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1 **Fructooligosaccharides improves growth performance and intestinal**
2 **epithelium function in weaned pigs exposure to Enterotoxigenic**
3 ***Escherichia coli***

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23

24 Abstract

25 This study was conducted to explore the protective potential of
26 Fructooligosaccharides (FOS) against Enterotoxigenic *Escherichia coli*
27 (ETEC)-induced inflammation and intestinal injury in weaned pigs. Twenty-four
28 weaned pigs were randomly assigned into three groups: (1) non-challenge (CON, fed
29 with basal diet), (2) ETEC-challenge (ECON, fed with basal diet), and (3) ETEC
30 challenge + FOS treatment (EFOS, fed with basal diet plus 2.5 g/kg FOS). On day 19,
31 CON group was orally infused with sterilized culture while pigs in ECON group and
32 EFOS group were orally infused with ETEC (2.5×10^{11} colony-forming units). After
33 3 days, pigs were slaughtered for sample collection. We showed that
34 ETEC-challenged significantly reduced average daily gain (ADG), however, FOS
35 improved the ADG ($P < 0.05$), apparent digestibility of crude protein (CP), gross
36 energy (GE), ash and reduced the diarrhea incidence ($P < 0.05$). FOS reduced plasma
37 concentrations of IL-1 β and TNF- α , down-regulated ($P < 0.05$) the mRNA expression
38 of IL-6 and TNF- α in jejunum and ileum as well as IL-1 β and TNF- α in duodenum.
39 Elevated the concentrations of plasma immunoglobulin A (IgA), immunoglobulin M
40 (IgM) and secreted IgA (SIgA) in the jejunum ($P < 0.05$). Interestingly, FOS elevated
41 villus height in duodenum, and elevated the ratio of villus height to crypt depth in the

42 duodenum and ileum in the EFOS group pigs ($P<0.05$). Moreover, FOS increased
43 lactase activity in the duodenum and ileum ($P<0.05$). The activities of sucrase and
44 alkaline phosphatase (AKP) were higher in EFOS group than in the ECON group
45 ($P<0.05$). Importantly, FOS up-regulated the expressions of critical genes in intestinal
46 epithelium function such as zonula occludens-1 (ZO-1), L-type amino acid
47 transporter-1 (LAT1), and cationic amino acid transporter-1 (CAT1) in the duodenum
48 and the expressions of ZO-1 and glucose transporter-2 (GLUT2) in the jejunum
49 ($P<0.05$). FOS also up-regulated the expressions of occludin, fatty acid transporter-4
50 (FATP4), sodium glucose transport protein 1 (SGLT1), and GLUT2 in the ileum
51 ($P<0.05$). FOS significantly increased the concentrations of acetic acid, propionic acid
52 and butyric acid in the cecal digesta. Additionally, FOS reduced the populations of
53 *Escherichia coli*, but elevated the populations of *Bacillus* and *Bifidobacterium* in the
54 caecal digesta ($P<0.05$). These results suggested that FOS could improve the growth
55 performance and intestinal health in weaned pigs upon ETEC challenge, which was
56 associated with suppressed inflammatory responses and improved intestinal
57 epithelium functions and microbiota.

58 **Keywords:** *Escherichia coli*, FOS; Intestine, Inflammation, Microbiota, Weaned pigs

59

60 **Introduction**

61 Small intestine is the main site of nutrient digestion and absorption, and can serve as
62 an important defense barrier against exogenous and endogenous harmful substances
63 or pathogens [1]. Previous studies indicated that a wide variety of enteric pathogens
64 such as pathogenic bacteria and viruses can induce acute diarrhea in mammalian
65 animals, especially at their weaning period, resulting in mortality, dehydration, weight
66 loss, and growth retardation [2-4]. In the last decades, antibiotics have been widely
67 used for the preventing of diarrhea; however, long-term or overdose utilization of
68 antibiotics increased the risk of developing drug resistance [5, 6]. Therefore,
69 alternatives for conventionally used antibiotic have attracted considerable research
70 interest worldwide. The most widely studied alternatives include probiotics,
71 prebiotics, enzymes, plant extracts, and nutraceuticals such as copper and zinc [7].

72 Oligosaccharides are carbohydrates of low degree of polymerization (DP) and
73 low molecular weight composed of monosaccharides [8]. Oligosaccharides usually
74 resist enzymatic hydrolysis and absorption in upper gastrointestinal tract, but can be
75 fermented in large bowel by a number of bacteria [9]. Interestingly, non-digestible
76 oligosaccharides have prebiotic activity, as they can promote body health by
77 increasing populations of beneficial microbes and/or their metabolic activity [10].
78 Fructooligosaccharides is composed of fructose and glucose units, which refers
79 specifically to short chains (3–6 units) of fructose units bound by β -(2–1) linkages
80 that are attached to a terminal glucose unit [11]. Oligosaccharides including the FOS
81 have been previously reported to improve growth performance and gut health in a

82 variety of animal species [12]. For instance, FOS not only promoted the growth of
83 weaning pigs, but also effectively reduced the colonization of pathogenic bacteria in
84 the intestine [10].

85 ETEC is a major cause of diarrhea in neonatal and weaning pigs [13]. ETEC
86 colonized in the intestine can release enterotoxins, which stimulates secretion of fluid
87 from the epithelial cells into the lumen, resulting in acute diarrhea [14]. On the other
88 hand, weaning deprives neonatal pigs from passive immune protection of mother's
89 milk, which increases their susceptibility to enterotoxigenic *E. coli* infection [15, 16].
90 Although, numerous studies indicated that FOS have prebiotic effects that can inhibit
91 the growth of pathogenic bacteria [17], only few reports indicated the influences of
92 FOS on the growth and intestinal health in weaned pigs exposure to ETEC challenge.
93 In this study, we explored the effect of dietary FOS supplementation on growth
94 performance, inflammatory response, intestinal epithelium function, and selected
95 bacterial populations in weaned pigs upon ETEC challenge. This study could also
96 provide convincing evidence on novel prebiotic effect of FOS and offer key insights
97 into its potential mechanisms of action.

98 **Methods**

99 **Animals, feeding and experimental design**

100 All the procedures used in the animal experiment were approved by the Institutional
101 Animal Care and Use Committee of Sichuan Agricultural University (no. 20181105).
102 Twenty-four weaned pigs with an average initial body weight (6.30 ± 0.30 kg) pigs

103 were randomly allotted into three treatment groups that included (1) non-challenged
104 control (CON; pigs fed a basal diet and infused with sterilized Luria–Bertani culture);
105 (2) ETEC-challenged control (ECON; pigs fed the same basal diet and infused with
106 ETEC); and (3) ETEC challenge + FOS treatment (EFOS, ETEC + FOS; pigs fed a
107 basal diet supplemented with 2.5 mg/kg FOS and infused with ETEC). FOS,
108 purchased from Shanghai Lanpu Biotechnology CO., LTD., (FOS \geq 20%). The basal
109 diet (Table 1) was formulated to meet the nutrient requirements recommended by the
110 National Research Council 2012 [18]. Pigs were individually housed in metabolism
111 cages (0.7 m \times 1.5 m) with room temperature maintained at 25–28 °C and relative
112 humidity controlled at 55–65%. All pigs were given an ad libitum access to fresh
113 water and feeding. The trial lasted for 21 d. On day 19, pigs in the ECON and EFOS
114 group were orally administered 150 mL of Luria–Bertani culture containing
115 approximately 2.5×10^9 CFU/mL of ETEC (serotype O149:K91:K88ac; China
116 Institute of Veterinary Drugs Control, Beijing, China) [19], while pigs in the CON
117 group were administered an equal volume of sterilized Luria–Bertani culture. All
118 pig's feeding consumption was measured daily and BW was measured on day 19 and
119 22 after 12-h fasting at the morning 8:00. The average daily body weight gain (ADG),
120 average daily feed intake (ADFI) and gain-to-feed ratio (G : F) were calculated.

121 **Sample collection and preparation**

122 At the beginning of the trial, representative feed samples of each group were sampled
123 and stored at -20 °C for chemical analysis. From day 12 to day 15 of the experiment,
124 fresh fecal samples were collected immediately after excretion from eight randomly

125 selected pigs in each group. After collection, the daily excreta of each pig was
126 weighted, and 10 mL of a 10% H₂SO₄ solution was added to each 100 g of wet fecal
127 sample, and subsequently stored in a sealed plastic bag at -20 °C. At the end of the
128 4-d period, all fecal samples of each pig were thawed at room temperature and mixed
129 thoroughly, and then dried at 65 °C for 48 h, after which they were ground to pass
130 through a 1-mm screen and stored at -20 °C for chemical analyses including DM, CP,
131 ether extract (EE), ash, and GE. Blood samples were collected by venepuncture at
132 8:00 after 12 h of fasting. Then the samples were centrifuged at 3500 × g at 4 °C for
133 10 min [20]. After centrifugation, the plasma were prepared by using heparin sodium
134 anticoagulant and frozen at -20 °C until analysis. After blood collection, pigs were
135 euthanized with an intravenous injection of sodium pentobarbital at a dose of 200
136 mg/kg BW [21] and then slaughtered by exsanguination protocols, the mid points of
137 the duodenum, jejunum and ileum of each pig were harvested and fixed in 4%
138 paraformaldehyde solution for morphological analyses. Besides, 2 g digesta from the
139 cecum was immediately snap-frozen in liquid nitrogen, and stored at -80 °C for
140 analysis of microbial DNA and Short-chain fatty acid. At the same time, mucosa
141 samples were scraped with a scalpel blade from duodenum, jejunum and ileum
142 segments and quickly-frozen using liquid N₂, following by the preservation at -80°C
143 until further analysis for related enzyme activity and gene expression.

144 **Measurement of growth performance**

145 Piglets were weighed before the morning feed on day 1, 19, and 22 after 12 h fasting
146 at the morning 8:00. Feed intake was recorded, orts were collected and weighed daily.
147 All feed was mash. ADFI, ADG, and G:F were calculated.

148 **Determination of the apparent total tract digestibility**

149 The apparent total tract digestibility (ATTD) was measured using acid-insoluble ash
150 (AIA) as indicator [22]. The AIA in diets and faeces samples was determined by a
151 method described by Chinese National Standard (GB/T23742). All samples were
152 analyzed for DM, CP, EE, ash. GE content of diets and fecal samples was
153 determined using an adiabatic bomb calorimeter. Gross energy was determined
154 using an automatic adiabatic bomb calorimeter (LECO, St. Joseph, Michigan, USA).
155 The ATTD was calculated as $(100 - A1F2/A2F1 \times 100)$, where A1 represents the AIA
156 content of the diet; A2 represents the AIA content of faeces; F1 represents the
157 nutrient content of the diet; F2 represents the nutrient content of faeces.

158 **Plasma biochemical analysis**

159 Plasma concentrations of D-lactate, Immunoglobulin (Ig), including IgA, IgG and
160 IgM were determined following the procedures outlined by the corresponding kit
161 manufacturer using commercially available swine Enzyme-Linked Immunosorbent
162 Assay (ELISA) kits (Jiangsu Jingmei Biotechnology Co., Ltd., Yancheng, China).

163 **Intestinal morphology analysis**

164 About 1-cm segment of the small intestine (duodenum, jejunum and ileum) were
165 mixed in 10% neutral buffered formaldehyde. The mixed tissue samples were

166 dehydrated with normal saline and then embedded in paraffin. Cross sections of each
167 sample were prepared, stained with haematoxylin and eosin (H&E) and then sealed by
168 a neutral resin size. Ultrathin sections of the duodenal, jejunal and ileal samples were
169 examined for the villus height and crypt depth with image processing and analysis
170 system (Media Cybernetics, Bethesda, MD, USA). Villus height was calculated from
171 the tip of the villi to the villus-crypt junction. Crypt depth was expressed as the
172 invaginated depth between adjacent villi. A total of 10 intact, well-oriented
173 crypt-villus units were analyzed in triplicate per segment. The ratio of villus height to
174 crypt depth (V/C) was calculated from the values described above.

175 **Determination of SIgA-Positive Cells**

176 Paraffin-embedded jejunal tissues were cut into 4 μm thin and deparaffinized with
177 xylene and then dehydrated using a descending alcohol gradient. Sections were
178 incubated with 3% hydrogen peroxide for 10 minutes at room temperature and then
179 rinsed three times for 5 minutes with phosphate-buffered saline (PBS). After that,
180 sections were heated up to a boil in 10 mM citrate buffer (pH 6.0), washed 2 times in
181 PBS for 5 minutes, and incubated with 10% goat serum albumin for 20 min at room
182 temperature. Subsequently, sections were incubated at 4°C overnight with primary
183 antibodies for anti-SIgA. PBS instead of primary antibody was employed as negative
184 control. After being washed with PBS, the sections were incubated with biotinylated
185 goat anti-rabbit secondary antibodies (Beijing ZhongShan Golded Bridge
186 Biotechnology Co., Ltd., Beijing, China) for 30 min at 37 °C. Sections were washed
187 three times with PBS for 5 minutes and incubated with 3,3-diaminobenzidine (DAB)

188 to visualize immune complexes. The sections were then counterstained with
189 hematoxylin and mounted with glycerol gelatin. Images were obtained using BA400
190 Digital microscope (Motic China Group Co., Ltd.). The integrated optical density was
191 detected using Image-Pro Plus 6.0 image analysis system (Media Cybernetics,
192 Bethesda, MD, USA), and the protein expression was reflected by the mean value of
193 the integrated optical density.

194 **Determination of intestinal enzyme activities**

195 After thawing, about 1 g of intestinal mucosa (including the duodenum, jejunum and
196 ileum) was homogenized in ice-cold physiological saline at 1:9 and then centrifuged
197 at $3000 \times g$, 4 °C for 15 min. The supernatant was assayed for protein content using
198 the Bradford method, followed by measurement of the sucrase, lactase maltase and
199 AKP activities, in accordance with the instructions accompanying the respective kit
200 (Nanjing Jiancheng Bioengineering Institute). The results were normalised to protein
201 concentration and expressed as U/mg protein.

202 **Intestinal microbial populations measurement**

203 Intestinal microbial populations measurement Microbial DNA was extracted from the
204 caecal digesta using the EZNA® Stool DNA kit (Omega Bio-Tek, Doraville, CA,
205 USA), according to the manufacturer's instructions. Based on the 16S rRNA
206 sequences of maximum species of each genus encountered in the pig intestinal tract
207 downloaded from the National Center for Biotechnology Information (GenBank),
208 European Molecular Biology Laboratory and DNA Data Bank of Japan, The

209 fluorescent quantitative specific primers and probe for total *E. coli*, *Lactobacillus*,
210 *Bifidobacterium*, and *Bacillus* (Table S1) were obtained from the Fierer et. al [23].
211 and Qi et. al [24] and were commercially synthesised by Invitrogen (Shanghai,
212 China). To quantify *Bacillus*, *Bifidobacterium*, *E. coli* and *Lactobacillus*, qPCR was
213 carried out on a CFX96 Real-Time PCR system (Bio-Rad Laboratories, Inc.,
214 Hercules, CA) using the RealMasterMix (Probe) kit (Tiangen Biotech Co., Ltd.,
215 Beijing, China). A volume of 25 μL , containing 12.5 μL of SuperReal PeMix (2.0 \times),
216 1 μL each of the forward and reverse primers, 1 μL of Fluorescent probe (20 \times), 1 μL
217 of ROX Reference*3 Dye (50 \times), 2 μL of DNA and 6.5 μL of RNase-Free ddH₂O
218 were used in each reaction. The reactions were subjected to 1 cycle at 95 $^{\circ}\text{C}$ for 15
219 min, followed by 39 cycles at 95 $^{\circ}\text{C}$ for 3 s, 58 $^{\circ}\text{C}$ for 25 s and 72 $^{\circ}\text{C}$ for 60 s. The
220 cycle threshold (Ct) values and baseline settings were determined by automatic
221 analysis settings, and the copy numbers of the target group for each reaction were
222 calculated from the standard curves, which were generated by constructing standard
223 plasmids by a 10-fold serial dilution of plasmid DNA (1×10^1 to 1×10^9 copies/ μL).

224 **Volatile fatty acid (VFA) analysis**

225 Samples of digesta from individual pigs were taken from the caecum to measure the
226 VFA concentration. The VFA concentrations in the digesta were determined using gas
227 liquid chromatography according to the method described by Pierce et al [25].
228 Approximately 1.0 g of thawed cecal digesta was suspended in 1.5 mL of sterile
229 milli-Q water in a centrifuge tube for 30 min. The entire sample was centrifuged at
230 $12,000 \times g$ for 10 min. 1 mL supernatant was transferred to a sterile tube, and then

231 mixed with 0.2 mL 25% metaphosphoric acid and 23.3 μ L 210 mmol/L crotonic acid
232 simultaneously, the sterile tubes were centrifuged again for 10 min after placed the
233 sterile tubes in ice-bath for 30 min. The mixture of 500 μ L supernatant and 500 μ L
234 methanol was homogenized for 10 min in another sterile tube. After that, the mixture
235 was centrifuged at $12,000 \times g$ for 10 min at 4°C. The supernatant was injected into a
236 gas chromatographic system (VARIAN CP-3800, America) to separate and quantify
237 the VFA.

238 **RNA isolation, reverse transcription, and real-time quantitative PCR**

239 Total RNA was extracted from duodenum, jejunum and ileum mucosa using the
240 Trizol Reagent (TaKaRa, Dalian, China). Meanwhile, the concentration and purity of
241 total RNA were assayed by spectrophotometer (Nano Drop, Gene Company Limited,
242 Guangzhou, China) at 260 and 280 nm following manufacturer's guidelines. The ratio
243 of OD 260/280 should vary between 1.8 and 2.0. Reverse transcription using the
244 Prime Script RT reagent kit (TaKaRa Biotechnology, Dalian, China) was exploited
245 following the manufacturer's instructions. The primers were synthesized
246 commercially by Life Technologies Limited and were exhibited in (Table S1).

247 Quantitative real-time polymerase chain reaction (PCR) was conducted to analyse
248 the mRNA expression abundance of TNF- α , IL-1 β , IL-6, ZO-1, Occludin, FATP1,
249 FATP4, CAT1, LAT1, SGLT1 and GLUT2 in the small intestinal mucosa using the
250 CFX-96 real-time PCR detection system (Bio-Rad) and SYBR Premix Ex Taq II (Tli
251 RNaseH Plus) reagents (TaKaRa, Dalian, China). The PCR reaction was run in a 10

252 μL reaction volume, which contained 5 μL of SYBR Premix Ex Taq II (Tli RNaseH
253 Plus), 0.5 μL of each primer, 1 μL of the cDNA sample, and 3 μL of nuclease-free
254 water. The PCR cycling parameters were as follows: initial denaturation at 95 °C for
255 30 s, followed by 40 cycles of 95 °C for 5 s, 57.5 °C for 30 s, and 72 °C for 5 min. A
256 melting curve analysis was performed following each real-time quantitative PCR
257 assay to confirm the Gene-specific amplification products had been generated. The
258 housekeeping gene β -actin was used as an internal control for normalization. The
259 target gene mRNA expression level was calculated using the $2^{-\Delta\Delta\text{Ct}}$ method [26]. Each
260 sample was simultaneously performed on the same PCR plate and three replicates
261 were set up.

262 **Statistical analysis**

263 All data were subjected to one-way analysis of variance for a completely randomised
264 design using the general linear model procedure of SPSS 24.0 (SPSS, Inc.), with each
265 pig representing one experimental unit. Statistical differences among treatments were
266 separated by Tukey's multiple-range test. The results are expressed as with their
267 standard errors. Statistical significance was set at $P < 0.05$ and $0.05 < P < 0.10$
268 indicating a trend.

269 **Results**

270 **Effect of FOS supplementation on growth performance and nutrient digestibility**

271 As shown in Table 2, there were no differences ($P > 0.05$) in ADG, ADFI, and G:F
272 among the three treatments during days 1–18 (pre-challenge). ETEC challenge

273 (during days 19–21) significantly reduced the ADG ($P < 0.05$) in the weaned pigs,
274 however, the ADG was significant higher in the EFOS group than in the ECON group
275 ($P < 0.05$). Moreover, FOS supplementation significantly decreased the diarrhea
276 incidence in the ETEC-challenge pigs (Table 2). As compared to the CON and ECON
277 group, dietary FOS supplementation significantly elevated the apparent digestibility
278 of CP, GE, and ash ($P < 0.05$). FOS supplementation did not significant influence on
279 the digestibility of DM (Table 3).

280 **Effect of FOS supplementation on concentrations of plasma cytokines and** 281 **immunoglobulins in weaned pigs upon ETEC challenge**

282 As shown in Table 4, ETEC challenge significantly increased the concentrations of
283 TNF- α in the plasma ($P < 0.05$). However, dietary FOS supplementation significantly
284 decreased the concentrations of plasma IL-1 β and TNF- α in the pigs upon ETEC
285 challenge ($P < 0.05$). Moreover, the concentrations of plasma IgA and IgG were lower
286 in the ECON group than in the CON group ($P < 0.05$). However, the concentrations
287 of plasma IgA and IgM were significantly higher in the EFOS group than in the
288 ECON group ($P < 0.05$).

289 **Effect of FOS supplementation on intestinal epithelium morphology in weaned** 290 **pigs upon ETEC challenge**

291 ETEC challenge significantly decreased the duodenal villus height, increased the
292 duodenal crypt depth in the ECON group compared to the CON group (Table 5,
293 Figure. 1). The duodenal villus height was significant higher in the EFOS group than

294 ECON ($P < 0.05$). Interestingly, ETEC significantly decreased the duodenal ratio of
295 villus height to crypt depth (V/C) in the ECON group compared to the CON group (P
296 < 0.05). As compared to the ECON group, FOS supplementation significantly
297 increased the V/C ratio in the duodenum and ileum ($P < 0.05$).

298 **Effect of FOS on intestinal permeability and the mucosal immunity**

299 As shown in Figure. 2A, compared to ECON group, the D-lactate content in the
300 plasma were significantly decreased ($P < 0.05$) in the EFOS group pigs. Moreover,
301 we determined the jejunal SIgA secretion by the immunohistochemical analysis,
302 indicating that FOS supplementation increased the counts of SIgA cell in the jejunum
303 of pigs (Fig. 2B). Besides, we detected the mRNA expression levels of cytokines in
304 the small intestine of pigs. As shown in Fig. 3, FOS supplementation down-regulated
305 ($P < 0.05$) the mRNA expression of IL-6 and TNF- α in jejunum and ileum as well as
306 IL-1 β and TNF- α in duodenum.

307 **Effect of FOS on enzyme activity of intestinal mucosa in weaned pigs upon** 308 **ETEC challenge**

309 As shown in Table 6, ETEC challenge significantly decreased the mucosal activities
310 of sucrase, lactase, and AKP in the duodenum ($P < 0.05$). The mucosal activity of
311 lactase in the ileum was also lower in the ECON group than in the CON group ($P <$
312 0.05). However, FOS supplementation not only elevated duodenal the activities of
313 sucrase, lactase, and AKP in the duodenum, but also elevated the ileal activity of
314 lactase in the ETEC challenged pigs ($P < 0.05$).

315 **Effect of FOS supplementation on microbial metabolites and intestinal**
316 **populations in weaned pigs upon ETEC challenge**

317 As shown in Figure.4, ETEC challenge decreased the concentrations of acetic acid
318 and propionic acid in cecal digesta ($P < 0.05$). However, FOS supplementation
319 significantly elevated their concentrations in the ETEC challenged pigs ($P < 0.05$).
320 The concentration of butyric acid was also higher in the EFOS group than in the
321 ECON group ($P < 0.05$). ETEC challenge increased the abundance of *E. coli* in the
322 cecal digesta; however, FOS supplementation significantly decreased the abundance
323 of *E. coli* in the cecal digesta (Table 7). Moreover, FOS supplementation significantly
324 elevated the abundance of beneficial microflora such as the *Bifidobacterium* and
325 *Bacillus* in the cecal digesta of ETEC-challenged pigs ($P < 0.05$).

326 **Effect of FOS supplementation on expressions of critical genes related to**
327 **intestinal epithelium functions**

328 As shown in Figure.5, ETEC challenge decreased the expression levels of
329 tight-junction protein ZO-1 in the duodenum. However, FOS supplementation not
330 only elevated the expression levels of ZO-1 in the duodenum and jejunum, but also
331 elevated the expression levels of occludin and FATP4 in the ileum ($P < 0.05$).
332 Moreover, FOS supplementation elevated the expression levels of LAT1 in the
333 duodenum and ileum ($P < 0.05$). The expression level of SGLT1 in the ileum, and the
334 expression levels of GLUT2 in the jejunum and ileum were higher in the EFOS group
335 than in the ECON group ($P < 0.05$).

336 **Discussion**

337 Antibiotics are widely used for preventing post-weaning diarrhea in pig production
338 industry [7]. However, long-term or overdose utilization of antibiotics may lead to
339 developing of bacterial resistance and drug residues in the product [5,6].
340 Fructooligosaccharides are highly interesting prebiotic fibres, resulting from the
341 transfructosylating action of specific microbial enzymes on sucrose, composed of one
342 molecule of glucose linked to 2 to 4 fructose units at position β -1,2 that have been
343 shown to improve weaned pigs' intestinal morphology and growth performance [10,
344 12]. In the present study, we investigated the protective potential of FOS against
345 ETEC-induced inflammation and intestinal injury in weaned pigs. We showed that
346 dietary FOS supplementation significantly improved the growth performance,
347 apparent nutrient digestibility and reduced the diarrhea incidence in pigs exposure to
348 ETEC challenge. The results are consistent with previous studies on pigs [27,28].

349 An integrated intestinal morphological structure is of critical importance for
350 nutrient digestion and absorption. D-Lactic acid is a metabolite produced by
351 fermentation of bacteria in the gastrointestinal tract that is rarely absorbed into the
352 bloodstream under normal circumstances [29]. However, large amounts of D-lactic
353 acid can be absorbed into the bloodstream through the damaged intestinal mucosa, so
354 plasma D-lactic acid can serve as a marker of maturity, integrity, and functional status
355 of intestinal epithelial cells [30,31]. In the present study, we found that FOS
356 supplementation reduced the D-lactic acid content in the plasma, indicating that FOS
357 can alleviate the detrimental effects of ETEC in pig's intestine which might be related

358 to the FOS enhanced the maturity, integrity, and functional status of intestinal. ETEC
359 challenge significantly decreased the duodenal villus height, and reduced the ratio of
360 V/C in the duodenum and ileum, indicating injury of the intestinal epithelium. The
361 result is consistent with previous study that ETEC strains function through producing
362 enterotoxins that act on the small intestines and lead to the secretion of fluids and
363 electrolytes, causing diarrhea and intestinal injury [32]. However, dietary FOS
364 supplementation attenuated the intestinal injury by increasing the villus height and the
365 ratio of V/C. The result is probably due to the prebiotic effects of FOS. For instance,
366 the FOS were found to prevent the adhesion of enteric pathogens to intestinal
367 epithelium, which significantly reduced their colonization and intestinal inflammation
368 and injury [33]. Moreover, FOS were found to improve the intestinal morphology and
369 significantly reduced the colonization of pathogenic bacteria in the intestine [34]. The
370 beneficial effects of FOS supplementation on intestinal epithelium function have also
371 been indicated by the mucosal enzyme activities, as FOS not only elevated duodenal
372 the activities of sucrase, lactase, and AKP, but also elevated the ileal activity of
373 lactase in the ETEC-challenged pigs. Sucrase and lactase are two important digestive
374 enzymes involved in carbohydrate digestion, play crucial roles in the process of
375 digestion in animals [35]. While, the AKP is an another crucial endogenous enzyme
376 expressed in the brush border, and considered as an excellent marker enzyme
377 involved in the primary digestive and absorptive processes of the small intestine [36].

378 SIgA is a 400 kDa molecule composed of the secretory components, J-chain and
379 dimeric IgA [37]. As a major component of the intestinal immune barrier, SIgA plays

380 an important role in maintaining intestinal health by clearing pathogenic
381 microorganisms and interacting with intestinal commensal microorganisms [38,39].
382 Furthermore, the host may discriminate symbionts from pathogens by recognizing the
383 coating of commensal bacteria by SIgA [40]. Thus, enough SIgA secretion in the gut
384 is essential for the intestinal homeostasis. However, piglets, especially weaning
385 piglets, are unable to get maternal immunoglobulins and usually cannot secrete enough
386 SIgA because of their underdeveloped intestinal immune system [41]. Interestingly, in
387 the current study, results of the immunohistochemistry analyses confirmed that FOS
388 supplementation increased the jejunal SIgA levels in EFOS group piglets, which
389 could be associated with a key role of supplementation of FOS stimulating the
390 development and maturation of the intestinal mucosal.

391 Cytokines act as crucial roles in the immune and inflammatory responses [42].
392 Over-production of TNF- α , IL-1 β and IL-6 have shown a series of proinflammatory
393 functions in inflammatory intestinal mucosa, further induces the dysfunction of the
394 intestinal barrier, causing the death of intestinal cells [43,44]. In the present study,
395 FOS supplementation down-regulated the mRNA expression of IL-6 and TNF- α in
396 jejunum and ileum as well as IL-1 β and TNF- α in duodenum, which suggested an
397 anti-inflammatory role of FOS in regulating the ETEC-induced inflammation. The
398 result is also consistent with previous study that mannan oligosaccharides added in the
399 sow diet enhance their offspring intestinal immunity by decreasing inflammation [45].
400 Both these results suggested a beneficial role of FOS in regulating the intestinal

401 immune function of pigs under ETEC injection. Our results are in line with the
402 present study that FOS suggest a positive effect on mucosal immunity. [46].

403 Both the IgA and IgG are crucial index of body immune capacity and contribute
404 to dealing with various pathogens. IgA is the most abundant immunoglobulin in the
405 body and is of critical importance in body immunity, plays a critical role in
406 maintaining intestinal mucosal immunity and preventing intestinal infection [47]. In
407 general, IgG enhances mucosal homeostasis and controls non-invasive and invasive
408 mucosal bacteria by reaching the lumen of mucosal organs upon binding to the
409 epithelial transporter FcRn [48]. Moreover, the plasma concentrations of IgA and IgG
410 were elevated by FOS in the ETEC-challenged pigs, which might be related to FOS
411 stimulate the growth and development of immune organs, promote the differentiation
412 of lymphocyte and enhance the cellular immunity and humoral immunity [49], our
413 result was consistent with the present study that FOS supplementation elevate the IgA
414 and IgG in the plasma [50].

415 In the present study, we also investigated the concentration of microbial
416 metabolites and abundance of major bacterial populations in the cecum. As expected,
417 FOS supplementation significantly elevated the concentrations of short chain fatty
418 acids (SCFAs, e.g. acetic acid, propionic acid, and butyric acid) in the cecal digesta of
419 ETEC-challenged pigs. The elevated SCFAs concentration is probably due to the fact
420 that oligosaccharides cannot be digested in the upper digestive tract, but can be
421 fermented by intestinal microorganisms in the lower diegstive tract [51]. Both the
422 acetic and propionic acids can be used as energy substrates for peripheral tissues.

423 While the butyrate not only acts as a critical energy source for intestinal epithelial
424 cells, but can also promote cell proliferation and differentiation [52]. In the present
425 study, we found FOS decreased the abundance of *E. coli*, but significantly increased
426 the abundance of *Bifidobacterium* and *Bacillus* in the cecum. This is probably due to
427 the fact that pathogenic bacteria such as the *E. coli* can not utilize the prebiotics
428 [53,54]. Moreover, the fermented products by beneficial bacteria provide an acidic
429 environment that is important for inhibiting the growth of harmful bacteria [55]. The
430 result is consistent with previous reports that FOS can regulate the balance of
431 intestinal microbial flora [56]. Both these results suggested a beneficial role of
432 prebiotics in regulating the intestinal microbial ecology and health.

433 To gain insights into the mechanisms underlying the FOS-regulated intestinal
434 health, we further investigated the expression levels of several critical genes involved
435 in the intestinal epithelium functions. Tight junctions play a critical role in
436 maintaining the intestinal barrier integrity, which can prevent the paracellular
437 diffusion of intestinal bacteria and other antigens across the epithelium [57].
438 Tight-junction proteins such as the ZO-1 and occludin are critical components for
439 structural and functional organization of the tight junctions. In this study, FOS
440 supplementation not only elevated the expression levels of ZO-1 in the duodenum and
441 jejunum, but also elevated the expression levels of occludin and FATP4 in the ileum.
442 FATP4 is a small molecule protein that is responsible for transportation of fatty acids
443 across the cell membranes [58]. Moreover, FOS supplementation significantly
444 elevated the expression levels of nutrient transporters (LAT1, CAT1, SGLT1, and
445 GLUT2) in the EFOS group. The LAT1 and CAT1 are transporters that is responsible
446 for transporting of the L-type and cationic amino acids, respectively [59]. While the

447 SGLT1 and GLUT2 are closely associated with glucose absorption [60]. The elevated
448 expressions of these nutrient transporters indicated an improved intestinal integrity
449 and epithelium functions in pigs exposure to ETEC challenge.

450 **Conclusions**

451 The present study suggested that dietary FOS supplementation can attenuate the
452 growth retardation in the weaned pigs exposure to ETEC challenge. Moreover, FOS
453 can alleviate the ETEC-induced intestinal injury in the weaned pigs, which was
454 associated with suppressing of the inflammatory responses and improving of the
455 intestinal epithelium functions and intestinal microbiota. The beneficial effects of
456 FOS on the growth performance and intestinal health should make it an attractive
457 prebiotic that can be tentatively used for the substitution of antibiotics.

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462 **Data Availability and materials**

463 The datasets used to support the findings of this study are available from the
464 corresponding author upon request.

465 **Authors' contributions**

466 LL, JH and HY conceived the study, performed the experiment, performed data
467 analysis, and contributed to drafting the manuscript. LL carried out the animal

468 experiment. DWC, BY, ZQH, YHL, PZ, XBM, JY, JQL, HY conceived the
469 experiment and proofread the manuscript. All authors read and approved the final
470 manuscript.

471 **Conflicts of Interest**

472 The authors declare that there are no conflicts of interest.

473 **Consent for publication**

474 Not applicable.

475 **Ethics approval and consent to participate**

476 All animal procedures were performed in accordance with the Guidelines for Care and
477 Use of Laboratory Animals of Sichuan Agricultural University and approved by the
478 Animal Ethics Committee of Sichuan Agricultural University (No. 20181105).

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482

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631

632 **Figure legends**

633 **Fig. 1 Effect of FOS on small intestinal morphology in weaned pigs.** (H&E; $\times 40$).

634 Non-challenged pigs (CON, fed with basal diet), ETEC-challenged pigs (ECON, fed
635 with basal diet), FOS and ETEC-treated pigs (EFOS, fed with basal diet containing
636 2.5 g/kg FOS challenged by ETEC).

637 **Fig. 2 Effect of FOS on intestinal permeability and the mucosal immunity** Values
638 are means \pm SEM, (n = 6), non-challenged pigs (CON, fed with basal diet),
639 ETEC-challenged pigs (ECON, fed with basal diet), and FOS and ETEC-treated pigs
640 (EFOS, fed with basal diet containing 2.5 g/kg FOS challenged by ETEC). ^{2 a, b, c}

641 Mean values within a row with unlike superscript letters were significantly different
642 ($P < 0.05$)

643 **Fig. 3 Effects of FOS on mRNA levels of intestinal inflammatory cytokines.**

644 Duodenum (A); Jejunum (B), and Ileum (C). Values are means \pm SEM, (n = 6),
645 non-challenged pigs (CON, fed with basal diet), ETEC-challenged pigs (ECON, fed
646 with basal diet), and FOS and ETEC-treated pigs (EFOS, fed with basal diet
647 containing 2.5 g/kg FOS challenged by ETEC). ^{2 a, b, c} Mean values within a row with
648 unlike superscript letters were significantly different ($P < 0.05$); IL-1 β , interleukin-1 β ;
649 IL-6, interleukin-6; TNF- α , tumour necrosis factor- α .

650 **Fig. 4 Effect of FOS on Concentration of Intestinal VFAs.** Acetic acid

651 concentration (A); Propionic acid concentration (B); Butyric acid (C);^{a,b,c} Mean values

652 with different letters on vertical bars indicate significant differences ($P < 0.05$).

653 Non-challenged pigs (CON, fed with basal diet), ETEC-challenged pigs (ECON, fed
654 with basal diet), FOS and ETEC-treated pigs (EFOS, fed with basal diet containing
655 2.5 g/kg FOS challenged by ETEC).

656 **Fig. 5 Relative expression levels of critical genes involved in the intestinal barrier**
657 **functions.** ZO-1, zonula occludens-1; FATP1, Fatty acid transport protein-1; FATP4,
658 Fatty acid transportprotein-4; LAT1, L-type amino acid transporter-1; CAT1, cationic
659 amino acid transporter-1; SGLT1, sodium glucose transport protein-1; GLUT2,
660 glucose transporter-2; ^{a,b,c} Mean values with different letters on vertical bars indicate
661 significant differences ($P < 0.05$). Non-challenged pigs (CON, fed with basal diet),
662 ETEC-challenged pigs (ECON, fed with basal diet), FOS and ETEC-treated pigs
663 (EFOS, fed with basal diet containing 2.5 g/kg FOS challenged by ETEC).

664

665 **Tables**666 **Table 1.** Composition and nutrient level of experimental diet

| Ingredients | % | nutrient level | contents |
|-----------------------------|-------|---------------------------------------|----------|
| Corn | 28.31 | Digestible energy (calculated, MJ/kg) | 14.78 |
| Extruded corn | 24.87 | Crude Protein (%) | 19.68 |
| Soybean meal | 8.50 | Calcium (%) | 0.81 |
| Extruded full-fat soybean | 10.30 | Available phosphorus (%) | 0.55 |
| Fish meal | 4.20 | Lysine | 1.35 |
| Whey powder | 7.00 | Methionine | 0.42 |
| Soybean protein concentrate | 8.00 | Methionine + cysteine | 0.60 |
| Soybean oil | 2.00 | Threonine | 0.79 |
| Sucrose | 4.00 | Tryptophan | 0.22 |
| Limestone | 0.90 | | |
| Dicalcium phosphate | 0.50 | | |
| NaCl | 0.30 | | |
| L -LysineHCl (78%) | 0.47 | | |
| DL -Methionine | 0.15 | | |
| L -Threonine (98.5%) | 0.13 | | |
| Tryptophan (98%) | 0.03 | | |
| Chloride choline | 0.10 | | |
| Vitamin premix ¹ | 0.04 | | |
| Mineral premix ² | 0.20 | | |
| Total | 100 | | |

667 ¹ The vitamin premix provided the following per kg of diet: 9000 IU of VA, 3000 IU of VD 3, 20 IU of
668 VE, 3 mg of VK 3, 1.5 mg of VB1, 4 mg of VB 2, 3 mg of VB6, 0.02 mg of VB12, 30 mg of niacin, 15
669 mg of pantothenic acid, 0.75 mg of folic acid, and 0.1 mg of biotin. ² The mineral premix provided the
670 following per kg of diet: 100 mg Fe, 6 mg Cu, 100 mg Zn, 4 mg Mn, 0.30 mg I, 0.3 mg Se.

671

672 **Table 2.** Effect of FOS on growth performance in weaned pigs challenged with
 673 enterotoxigenic *Escherichia coli*

| Items | Treatments | | | <i>P</i> -Value |
|------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------|
| | CON | ECON | EFOS | |
| pre-challenged | | | | |
| (1–19 days) | | | | |
| ADFI(g/day) | 430.70 ± 30.80 | 403.49 ± 26.27 | 417.71 ± 38.79 | 0.96 |
| ADG (g/day) | 296.56 ± 14.63 | 270.67 ± 16.74 | 285.00 ± 27.32 | 0.68 |
| F:G (g/g) | 1.45 ± 0.05 | 1.60 ± 0.16 | 1.47 ± 0.04 | 0.76 |
| post-challenged | | | | |
| (19–21 days) | | | | |
| ADFI (g/day) | 501.40 ± 13.40 | 463.73 ± 3.45 | 484.67 ± 8.10 | 0.05 |
| ADG (g/day) | 396.00 ± 23.98 ^a | 314.67 ± 48.42 ^b | 381.80 ± 22.14 ^a | 0.04 |
| Diarrhea incidence (%) | 33.00 ± 2.15 ^b | 61.07 ± 5.89 ^a | 23.33 ± 4.76 ^b | < 0.01 |

674 ¹ Values are means ± SEM, (n = 6), non-challenged pigs (CON, fed with basal diet), ETEC-challenged
 675 pigs (ECON, fed with basal diet), and FOS and ETEC-treated pigs (EFOS, fed with basal diet
 676 containing 2.5 g/kg FOS challenged by ETEC). ² ^{a, b, c} mean values within a row with unlike superscript
 677 letters were significantly different (*P* < 0.05). ³ ADFI = Average daily feed intake; ADG = Average
 678 daily gain; G/F = the ratio of gain to feed intake. ⁴ Incidence of diarrhea was calculated as follows:
 679 diarrhea incidence (%) = (total number of pigs with diarrhea)/(number of pigs × 3) × 100%, where the
 680 number of pigs with diarrhea was the summation of the number of pigs with diarrhea every day.

681 .

682 **Table 3.** Effect of FOS on ATTD of nutrients in weaned pigs

| Item, % | Treatments | | | <i>P</i> -Value |
|-----------|---------------------------|---------------------------|---------------------------|-----------------|
| | CON | ECON | EFOS | |
| DM | 81.58 ± 1.41 | 83.65 ± 2.51 | 85.36 ± 0.61 | 0.21 |
| CP | 83.38 ± 2.40 ^b | 85.33 ± 0.84 ^b | 90.95 ± 0.86 ^a | 0.01 |
| GE | 87.98 ± 2.31 ^b | 86.91 ± 0.59 ^b | 93.04 ± 0.58 ^a | 0.03 |
| Crude fat | 79.62 ± 2.82 | 78.38 ± 5.06 | 88.60 ± 0.70 | 0.07 |
| Ash | 74.19 ± 0.66 ^b | 68.81 ± 1.16 ^b | 79.54 ± 0.38 ^a | < 0.01 |

683 Values are means ± SEM, (n = 6), non-challenged pigs (CON, fed with basal diet), ETEC-challenged
684 pigs (ECON, fed with basal diet), and FOS and ETEC-treated pigs (EFOS, fed with basal diet
685 containing 2.5 g/kg FOS challenged by ETEC). ² a, b, c mean values within a row with unlike superscript
686 letters were significantly different (*P* < 0.05). ³ DM, dry matter; CP, crude protein; GE, gross energy.

687

688 **Table 4.** Effect of FOS on plasma Immunoglobulin and Cytokine Concentrations in
 689 weaned pigs challenged with enterotoxigenic *Escherichia coli*

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| Items | Treatments | | | <i>P</i> -Value |
|-----------------------|----------------------------------|---------------------------------|---------------------------------|-----------------|
| | CON | ECON | EFOS | |
| IL-1 β (ng/L) | 456.20 \pm 8.28 ^a | 508.93 \pm 31.71 ^a | 392.40 \pm 10.19 ^b | < 0.01 |
| IL-6 (ug/L) | 1.34 \pm 0.14 | 1.44 \pm 0.13 | 1.35 \pm 0.08 | 0.76 |
| TNF- α (pg/mL) | 487.60 \pm 11.09 ^b | 589.52 \pm 52.31 ^a | 454.61 \pm 13.27 ^b | 0.03 |
| Ig A (ug/L) | 432.48 \pm 5.78 ^a | 357.52 \pm 15.60 ^b | 481.92 \pm 32.26 ^a | < 0.01 |
| Ig G (ug/L) | 515.70 \pm 28.15 | 447.79 \pm 17.40 | 508.71 \pm 14.55 | 0.08 |
| Ig M (ug/L) | 464.96 \pm 26.63 ^{ab} | 393.21 \pm 21.40 ^b | 518.15 \pm 30.48 ^a | 0.02 |

690 ¹ Values are means \pm SEM, (n = 6), non-challenged pigs (CON, fed with basal diet), ETEC-challenged
 691 pigs (ECON, fed with basal diet), FOS and ETEC-treated pigs (EFOS, fed with basal diet containing
 692 2.5 g/kg FOS challenged by ETEC). ² ^a, ^b, ^c mean values within a row with unlike superscript letters
 693 were significantly different (*P* < 0.05). ³ IgA, immunoglobulin A; IgG, immunoglobulin G; IgM,
 694 immunoglobulin M; IL-1 β , interleukin-1 β ; IL-6, interleukin-6 and TNF- α , tumor necrosis factor- α .

695

696 **Table 5.** Effect of FOS on intestinal morphology in weaned pigs challenged with
 697 enterotoxigenic *Escherichia coli*

| Items | Treatments | | | <i>P</i> -Value |
|--------------------|-----------------------------|-----------------------------|-----------------------------|-----------------|
| | CON | ECON | EFOS | |
| Duodenum | | | | |
| Villus height (µm) | 486.81 ± 22.81 ^a | 424.46 ± 62.95 ^b | 524.71 ± 67.59 ^a | 0.02 |
| Crypt depth (µm) | 345.09 ± 18.10 ^b | 372.68 ± 17.11 ^a | 301.78 ± 5.95 ^b | 0.04 |
| V/C | 1.42 ± 0.07 ^a | 1.12 ± 0.10 ^b | 1.73 ± 0.11 ^a | < 0.01 |
| Jejunum | | | | |
| Villus height (µm) | 430.81 ± 14.48 | 399.71 ± 12.26 | 438.98 ± 12.79 | 0.29 |
| Crypt depth (µm) | 173.29 ± 8.39 | 194.45 ± 11.36 | 194.97 ± 11.82 | 0.29 |
| V/C | 2.50 ± 0.11 | 2.28 ± 0.09 | 2.10 ± 0.18 | 0.13 |
| Ileum | | | | |
| Villus height (µm) | 341.94 ± 26.81 | 357.14 ± 20.34 | 366.00 ± 7.03 | 0.69 |
| Crypt depth (µm) | 178.42 ± 6.23 ^b | 203.97 ± 6.12 ^a | 159.78 ± 9.65 ^b | < 0.01 |
| V/C | 1.92 ± 0.13 ^{ab} | 1.84 ± 0.17 ^b | 2.32 ± 0.11 ^a | 0.04 |

698 ¹ Values are means ± SEM, (n = 6), non-challenged pigs (CON, fed with basal diet), ETEC-challenged
 699 pigs (ECON, fed with basal diet), and FOS and ETEC-treated pigs (EFOS, fed with basal diet
 700 containing 2.5 g/kg FOS challenged by ETEC). ² ^{a, b, c} mean values within a row with unlike superscript
 701 letters were significantly different (*P* < 0.05).

702

703 **Table 6.** Effect of FOS on enzyme activity of small intestine in weaned pigs challenged with enterotoxigenic *Escherichia coli*
 704

| Items | Treatments | | | P-Value |
|------------------------|-----------------------------|----------------------------|-----------------------------|---------|
| | CON | ECON | EFOS | |
| Duodenum | | | | |
| Sucrase, U/mg protein | 155.07 ± 20.71 ^a | 83.30 ± 22.67 ^b | 156.70 ± 16.11 ^a | 0.04 |
| Lactase, U/ mg protein | 12.07 ± 3.59 ^a | 6.21 ± 1.88 ^b | 13.51 ± 4.16 ^a | 0.28 |
| Maltase, U/mg protein | 21.11 ± 2.65 | 13.17 ± 2.23 | 23.71 ± 6.33 | 0.17 |
| AKP, U/mg protein | 1.44 ± 0.35 ^a | 0.65 ± 0.08 ^b | 1.50 ± 0.19 ^a | 0.04 |
| Jejunum | | | | |
| Sucrase, U/mg protein | 536.76 ± 43.76 | 475.36 ± 88.10 | 526.5 ± 84.74 | 0.83 |
| Lactase, U/ mg protein | 94.21 ± 26.96 | 57.84 ± 13.22 | 96.34 ± 21.48 | 0.39 |
| Maltase, U/mg protein | 576.75 ± 78.80 | 423.32 ± 113.97 | 544.7 ± 131.62 | 0.76 |
| AKP, U/mg protein | 1.21 ± 0.09 | 1.15 ± 0.08 | 1.22 ± 0.09 | 0.83 |
| Ileum | | | | |
| Sucrase, U/mg protein | 210.25 ± 34.41 | 153.42 ± 31.10 | 208.23 ± 17.69 | 0.32 |
| Lactase, U/ mg protein | 35.19 ± 6.69 ^a | 15.37 ± 3.10 ^b | 32.38 ± 5.39 ^a | 0.04 |
| Maltase, U/mg protein | 131.49 ± 18.75 | 110.36 ± 25.63 | 135.41 ± 8.82 | 0.62 |
| AKP, U/mg protein | 2.34 ± 0.20 | 2.06 ± 0.21 | 2.19 ± 0.11 | 0.57 |

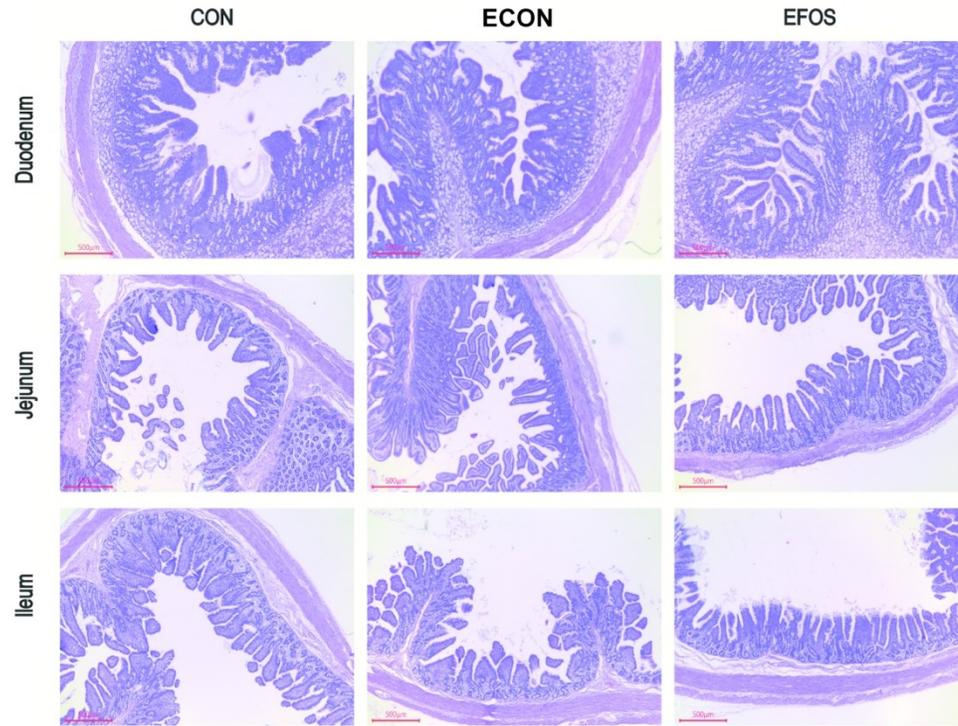
705 Values are means ± SEM, (n = 6), non-challenged pigs (CON, fed with basal diet), ETEC-challenged
 706 pigs (ECON, fed with basal diet), and FOS and ETEC-treated pigs (EFOS, fed with basal diet
 707 containing 2.5 g/kg FOS challenged by ETEC). ² a, b, c mean values within a row with unlike superscript
 708 letters were significantly different ($P < 0.05$). ³ AKP = alkaline phosphatase

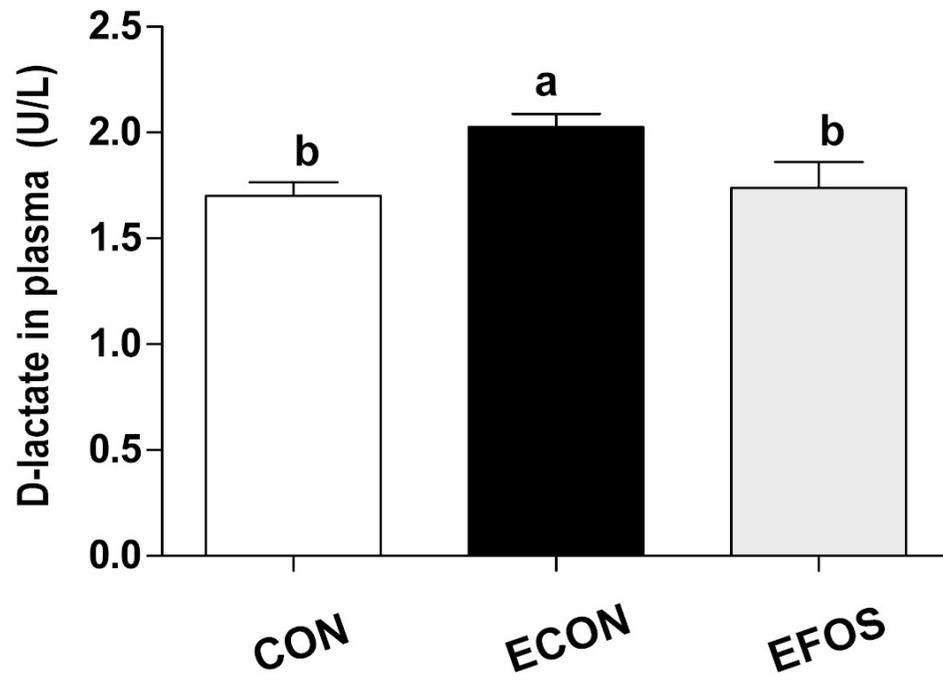
709

710 **Table 7.** Effect of FOS on intestinal bacteria in the cecal digesta of weaned pigs
 711 challenged with enterotoxigenic *Escherichia coli*

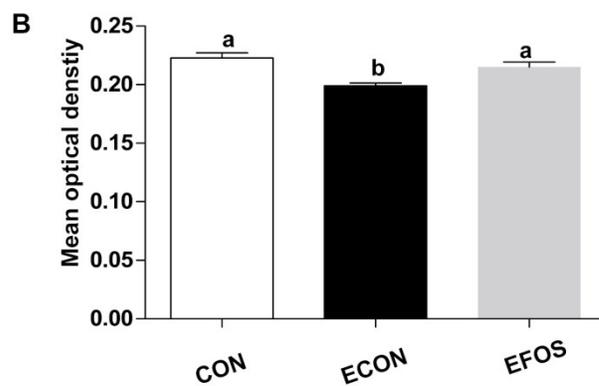
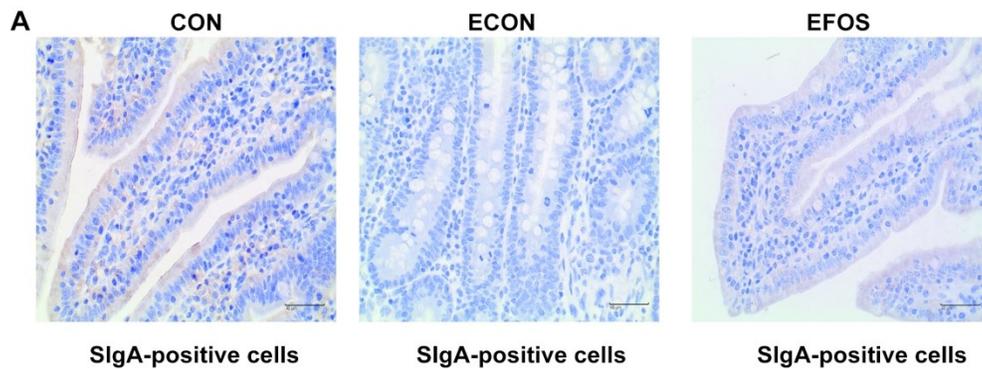
| Items | Treatments | | | <i>P</i> -Value |
|--|--------------------------|---------------------------|--------------------------|-----------------|
| | CON | ECON | EFOS | |
| Total bacteria 1g (copies/g) | 10.43 ± 0.13 | 10.21 ± 0.08 | 10.14 ± 0.19 | 0.37 |
| <i>Escherichia coli</i> 1g (copies/g) | 8.62 ± 0.05 ^b | 10.17 ± 0.23 ^a | 8.68 ± 0.26 ^b | 0.01 |
| <i>Bifidobacterium</i> 1g (copies/g) | 6.86 ± 0.47 ^a | 5.24 ± 0.42 ^b | 6.54 ± 0.39 ^a | 0.046 |
| <i>Bacillus</i> 1g (copies/g) | 8.57 ± 0.17 ^a | 8.03 ± 0.06 ^b | 8.59 ± 0.10 ^a | 0.01 |
| <i>Lactobacillus</i> 1g (copies/g) | 5.21 ± 0.37 | 4.57 ± 0.23 | 5.03 ± 0.12 | 0.17 |

712 ¹ Values are means ± SEM, (n = 6), non-challenged pigs (CON, fed with basal diet), ETEC-challenged
 713 pigs (ECON, fed with basal diet), FOS and ETEC-treated pigs (EFOS, fed with basal diet containing
 714 2.5 g/kg FOS challenged by ETEC). ² ^a, ^b, ^c mean values within a row with unlike superscript letters
 715 were significantly different (*P* < 0.05).

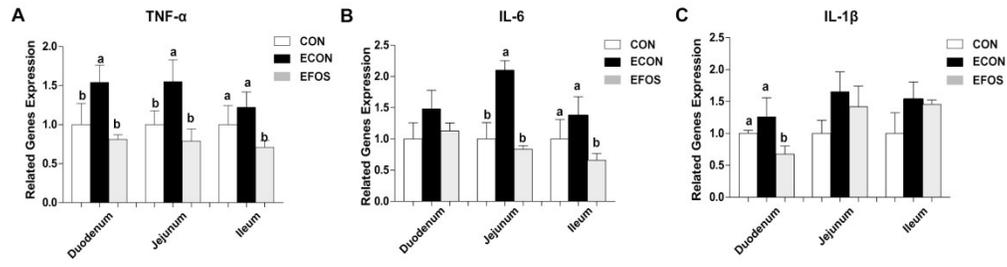




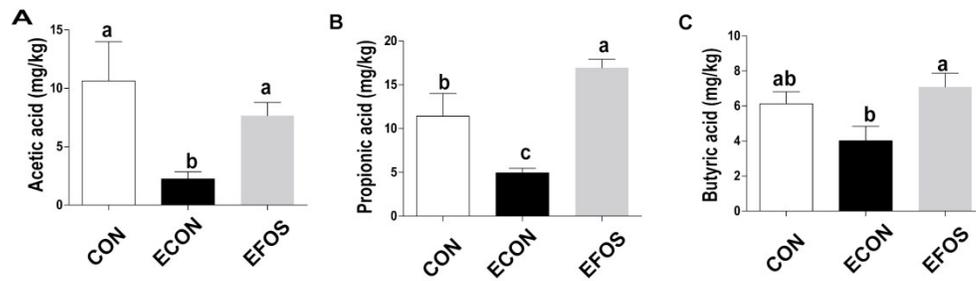
95x65mm (300 x 300 DPI)



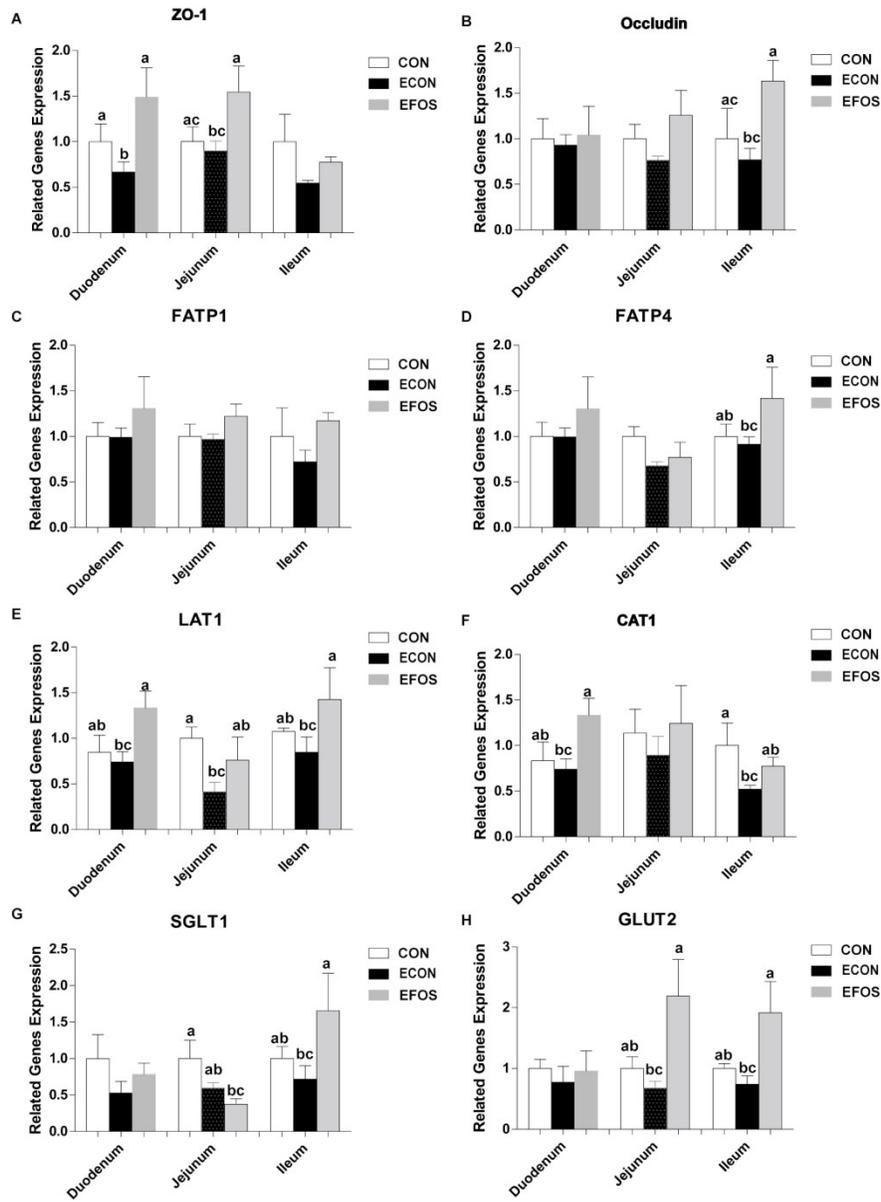
177x141mm (300 x 300 DPI)



259x70mm (300 x 300 DPI)



204x60mm (300 x 300 DPI)



245x327mm (200 x 200 DPI)