

Supplementary materials

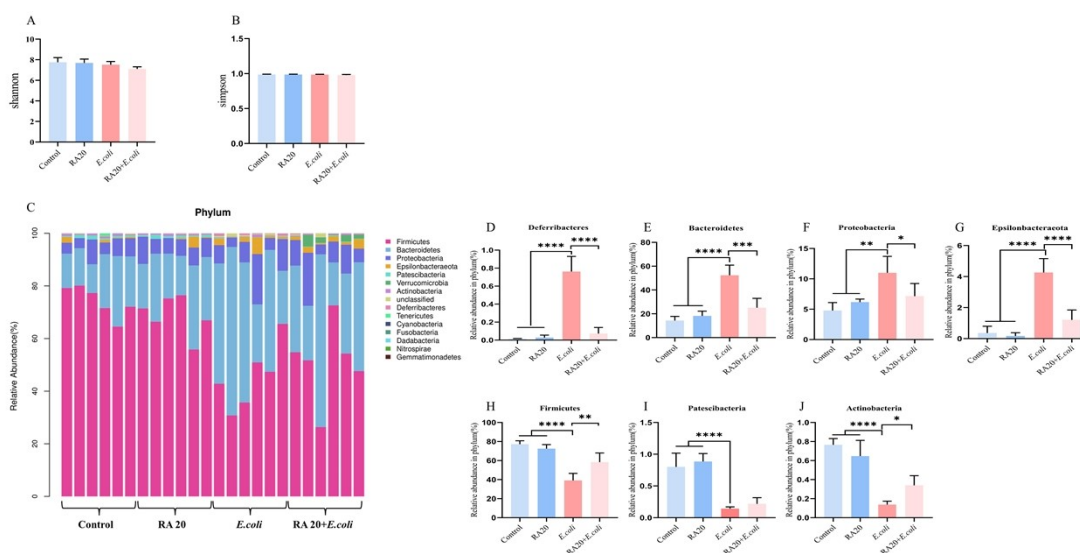
Table 1. Specific primers for real-time PCR

target gene	Forward (5'-3')	Reverse (5'-3')
TNF-α	F: CGCTGAGGTCAATCTGC	R: GGCTGGGTAGAGAATGGA
IL-6	F: ACAGAAGGAGTGGCTAAGGA	R: AGGCATAACGCACTAGGTTT
ZO-1	F: GCTTTAGCGAACAGAAGGAGC	R: TTCATTTTCCGAGACTTCACCA
Occludin	F: TGAAAGTCCACCTCCTTACAGA	R: CCGGATAAAAAAGAGTACGCTGG
Claudin-1	F: TGCCCCAGTGGAAAGATTTACT	R: CTTTGCGAAACGCAGGACAT
β-actin	F: CCTCACTGTCCACCTCC	R: GGGTGTA AAAACGCAGCTC

Table 2 The minimum inhibitory concentration of RA against *E. coli*

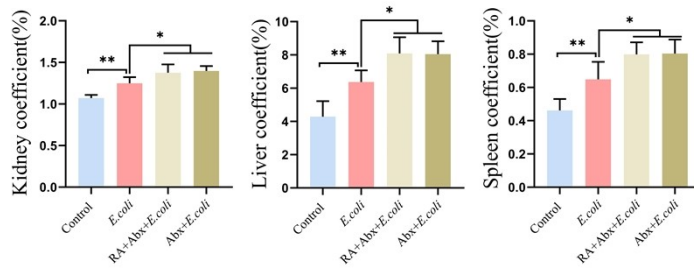
drug	4.00	2.00	1.00	0.500	0.250	0.125	0.0625	0.0313	0.0157	0.00785	Positive control	negative control
concentration (mg/mL)												
1	-	-	+	+	+	+	+	+	+	+	+	-
2	-	-	+	+	+	+	+	+	+	+	+	-
3	-	-	+	+	+	+	+	+	+	+	+	-

+ represents positive, indicating the presence of bacterial precipitation, - represents negative, and no bacterial precipitation appears.

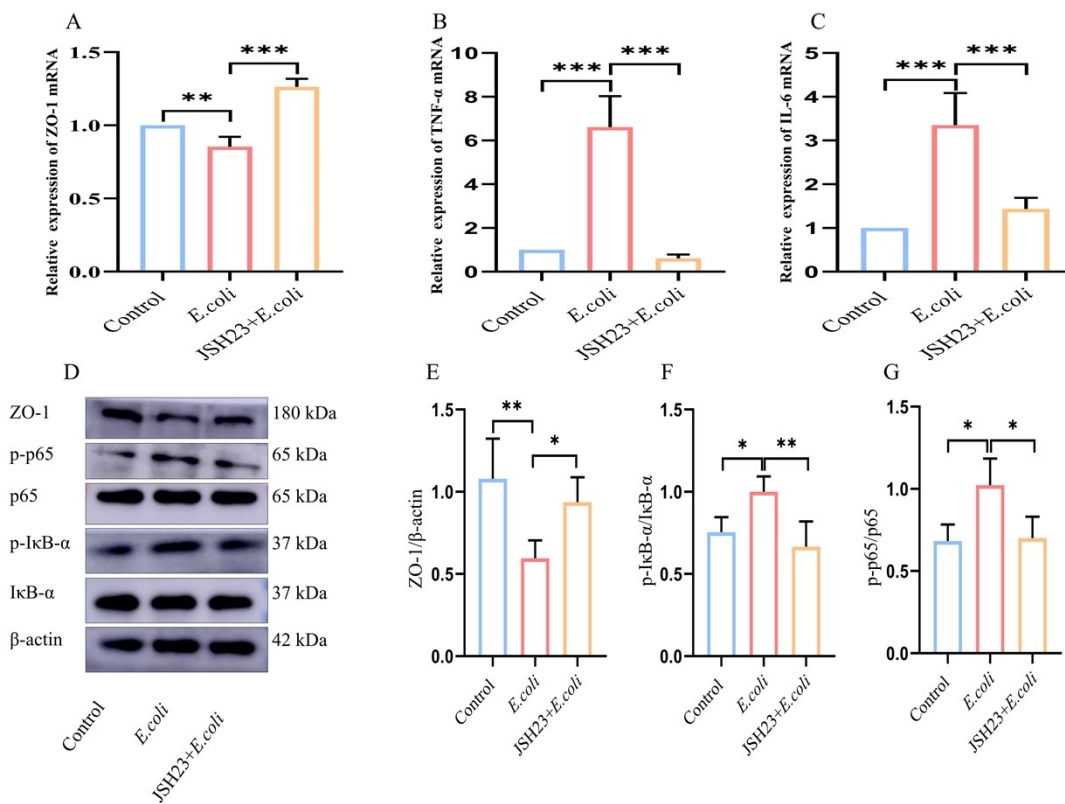


Supplementary Figure 1 RA maintained dysbiosis of the bacterial community caused by *E. coli*. (A) Alpha diversity analysis- shannon (n = 6); (B) Alpha diversity analysis-simpson (n = 6); (C) Relative abundance of the gut microbiota at the Phylum level (n = 6); (D) the relative abundance of p__*Deferribacteres* (n = 3); (E) the relative abundance of p__*Bacteroidetes* (n = 3); (F) the relative abundance of p__*Proteobacterin* (n = 3); (G) the relative abundance of p__*Epsilonbacteraeota* (n = 3); (H) the relative abundance of p__*Firmicutes* (n = 3); (I) the relative

abundance of p__*Patiscibacteria* (n = 3); (J)the relative abundance of p__*Actinobacteria* (n = 3).



Supplementary Figure 2 Measurement of organ coefficient. (A) Mice liver coefficient (n = 6); (B) Mice kidney coefficient (n = 6); (C) Mice spleen coefficient (n = 6).



Supplementary Figure 3 RA reduced inflammation caused by *E. coli* by inhibiting the NF-κB signaling pathway to restore ZO-1 expression. (A) The mRNA expression of ZO-1 (n = 6) ;(B) The mRNA expression of TNF-α (n = 6); (C) The mRNA expression of IL-6 (n = 6); (D-G) Expression of protein of ZO-1, p65, p-p65, IκB-α and p-IκB-α (n = 3).