

**Modulating effects of polysaccharides from the fruits of *Lycium barbarum* on
immune response and gut microbiota in cyclophosphamide-treated mice**

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Table S1. Target Genes and Primers Sequence

Gene	Sequence (5'-3')
TNF- α	F: CTCATGCACCACCATCAAGG R: ACCTGACCACTCTCCCTTG
IL-1 β	F: AGCTTCAAATCTCGCAGCAG R: TCTCCACAGCCACAATGAGT
TLR4	F: GCTCTCAGCCATCCACAAAG R: GAGTCGGGAAGAGGAAGAGG
MyD88	F: TTTCCCAGTATCCTGCGGTT R: GCGBAAGAACACAGACAGAC
p65	F: GACACAGATGTGGTGGAACG R: GTGAGGTGGATCCTGGTGA
Occludin1	F: AGCACTTAACCTGCCTGGAT R: AGCCTGTGGAAGCAAGAGAT
Claudin5	F: TTCTTCTATGCGCAGTTGGC R: TTGGTGCCACTTCACCGAT
MUC2	F: GCTCTCAAATGGTGCAGAGG R: AGTGGGAAGGGATGAATGGG
GAPDH	F: GGACTTACAGAGGTCCGCTT R: CTATAAGGCCTGGTCAGTG

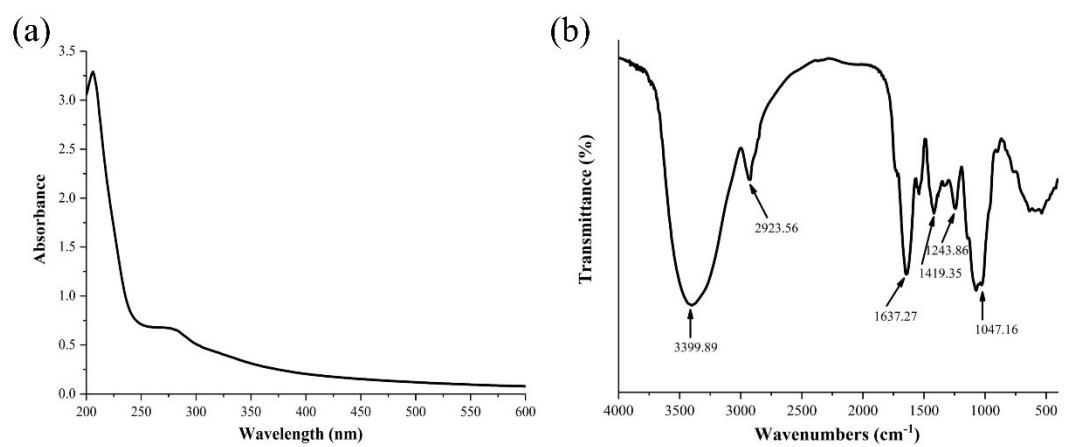


Fig. S1. UV (a) and FT-IR (b) spectra of LBPS

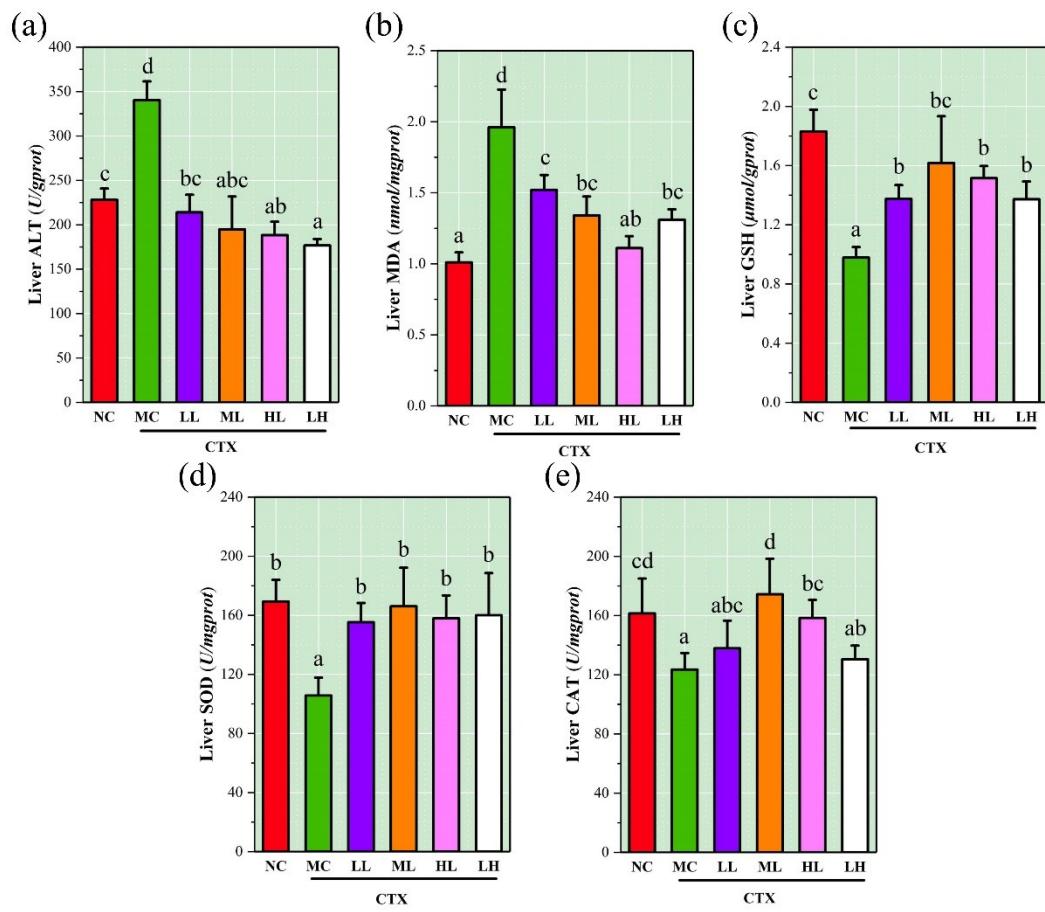


Fig. S2. Effects of LBPS administration on liver parameters including ALT (a), MDA (b), GSH (c), SOD (d), CAT (e). The different letters represent significant differences between different groups ($p < 0.05$).

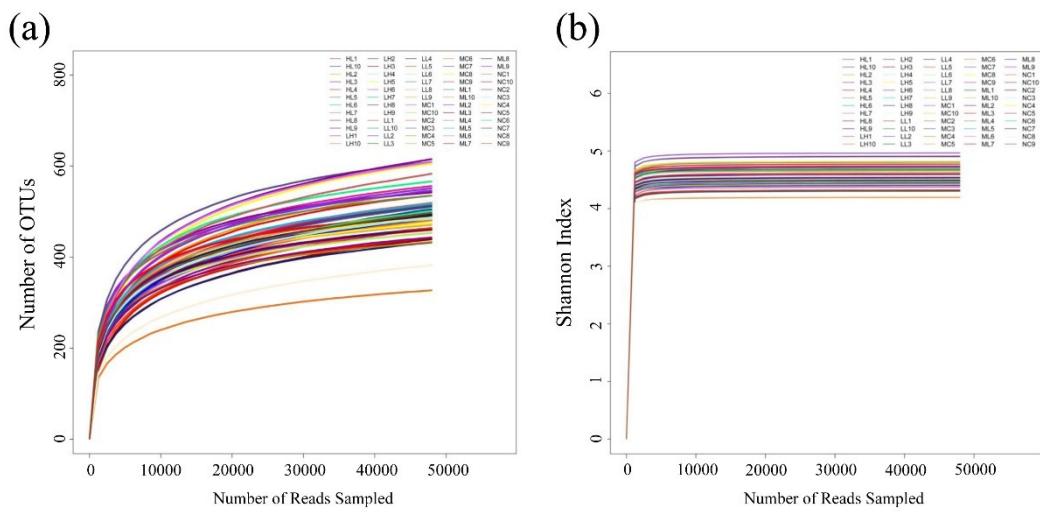


Fig. S3. Alpha diversity indexes of microbiota in different groups. Observed species

(a), shannon index (b) of fecal samples from NC, MC, LL, ML, HL and LH groups.

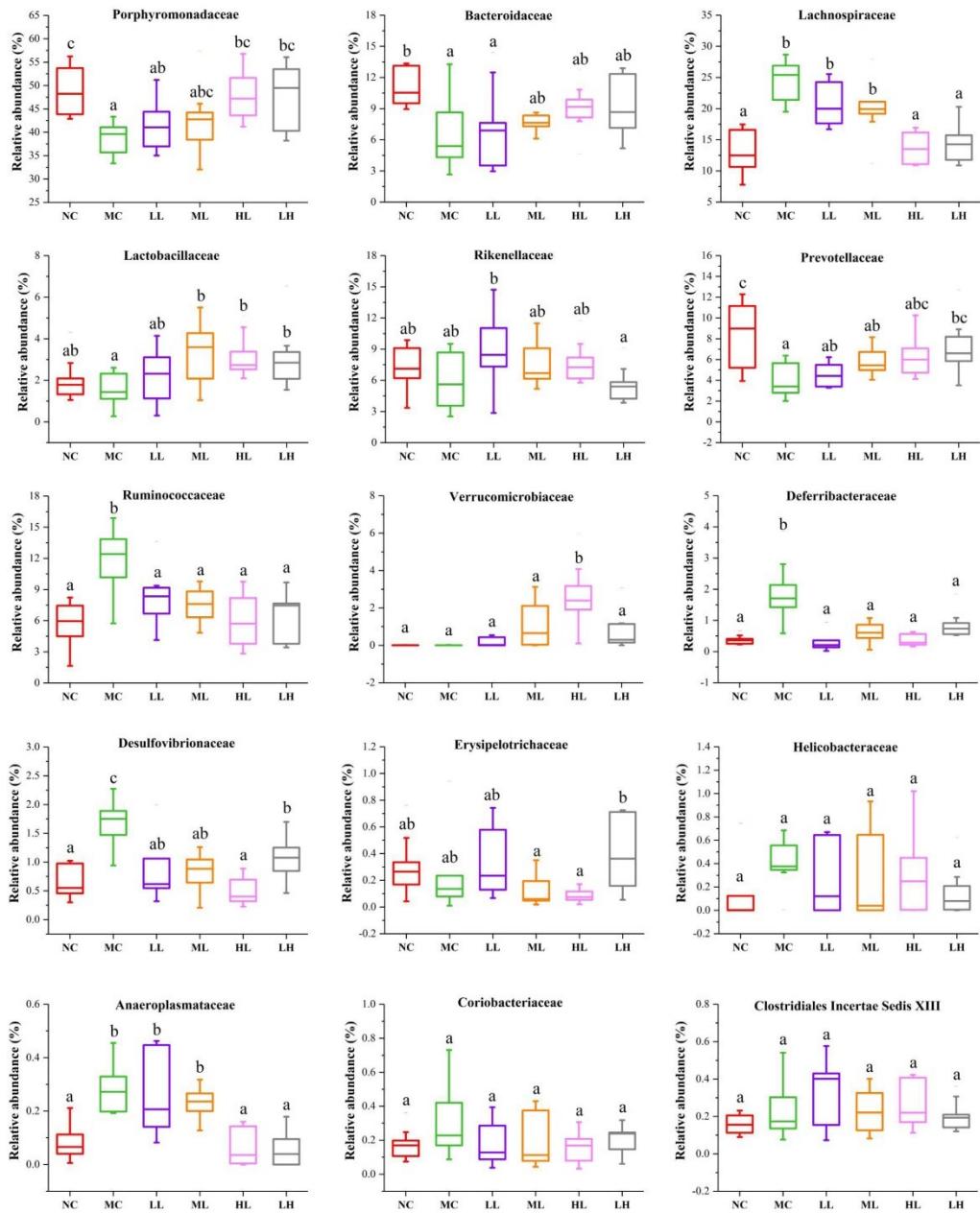


Fig. S4. Comparative analyses of 15 gut microbiota with most relative abundances at the family level. One-way ANOVA procedure followed by Tukey test was used to evaluate the statistical significance. The different letters represent significant differences between different groups ($p < 0.05$).