Supporting Information

Gene	Primer Sequence (5'-3')	GenBank NO.
IL-6	F: CCCACCAAGAACGATAGTCA	NM_001314054.1
	R: CTCCGACTTGTGAAGTGGTA	
TNF-α	F: CAGGCGGTGCCTATGTCTC	NM_001278601.1
	R: CGATCACCCCGAAGTTCAGTAG	
IL-10	F: CAAGGAGCATTTGAATTCCC	NM_010548.2
	R: GGCCTTGTAGACACCTTGGTC	
Shank3	F: GATCTGCCATCCCTACAAC	NM_021423.4
	R: AGCTAAGGGTGAGCTAGGAT	
Pten	F: TGGATTCGACTTAGACTTGACCT	NM_008960.2
	R: GCGGTGTCATAATGTCTCTCAG	
Claudin-1	F: CATCAATGCCAGGTATGAATT	NM_016674.4
	R: TGTTGGGTAAGAGGTTGTTT	
GAPDH	F: AGGTCGGTGTGAACGGATTTG	NM_001289726.2
	R: GGGGTCGTTGATGGCAACA	

Table S1. The primer sequence of target gene and housekeeping gene.

Figure S1



Figure S1 Synbiotic intervention improved behavioral impairment of VPA-induced ASD mice.

(A) Number of marbles buried in marble burying test. (B) In the three-chamber social test, the time spent by the test mice interacting with the empty cup and Mouse 1 was measured. (C) In the three-chamber social test, the time spent by the test mice interacting with Mouse 1 and Mouse 2 was measured. (D) Representative images of Iba-1 immunofluorescence staining in the brain, the Iba-1 were stained red, and nuclei were counter-stained blue. (E) Representative images of BDNF immunofluorescence staining in the cortex, the BDNF were stained red, and nuclei were counter-stained blue. (F) The fluorescence intensity of Iba-1 in the cortex. (G) The fluorescence intensity of BDNF in the cortex. Data are presented as mean \pm SEM and statistical analyses were determined by one-way ANOVA with Tukey's test. *p < 0.05, **p < 0.01, ns p > 0.05.

Figure S2



Figure S2 Effects of synbiotic treatment on colon morphology and barrier function in VPA-induced ASD mice.

(A) Representative images of H&E staining of the colon for each group (n=4). (B) Representative images of Alcian blue staining of the colon for each group (n=4). (C) Representative images of Claudin-1 immunofluorescence staining in the colon, the Claudin-1 were stained red, and nuclei were counter-stained blue (n=3-4). (D) Crypt depth based on H&E-stained sections. (E) Percentage of Alcian blue-stained area in the colon (%). (F) The fluorescence intensity of Claudin-1 in colon. (G) The relative mRNA expression levels of *Claudin-1* in the colon of each group. Data are presented as mean \pm SEM and statistical analyses were determined by one-way ANOVA with Tukey's test. *p < 0.05, **p < 0.01.

Figure S3





(A) A heatmap displaying the relative abundance of gut microbiota at the phylum level across different experimental groups. (B) A heatmap displaying the relative abundance of gut microbiota at the genus level across different experimental groups. (C) Relative abundance of Bacillota. (D) Relative abundance of Bacteroidota. (E) Relative abundance of *Helicobacter typhlonius*. (F) Relative abundance of *Akkermansia muciniphila*. (G) Relative abundance of *Lawsonibacter asaccharolyticus*. (H) Relative abundance of *Barnesiella intestinihominis*. (I) Relative abundance of *Dorea formicigenerans*. (J) Relative abundance of *Allobaculum stercoricanis*. Data are presented as mean ± SEM and statistical analyses were determined by one-way ANOVA with Tukey's test. *p < 0.05, **p < 0.01.