGGT RV: TTGCCGAAGTTGTAGCCAAA	59
PFK	FW: ACCGTGGACCTGGAGAAA RV: AGCCCTGACAGCAGCATT	60
PPAR $\alpha$	FW: TCATCAAGAAGACCGAGTCC RV: CCTCTTCATCCCCAAGCGTA	60
PPAR $\gamma$	FW: GCTGAACGTGAAGCCCATCG RV: GGCAGAACAGCTGAGAGGACT	60
FAS	FW: GGCACCTATGGCGAGGACTT RV: GCCCTCCCGTACACTCACTC	60
ACC	FW: AAAGGCTATGTGAAGGATG RV: TCTGAAGAGGTTAGGGAAG	59
SREBP-1c	FW: GGCACTAAGTGCCCTAACCT RV: GCCACATAGATCTCTGCCAGTGT	60

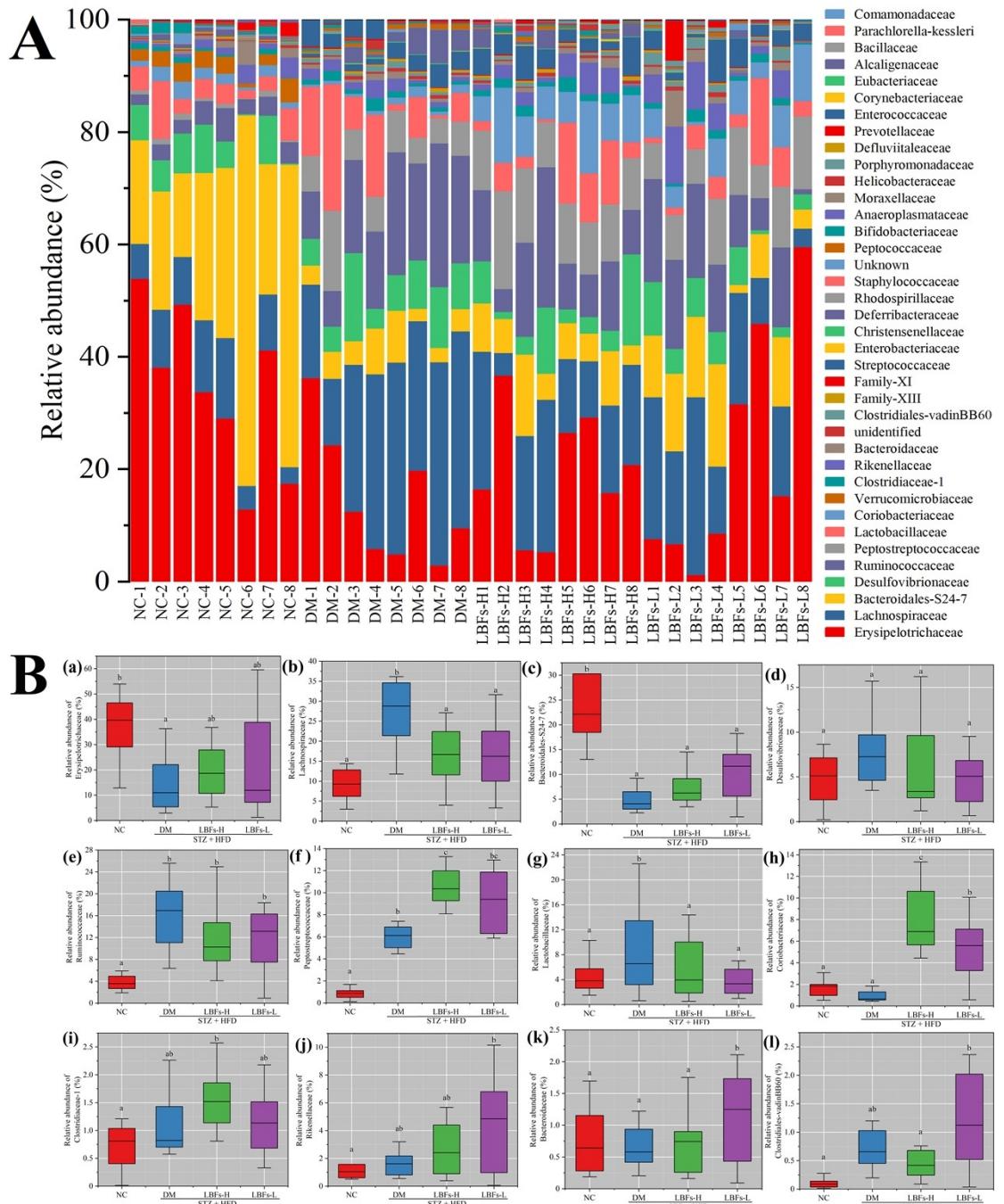
**Table S1.** Primer sequences used in RT-qPCR assays in colonic tissue.



**Fig. S1.** Water consumption (A) and food consumption (B) of the four groups.



**Fig. S2.** Rarefaction curves (A) and Shannon-Wiener curves (B) of the four groups.



**Fig. S3.** Bacterial taxonomic profiling at the family level of gut microbiota. (A) bacterial taxonomic profiling at the family level of gut microbiota; (B) comparative analysis of 12 most relative abundance of gut microbiota at the family level. a-d represent significant differences among different groups by one-way ANOVA procedure followed by Tukey's HSD test ( $p < 0.05$ ).