Bacteroides vulgatus attenuates experimental mice colitis through modulating gut microbiota and immune responses

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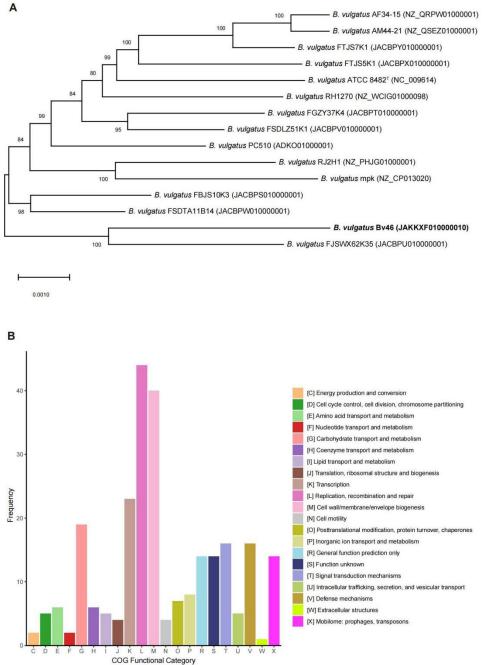


FIGURE S1 Genome analysis of B. vulgatus strains. (A) The phylogenetic relationship of 15 B. vulgatus strains based on core genes. (B) The number of B. vulgatus Bv46 strain-specific genes assigned in COGs (Clusters of Orthologous Groups) categories.

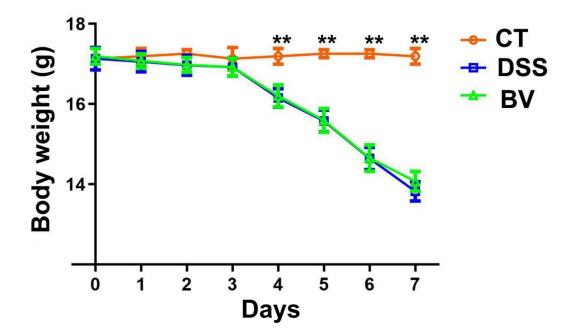


FIGURE S2 Changes in body weight of DSS-induced mice treated with *B. vulgatus* Bv46 (in BV group), or phosphate buffer saline (in DSS group), and control (CT, no DSS induction) for 7 days. Values represent the mean \pm standard deviation. **, *P* < 0.01 indicated the significant differences of DSS group versus control group.

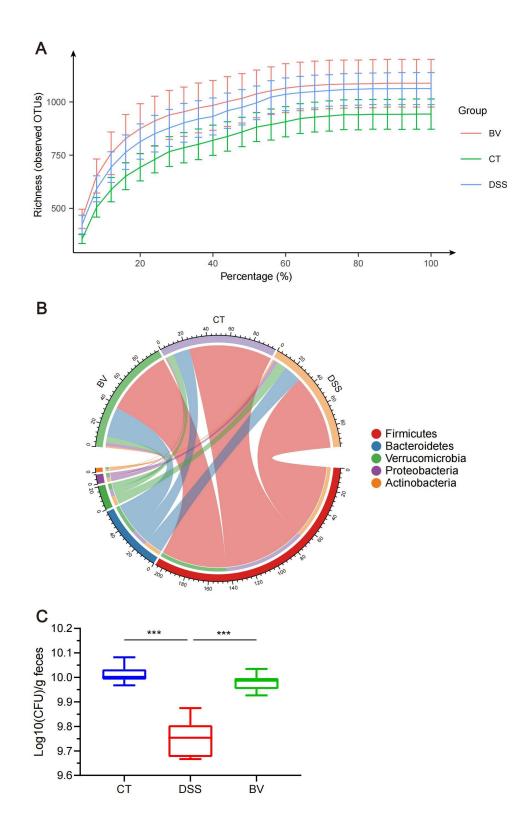


FIGURE S3 (A) Rarefaction curve analysis of observed OUTs from the three groups based on random sampling. (B) Overview of the dominant bacterial phyla in the three groups. (C) Species specific qPCR-tracked *B. vulgatus* in feces of mice. Statistical comparison was implemented by a one-way ANOVA followed by Tukey's multiple-comparison test. Values are expressed as quartiles in each group, ***, P < 0.001.

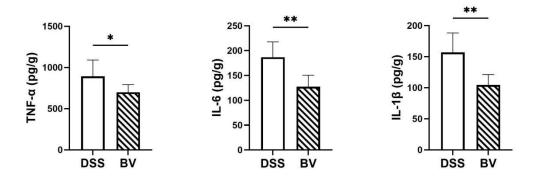


FIGURE S4 The expression of colonic inflammatory cytokines (TNF- α , IL-6 and IL-1 β) in DSS and BV groups. Values represent the mean \pm standard deviation. Significance levels are shown as *, *P* < 0.05; **, *P* < 0.01.