## RESEARCH





Arula-7 powder improves diarrhea and intestinal epithelial tight junction function associated with its regulation of intestinal flora in calves infected with pathogenic *Escherichia coli* O<sub>1</sub>

Hao Chen<sup>1†</sup>, Zhifeng Jia<sup>2,3†</sup>, Meiling He<sup>1†</sup>, Aorigele Chen<sup>1</sup>, Xin Zhang<sup>4</sup>, Jin Xu<sup>5</sup> and Chunjie Wang<sup>2\*</sup>

### Abstract

**Background** The effects of Arula-7 powder (ASP) on diarrhea and intestinal barrier function associated with its regulation of intestinal microflora in calves infected with pathogenic *Escherichia coli*  $O_1$  (*E. coli*  $O_1$ ) were studied.

**Method** Twenty Holstein calves were randomly divided into four treatment groups: normal control (NC), model control (MC), 0.5 mg/kg ciprofloxacin (CIP) and 2.50 g/kg ASP groups.

**Results** ASP inhibited the relative abundance of *Proteobacteria*, *Selenomonadales*, and *Enterobacteriales*, and increased the relative abundance of *Lactobacillus*, *Faecalibacterium*, and *Alloprevotella*. Moreover, we demonstrated for the first time that the ASP and CIP promoted weight gain, reduced the diarrhea rate (P < 0.05), and enhanced antioxidant capacity (P < 0.05) due to the increase in average daily gain (ADG), total protein (TP), and albumin (ALB). In addition, ASP and CIP increased the expression of Zunola occludens-1 (ZO-1), Occludin, and Claudin-1 in the ileum (P < 0.05), and improved immunity due to increase levels of interleukin-2 (IL-2), interleukin-4 (IL-4), interferon- $\gamma$  (IFN- $\gamma$ ), immunoglobulin A (IgA), and immunoglobulin G (IgG) in the serum, strengthened CD4<sup>+</sup>T levels in the ileal mucosa and reducing CD8<sup>+</sup>T and CD11c<sup>+</sup>T (P < 0.05).

**Conclusion** Hence, The intestinal microbiota environment formed by early intervention of ASP powder has a protective effect on the intestinal mucosal function of calves infected with pathogenic *E. coli*.

Keywords Arula-7 powder, Pathogenic Escherichia coli O1, Diarrhea, Intestinal flora, Intestinal barrier function, Calf

<sup>†</sup>Hao Chen, Zhifeng Jia, and Meiling He contributed equally to this work.

\*Correspondence: Chunjie Wang chunjiewang200@sohu.com Full list of author information is available at the end of the article



© The Author(s) 2023. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativeccommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

Content courtesy of Springer Nature, terms of use apply. Rights reserved.

### Introduction

Pathogenic Escherichia coli (E. coli) harms humans and livestock health and causes economic losses to the breeding industry [1]. Pathogenic E. coli, the clinical symptom of diarrhea, is very harmful to the health of infants (diarrheal disease leads to 15% of deaths in children under 5 years of age) and newborn poultry, and antibiotics are the first choice for clinical prevention [2]. Currently, many toxicities and side effects (such as destruction of the intestinal flora) caused by the abuse of antibiotics in the livestock industry has led to the refusal of people to consume the foods or products from livestock treated with antibiotics [3]. Therefore, to reduce and prohibit the use of antibiotics, researchers have begun to work on the research and development of alternatives to traditional antibiotics [4]. In this process, researchers have found that herbal medicines have antibacterial, and anti-inflammatory, properties, improve body immunity, and regulate intestinal flora, and herbal medicines maintain the dynamic balance of the internal environment to achieve disease prevention and treatment [5]. It has already been proven that in the process of infection with pathogenic *E*. *coli* O1, animals suffer from atrophy of villi in the small intestine and diarrhea, which affect the weight gain [6]. Some studies have shown that the two important effects of infection with pathogenic E. coli are the change in animal intestinal structure and imbalance of intestinal flora [7]. In fact, pathogenic *E. coli* leads to impaired intestinal epithelial barrier function and induces the destruction of calf gut microbiota balance. Once the microbiota balance is broken, potential pathogens in the intestine invade and colonize the gut, causing various diseases, especially inflammation and diarrhea [8]. Defects in microbial and immune function in the gut of newborn calves, and pathogenic microbes colonize the intestine to increase the risk of diarrhea at this stage [9]. In dairy farming, the pathogenic microbial-induced diarrhea rate reaches  $60 \sim 70\%$ , the mortality rate is more than 50%, and sick calves also emit a large amount of pathogenic bacteria to the outside world [10]. In addition to gut maturation, the calf has to adapt to the new environment, which is loaded with pathogenic microbes that may cause the calf to have diarrhea and die [11, 12].

Mongolian medicine plants, including seeds, berries, roots, stems, leaves, bark, and/or flowers from one or more plants, have therapeutic or other human health benefits. In some traditions, the scope of Mongolian medicine has been extended to cover materials of fungal, mineral and animal origin [13]. Herb compound intervention provides a feasible and practical method for mitigating intestinal inflammation and regulating animal intestinal flora [14]. There is an overwhelming consensus based on countless evidence that the herbal compound has the beneficial effects of improving the weight gain, enhancing immunity of livestock, protecting the intestinal mucosal barrier, and reducing disease [15]. Additionally, the favorable effects of curcumin on the animal health include the production of metabolites that inhibit the colonization of pathogenic microorganisms, which regulate the immunity of the animal's body, alleviate intestinal inflammation, and improve the intestinal barrier [16, 17]. Arula-7 powder (ASP), a traditional Mongolian medicine formula chronicled in the Mongolian medicine division, is widely used to improve intestinal diseases, such as diarrhea. Because ASP has long been used for the prevention and treatment of human diseases, research on its use by veterinarians is in a low-level development state.

Supplementation with short-chain fatty acids produced by specific bacteria significantly increased the production of Treg cells and enhanced the antibacterial activity of macrophages to prevent colitis [18]. Functionally, there is overlap in Treg inhibitory functions, and a subpopulation of T cells co-expressing CD8 and CD11c that represent an important category of adaptive immunomodulators with the ability to inhibit and exert effects in vivo was recently reported [19]. Numerous studies have confirmed that CD4+and CD8+T cells accelerate the activation and proliferation of T cells and produce various cytokines [20]. It is generally believed that inflammation plays a key role in interleukin (IL) and immunoglobulin (Ig) production to generate body immunity [21]. In recent years, there have been many studies on CD4+T lymphocyte subsets and their representative factors in diarrheal diseases, but few studies have been conducted on bacterial diarrhea. Therefore, this study will evaluate the effects of ASP on the status of CD4+T lymphocytes and their representative factors in mammals.

It has long been known that infection with pathogenic *E. coli* can damage the structure and function of the gut of the gut microbiota of an animal. However, there are few reports on the effect of herbal powder on intestinal epithelial tight junction function and intestinal flora, and the beneficial effect of improvement of diarrhea in calves infected with pathogenic *E. coli*  $O_1$  has rarely been reported. Therefore, in this study, ASP was used to interfere with the effective colonization of early calf gut microbiota, and the effective colonization of pathogenic microbes in the calf intestine was controlled from the source, thus providing suitable colonization conditions for probiotics in the calf.

The purpose of this study was examine the effects of Arula-7 powder on the diarrhea, immunity, intestinal epithelial tight junction function and intestinal flora of calves infected with pathogenic *E. coli* O<sub>1</sub>, thus providing a theoretical basis and experimental evidence for

understanding the biological effects of Arula-7 powder and its potential for regulating intestinal flora.

### Materials and methods

### Setting of groups and drug dose

All experiment animals were approved and performed in accordance with the guidelines of Institutional Animal Care and Use Committee of Inner Mongolia Agricultural University (IMAUG00001110). This study used Holstein bull calves (n=8) and female calves (n=12) reared at HeTan Crassland and Cattle 301 Ranch of the Shanxi, China.

The calves used in this study received 4 L of whole milk every day from the first day to the seventh day of the trial. The calves used in the study had no records of respiratory or enteric diseases or use of treatments for any disease. Twenty healthy Holstein calves were randomly divided into the natural control group (NC; bull calves = 2; female calves=3); model group (MC; bull calves=2; female calves=3); Ciprofloxacin (CIP; bull calves=2, female calves=3; CIP) and Arula-7 powder group (ASP; bull calves = 2; female calves = 3). Calves in the CIP and ASP groups received the following treatment twice per day for 4 days: 0.5 mg/kg Ciprofloxacin and compound Mongolian medicine 2.50 g/kg Arula-7 powder at a dosage. The NC group did not receive E. coli O1 or medicinal treatment, while the calves in all other groups received oral *E. coli* O1  $(2.50 \times 10^{11} \text{ CFU/mL}, 100 \text{ mL})$  on day 5, the E. coli O1 strain was provided by the Cow Production Laboratory, College of Animal Science, Inner Mongolia Agriculture University. And the calves in the CIP and ASP groups received the medicine treatment until day 7. At the end of the experiment, the calves were humanely euthanized using a captive bolt gun stunning method and sampling was completed within 30 min after euthanasia. According to the color and texture of the calf's feces, a score of 3 or more is determined as diarrhea based on the fecal scoring standard [22].

### Preparation of ASP

All seven medical plants were purchased from Hohhot Tianli Co., Ltd. (Hohhot, China); the detailed composition of ASP is listed in Table 1. The ASP was crushed, and then the powder was filtered through 200 mesh gauze.

### DNA isolation and 16S rRNA IonS5<sup>TM</sup>XL sequencing

For microbiota analysis, fresh cecal and colon contents (n=5) were collected at days 7 and RNA extraction (1.0 ng/mL) was performed. The extracted total RNA was amplified using CTAB/SDS method and sequencing (the RNA was stored at -78.5 °C in dry ice and performed at the Novogene Bioinformatics Technology Co., Ltd. Briefly). The 16S rRNA V34 341F 5'-CCT

Table 1	The comp	osition	of Arula-7	powder
---------	----------	---------	------------	--------

Chinese name	Latin name	Ratio
He Zi	Terminalia chebula Retz	3
Qian Cao	Rubia wallichiana Decne	2
Cao Wu Ye	Folium aconiti kusnezoffii	2
Gan Song	Nardostachys chinensis Batal	2
Ru Xiang	Boswellia carteri	2
Qu Mai	Dianthus superbus L	1
Fan Bai Cao	Potentilla discolor Bge	2

AYGGGRBGCASCAG-3 and 805R 5'-GGACTA CNNGGGTATCTAAT-3' amplicon was amplified by PCR and sequenced in the IonS5<sup>TM</sup>XL platform using the 2×250 bp paired-end protocol [23]. PCR products were purified using the QIA quick Gel Extraction Kit (QIAGEN, Dusseldorf, Germany) [24]. Single-end was used to evaluate the library quality and subsequently sequenced on an IonS5TMXL platform [25]. The 97% similarity sequence was analyzed by Alpha diversity including rarefaction analysis [26]. In brief, the OTU were influenced by IonS5TMXL [27] for the following analysis: alpha diversity, including observed species and Chao1 indices, number of observed OTUs and Shannon and Simpson diversity based on unweighted Uni-Frac distance of IonS5TMXL sequencing depth [28].

### Analysis of intestinal morphology

After a 7-day trial, all the jejunal and ileal segments of calves were stored at 10% formalin (Bkmam Biotechnology) and embedded in paraffin (Biosharp, BL957A) for histological and immunofluorescence analysis [29]. The paraffin-embedded jejunal and ileal tissue samples were sliced, stained by hematoxylin and eosin (H&E) methods, and then observed under an optical microscope.

### **Biochemical analysis**

After a 7-day trial, 30 min before euthanasia, Blood samples collected from the venous vein were centrifuged at  $3000 \times g$  for 15 min at 4 °C, and the clear supernatant was collected, dispensed into 1.5 mL centrifugal tubes and stored at – 80 °C for biochemical analysis. The enzyme-linked immunosorbent assay (ELISA, Bethesda, MD, USA) was used to quantitatively detect the level of SIgA, IgG, IgA, IFN- $\gamma$ , TNF- $\alpha$ , IL-6, IL-2, and IL-4 in calf serum. Strictly follow the instructions of the kit for each step, and conduct a preliminary study for each experiment.

### Analysis of serum biochemical indices

After a 7-day trial, the levels of serum urea nitrogen (BUN), serum alanine aminotransferase (ALT); aspartate aminotransferase (AST), alkaline phosphatase (ALP), total protein (TP), albumin (ALB), glucose (GLU), and total cholesterol (TC) were analyzed by BS-180VET (Mindray BS-2600 Vet, China), according to the manufacturer's instructions (Bethesda, MD, USA).

### Intestinal cytokines determined by ELISA

After a 7-day trial, after grinding in liquid nitrogen, total protein was extracted from ileal mucosa samples with lysis buffer (BaoXin, Hohhot, China), and then clarified by centrifugation  $(3000 \times g \text{ for } 15 \text{ min at } 4 \text{ °C})$ . The levels of CD4<sup>+</sup>T, CD8<sup>+</sup>T, CD11c<sup>+</sup>, ZO-1, Occludin, and Claudin-1 were measured using ELISA kit (Bethesda, MD, USA).

### Immunofluorescence and confocal microscopy

At the end of the experiment, the calf jugular vein was bled to death and the ileal was dissected for immunofluorescence experiments. The paraffin sections of the ileal were baked in an oven at 60 °C for 1 h, and then deodorized with xylene I, xylene II and xylene III for 5 min after soaking in 100%, 95%, and 80% ethanol for 5 min. The slides were incubated in biotin-conjugated primary antibody CD4<sup>+</sup>T (1:800 dilution, GeneTex), CD8<sup>+</sup>T (1:300 dilution, Bioss), and CD11c<sup>+</sup>T (1:500 dilution, Bioss), Occludin (1:500 dilution, Proteintech), ZO-1 (1:400 dilution, Proteintech) and Claudin-1 (1:300 dilution, Proteintech) overnight at 4 °C, and secondary antibody (1:200 dilution, Donkey anti Rat 594, Jackson) at room temperature for 1 h. The following steps are described by Xiong and Jin et al. [30, 31]. Finally, fluorescence was visualized using Olympus CK40 laser scanning confocal microscope.

### Statistical analysis

All the data from experiment were analyzed by using the one-way ANOVA. All the results obtained from replicate experiments are showed as means  $\pm$  s.e.m where "*n*" represents the number of animals/samples, and other figure results are presented with means  $\pm$  SEM, and histogram using GraphPad Software (La Jolla, CA, USA, www.graphpad.com) [32]. Significance was set at *P* < 0.05.

### Results

# Effects of treatments with ASP and CIP on the intestinal microbiome alpha diversity

Table 2 shows the results of the Chao1 index and Shannon index in the contents of cecum and colon.

 Table 2
 Effect of ASP on intestinal flora alpha diversity in diarrhea calves

ltems	NC	МС	0.50 mg/kg CIP	2.50 g/kg ASP
Cecal conten	t			
Chao1	172.35 <sup>a</sup>	128.42 <sup>c</sup>	138.05 <sup>bc</sup>	160.30 <sup>ab</sup>
Shannon	4.00 <sup>a</sup>	3.29 <sup>c</sup>	3.61 <sup>b</sup>	2.54 <sup>d</sup>
Colonic conte	ent			
Chao1	223.34 <sup>a</sup>	141.61 <sup>b</sup>	193.50 <sup>a</sup>	132.30 <sup>b</sup>
Shannon	3.67 <sup>a</sup>	3.40 <sup>a</sup>	3.87 <sup>a</sup>	3.47 <sup>a</sup>
Rectal feces				
Chao1	136.18 <sup>bc</sup>	211.61 <sup>ab</sup>	115.63 <sup>c</sup>	231.52ª
Shannon	3.71 <sup>ab</sup>	2.83 <sup>b</sup>	2.84 <sup>b</sup>	4.0 <sup>a</sup>

 $^{\rm a,b,c}$  Means bearing different superscripts in the same row differ significantly  $(P\,{<}\,0.05)$ 

Compared with MC group, Chao1 index of the cecal contents samples in the 2.50 g/kg ASP were significantly higher (P < 0.05). Compared with NC group, the Chao1 index of the cecal and colonic contents samples in MC group decreased apparently (P < 0.05), and the Shannon index of the cecal contents in MC group decreased apparently (P < 0.05). Compared with the NC group, the Shannon index of the of the cecal contents samples in 0.5 mg/kg CIP group increased apparently (P < 0.05). Chao1 index of the samples in the 0.5 mg/kg CIP group was significantly higher than those in the MC group (P < 0.05). Compared with NC and CIP groups, Chao1 index of the rectal feces samples in the ASP group were significantly higher (P < 0.05). Compared with the MC group, the Shannon index of the of the rectal feces in ASP group increased apparently (P < 0.05).

# Overall structure regulation of intestinal flora of calf diarrhea caused by E. coli $O_1$ after ASP treatment

Histograms illustrating the relative abundances of the three different taxonomic levels of calf cecum and colon contents is shown in Fig. 1 (Phylum, Order, and genus). With the different treatments, we observed a profound change in intestinal flora at the phylum, order and genus levels. In the cecal contents, compared with MC group, 2.50 g/kg ASP increased the relative abundance of Firmicutes and Bacteroidetes, and decreased Proteobacteria in phylum, but 0.50 mg/kg CIP increased the relative abundance of Bacteroidetes, and decreased Firmicutes and Proteobacteria (Fig. 1A). Compared with MC group, the relative abundance of Clostridiales and Bacteroidales was increased in the ASP group (P < 0.05). The relative abundance of Faecalibacterium, Alloprevotella and Bacteroides was significantly different among the ASP and CIP groups (P < 0.05). Compared with NC group, the



**Fig. 1** Effect of ASP on the composition of cecal and colonic contents microbiota in calves infected with pathogenic *E. coli*  $O_1$ . Graph  $\mathbf{A} \sim \mathbf{C}$  represents the OTUs at different taxonomical levels in the contents of the cecal: phylum, Order, and Genus. Graph  $\mathbf{D} \sim \mathbf{F}$  represents the OTUs at different taxonomical levels in the colonic: phylum, Order, and Genus. Red Latin indicates a significant increase (*P* < 0.05), and blue indicates a significant decrease (*P* < 0.05)

*Proteobacteria* and *Clostridiales* abundance of calf in cecal content was the higher in MC group at day 7.

The relative abundance of *Firmicutes* and *Proteobacteria* decreased in ASP and CIP groups at days 7 in colonic content if compared with MC group (Fig. 1D, P < 0.05). At the order level, The relative abundance of *Clostridiales* increased in MC group at days 7 in colonic content if compared with NC group. Compared with MC group, ASP and CIP groups decreased the relative abundance of *Clostridiales* and *Enterobacteriales*, but increased *Bacteroidales* (Fig. 1E, P < 0.05).

# Arula-7 powder alters the intestinal flora of calves infected with pathogenic E. coli $O_1$

To assess how ASP administration affected the calf rectal gut microbial composition, the phylum and genuslevel bacterial relative abundance of the control and treatment groups at days 7 was compared (Fig. 2). At the phylum level, compared with MC group, the Firmicutes, Actinobacteria, and ratio of Firmicutes/Bacteroidetes was reduced in ASP group from day 7 after ceased ASP application, and the significantly higher abundance in Firmicutes, Actinobacteria, and ratio of Firmicutes/Bacteroidetes (P < 0.05) in MC group was detected, and the markedly more *Faecalibacterium* (P < 0.01) was detected after ceased CIP application. At day 7, there is significantly higher *Fusobacterium* and *Bacteroides* (P < 0.05).

# Effects of different doses of Arula-7 powder on the content of E. coli in the intestine of calves

We quantified these two important bacterial species in the cecal, colonic, and rectal contents digesta by using quantitative real-time PCR analysis. The results are shown in Table 3. The ASP, CIP, and NC groups had a similar relative abundance of *E. coli* in cecal and rectal contents. Moreover, the number of *E. coli* in colonic contents from the ASP and MC groups was significantly higher while the number of *E. coli* was significantly lower than those in the other groups (P < 0.05), and the number of *E. coli* in the MC group was significantly higher than those in the other groups (P < 0.05).



**Fig. 2** Effect of Arula-7 powder on the composition of rectal microbiota in calves infected with pathogenic *E. coli*  $O_1$ . **A** Graph represents the OTUs at different taxonomical levels: order (n=5). **B** Graph represents the OTUs at different taxonomical levels: order (n=5). **C** The change in the relative abundance of phylum Firmicutes, Bacteroidetes, Fusobacteria, and Actinobacteria. **D** The ratio of Firmicutes/Bacteroidetes was significantly low in 2.50 g/kg ASP group. **E** The change in the relative abundance of genus *Faecalibacterium*, *Fusobacterium*, *Alloprevotella*, *Megamonas*, and *Bacteroides*. Metastats analysis was applied to identify the significantly differentially abundant phylum and genera among groups. Different letters above the bars denotes significantly differentially abundant genera among groups. (P < 0.05). n=5

Table	3 Q	uantitativ	e real-time	e PCR	analysis	of	total	Escheric	hia
<i>coli</i> in	ceca	l, colonic,	and rectal	conte	ents [lgCF	U/g	g wet	weight)	]

ltems	NC	МС	0.5 mg/kg CIP	2.50 g/kg ASP
Cecal content	5.73 <sup>c</sup>	12.46 <sup>a</sup>	7.06 <sup>c</sup>	8.48 <sup>b</sup>
Colonic content	6.44 <sup>b</sup>	12.47 <sup>a</sup>	6.80 <sup>b</sup>	10.98 <sup>a</sup>
Rectal feces	5.24 <sup>c</sup>	9.67 <sup>a</sup>	5.42 <sup>bc</sup>	6.82 <sup>b</sup>

 $^{\rm a,b,c}$  Means bearing different superscripts in the same row differ significantly  $(P\!<\!0.05)$ 

Table 4         Effects of ASP on weight gain and diarrhea rate of ca
---

ltems	NC	мс	0.5 mg/kg CIP	2.50 g/kg ASP	
Initial weight, kg	$29.75 \pm 0.73^{a}$	$29.02 \pm 1.04^{a}$	$30.85 \pm 1.24^{a}$	$29.78 \pm 1.04^{a}$	
Final weight, kg	$33.37 \pm 0.83^{a}$	$31.90 \pm 1.25^{b}$	$33.89 \pm 0.86^{a}$	$34.15 \pm 1.37^{a}$	
ADG, kg/day	$0.56 \pm 0.08^{b}$	$0.25 \pm 0.04^{\circ}$	$0.46 \pm 0.11^{b}$	$0.71 \pm 0.05^{a}$	
Diarrhea ratio, %	0.00 <sup>c</sup>	80.00 <sup>a</sup>	20.00 <sup>b</sup>	20.00 <sup>b</sup>	

 $^{\rm a,b,c}$  Means bearing different superscripts in the same row differ significantly  $(P\!<\!0.05)$ 

# Effects of ASP on weight gain, diarrhea rate, and blood biochemical indexes of calves with diarrhea

After a 7-day trial, the growth performance and diarrhea rate of calves with *E. coli* diarrhea are shown in Table 4. Compared with the NC group, MC group significantly decreased final weight (P<0.05) and decreased average daily gain (ADG, P<0.05), but ASP increased ADG (P<0.05), CIP group had no effect on ADG (P>0.05). ASP and CIP groups significantly decreased the diarrhea rate compared with that in MC group (P<0.05). In addition, the ASP and CIP groups restored the calf body weight to the NC group level.

# Effect of ASP on blood biochemical indexes of calves with Escherichia coli diarrhea

The blood indexes of calves with *E. coli* diarrhea are shown in Fig. 3. After a 7-day trial, compared with the NC group, MC group had no effect on Glu (Fig. 3F) and AST (Fig. 3B) in serum. Moreover, compared



**Fig. 3** Effect of ASP on blood biochemical indexes of calves with *E. coli* diarrhea. **A** BUN, blood urea nitrogen. **B** AST, aspartate aminotransferase. **C** TP, serum total protein. **D** ALB, albumin. **E** ALP, alkaline phosphatase. **F** Glu, glutamic. n = 5. In the same histogram, the values with the same letter superscripts mean no significant difference (P > 0.05), while with different letter superscripts mean significant difference (P < 0.05)

with the MC group, ASP, and CIP groups significantly increased TP (Fig. 3C) and ALB (Fig. 3D) and CIP decreased the ALP (Fig. 3E) content (P < 0.05), and the level of TP and Glu in ASP group were significantly higher than NC group.

## Effect of ASP on the intestinal morphology of calves with Escherichia coli diarrhea

As shown in Fig. 4, compared with the NC group, the intestinal villi in the jejunum and ileum of the MC group with *E. coli*  $O_1$  were severely shed, especially the ileum



Fig. 4 Effects of ASP on histopathological changes of jejunum and ileum with H&E staining (original magnification of 100×). A Stroke-physiological saline solution + NC group, B *E. coli* O<sub>1</sub> + MC group, C *E. coli* O<sub>1</sub> + 0.5 mg/kg CIP group, D *E. coli* O<sub>1</sub> + 2.50 g/kg ASP group

injury was the most serious. while the villi in the ASP group were intact and tight.

## Effect of ASP on tight junction protein and lymphocyte factor in the ileum of calves with Escherichia coli diarrhea

To investigate the effects of ASP treatment on calf tight junction proteins and lymphocytes, the ileal ZO-1, Occludin, and Claudin-1, as well as ileal CD4<sup>+</sup>T, CD8<sup>+</sup>T, and CD11c<sup>+</sup>T, were monitored (Table 5). Compared with the NC and MC groups, the level of ileal ZO-1, Occludin, and Claudin-1 were enhanced in ASP group at day 7 (P<0.05). Compared with the NC group, pathogenic *E. coli* O<sub>1</sub> did result in a significant reduction in the ileal CD4<sup>+</sup>T levels for MC group. However, the level of ileal CD4<sup>+</sup>T was increased in ASP and CIP groups (P<0.05). Compared with MC group, CIP and ASP group significantly reduced the content of CD8<sup>+</sup>T and CD11c<sup>+</sup>T (P<0.05), and the levels of CD4<sup>+</sup>T, CD8<sup>+</sup>T, and CD11c<sup>+</sup>T in the ASP and CIP groups were restored to the NC group level.

# Effects of ASP on the expression of intestinal mucosa tight junction protein

There was a significant difference in tight junctionrelated components among the 4 groups at the ending of day 7 (Fig. 5). Pathogenic *E. coli*  $O_1$  reduced the expression of ZO-1 (Fig. 5A), Occludin (Fig. 5B), and Claudin-1 (Fig. 5C) in the ileum. The expressions of ZO-1 (Fig. 5D), Occludin (Fig. 5E), and Claudin-1 (Fig. 5F) in the ASP and CIP groups increased compared with the MC group (P<0.05). ASP and CIP administration restored ZO-1, Occludin, and Claudin-1 expression in the ileum. The distributions of ZO-1, Occludin, and Claudin-1 were also estimated by immunofluorescence staining. Compared with the MC group, which showed clear and uniform positive staining of the ZO-1 (Fig. 5D), Occludin (Fig. 5E), and Claudin-1 (Fig. 5F) in the intestinal epithelium if compared with MC and NC groups at 7-day (P < 0.05), the MC group showed a decrease staining of both ZO-1, Occludin, and Claudin-1 in the ileum if compared with NC group (P < 0.05).

# Effect of ASP administration improved intestinal CD4<sup>+</sup>T lymphocytes and their representative cytokines

The distributions of CD4<sup>+</sup>T, CD8<sup>+</sup>T and CD11c<sup>+</sup>T were also estimated by immunofluorescence staining (Fig. 6). Oral E. coli O1 significantly reduced the expression of  $CD4^+T$  in ileum (Fig. 6A), while the expression of  $CD8^+T$ (Fig. 6B) and CD11c<sup>+</sup>T (Fig. 6C) significantly increased (P < 0.05). Compared with NC calves, the MC calves showed significantly more CD8<sup>+</sup>T (P<0.05; Fig. 6E) and CD11c<sup>+</sup>T (P < 0.05; Fig. 6F) deposition in the ileum epithelium, whereas a dramatic reduction in the ileum accumulation was noted with ASP and CIP administration. Compared with the MC group, ASP group significantly increased CD4<sup>+</sup>/CD8<sup>+</sup>T (P<0.05; Fig. 6G) and increased  $CD4^+T$  (*P*<0.05; Fig. 6D). In addition, ASP significantly increased the CD4<sup>+</sup>T and decreased CD11c<sup>+</sup>T (P < 0.05) but had no effect on  $CD8^+T$  (P>0.05) compared with that in NC group (P < 0.05). The CD4<sup>+</sup>T in the ASP group was significantly higher than those in the CIP group (P < 0.05). Our results showed that ASP and CIP administration reversed CD4<sup>+</sup>T expression in the ileum.

### ASP improved calf immune responses

Firstly, we investigated the effects of the ASP treatment on animal immunity, the serum IgA, IgG, SIgA, IL-4, IL-2, IFN- $\gamma$ , IL-6, and TNF- $\alpha$ , were monitored (Fig. 7). ASP and CIP significantly increased IgA (Fig. 7A), IgG (Fig. 7B), IL-4 (Fig. 7D), IL-2 (Fig. 7E), and IFN- $\gamma$ (Fig. 7F), and decreased IL-6 and TNF- $\alpha$  compared with that in MC group (P<0.05). For the MC group, significantly reduced the levels of serum IL-6 (Fig. 7G) and TNF- $\alpha$  (Fig. 7H) comparing with NC group.

### Discussion

Intestinal development is a critical stage of growth for infants and newborn calves, and infection with pathogenic *E. coli* during intestinal development can cause intestinal barrier dysfunction, gut microbiota disruption, and inflammation, accompanied by a decline in the absorption capacity of the small intestine, and eventually leading to an increase in the incidence of

Table 5 Effect of ASP on tight junction protein and lymphocyte factor in calf ileum

ltems	NC	МС	0.5 mg/kg CIP	2.50 g/kg ASP
ZO-1, ng/0.2 g	130.36±6.50 <sup>b</sup>	110.0±6.83 <sup>c</sup>	133.81±10.42 <sup>b</sup>	156.90±11.97 <sup>a</sup>
Occludin, ng/0.2 g	707.23±16.37 <sup>b</sup>	$636.56 \pm 20.91^{b}$	$650.98 \pm 30.90^{b}$	$1062.52 \pm 97.94^{a}$
Claudin-1, pg/0.2 g	$220.28 \pm 2.03^{b}$	$214.32 \pm 5.28^{b}$	211.51±6.34 <sup>b</sup>	$232.04 \pm 3.17^{a}$
CD4+, ng/0.2 g	$72.39 \pm 4.35^{a}$	$59.03 \pm 3.80^{b}$	$70.32 \pm 1.04^{a}$	$73.39 \pm 4.80^{a}$
CD8 <sup>+</sup> , ng/0.2 g	$28.50 \pm 2.84^{ab}$	$32.19 \pm 2.67^{a}$	$25.75 \pm 1.21^{b}$	25.20±3.11 <sup>b</sup>
CD11c <sup>+</sup> , ng/0.2 g	$45.42 \pm 0.54^{ab}$	43.61±1.79 <sup>ab</sup>	$47.50 \pm 1.17^{a}$	37.31±0.21 <sup>b</sup>

 $^{a,b,c}$  Means bearing different superscripts in the same row differ significantly (P<0.05)



**Fig. 5** Intestinal tight junction proteins in pathogenic *E. coli* O<sub>1</sub> induced diarrhea calf with ASP and CIP for 7 days (400 ×). Immunofluorescence staining of (**A**) ZO-1, (**B**) Occludin, and (**C**) Claudin-1 in the ileum. **D**, **E**, **F** Each field of ZO-1, Occludin, and Claudin-1 from every group was quantified by using ImageJ. Scale bar = 20 µm; red, ZO-1, Occludin, and Claudin-1; blue, 4,6-diamidino-2-phenylindole counterstaining of the nuclei. Control group was denoted as CG; NC, Normal control group; MC, *E. coli* group; CIP, Ciprofloxacin group; ASP, Arula-7 powder group. ZO-1, Occludin and Claudin-1

diarrhea [7]. Pathogenic *E. coli* infection is associated with impairment of cellular immunity and an increased prevalence of gastrointestinal infection in newborn calves [2]. Therefore, the appropriate addition of plantbased drugs in the early stage seems to be crucial for enhancing the gut health of animals. Gan et al. reported that some plant extracts (e.g., curcumin) were highly effective in protecting the intestinal barrier function of mice [17]. In this work, we found that ASP powder as a natural antibacterial, anti-inflammatory, and immune enhancer, improved calf growth performance, reduced diarrhea, strengthened immunity, ameliorated intestinal barrier dysfunction and inflammation, and blocked the discharge of pathogenic microbes. First, the calf was supplemented with Arula-7 powder in breast milk for 5 days after birth, then the calf was orally infected with pathogenic *E. coli*  $O_1$  and the drug was administered for another 2 days. After the 7-day trial, the incidence of diarrhea in the ASP group was low, which may be related to decreased colonization of pathogenic *E. coli* O1 [6]. Compared with the MC group, ASP increased the ADG, indicating that the weight was improved. In addition, ASP significantly reduced the rate of diarrhea in calves infected with pathogenic *E. coli*  $O_1$ . Compared with the NC group, in the MC group, the ADG and final weight were reduced due to the unbalanced intestinal flora. In the current study, the weight gain, immunity, and diarrhea rate of



**Fig. 6** Intestinal lymphocyte surface molecular proteins in pathogenic *E. coli*  $O_1$  induced diarrhea calf with ASP and CIP for 7 days (400×). Immunofluorescence staining of **A** CD4<sup>+</sup>, **B** CD8<sup>+</sup>, and **C** CD11c<sup>+</sup> in the ileum. **D**, **E**, **F** Each field of CD4<sup>+</sup>, CD8<sup>+</sup>, and CD11c<sup>+</sup> from every group was quantified by using Image J. **G** The ratio of CD4<sup>+</sup>/CD8<sup>+</sup>T was significantly high in ASP group. Scale bar = 20 µm; red, CD4<sup>+</sup>, CD8<sup>+</sup>, and CD11c<sup>+</sup>; blue, 4,6-diamidino-2-phenylindole counterstaining of the nuclei. Control group was denoted as CG; NC, Normal control group; MC, *E. coli* group; CIP, Ciprofloxacin group; ASP, Arula-7 powder group

animals were related to intestinal mucosa permeability, intestinal morphology, and intestinal flora balance [33]. Further analysis showed that milk supplementation with ASP significantly increased the contents of TP, ALB, and Glu in the serum, and reduced the content of BUN. These findings suggest that ASP supplementation affected the growth and production performance of calves. The effect of ASP on antioxidation indices is consistent with a prior study of diarrhea in rabbits [34].

In addition, the structure of the epithelial barrier was impaired, and the tight junctions of the intestinal epithelium were destroyed, thereby destroying the polarity of the intestinal epithelial cells and resulting in the substance spilling into the epithelial cell gap [35]. Protecting the ZO-1 protein in the intestine plays an important role in maintaining the tight junctions of intestinal epithelial cells [36]. Occludin, a transmembrane protein, is positively correlated with the structural integrity of the intestinal epithelium and intestinal epithelial barrier function [37]. After EPEC infection of T84 human intestinal epithelial cells, the close relationship between claudin-1, occludin and ZO-1 gradually decreased, their positions gradually changed, and the structure was damaged [38]. Huangqin decoction has been reported to significantly inhibit colitis and improve tissue damage [5]. In this study, ASP powder improved the intestinal morphology of calves infected with pathogenic *E. coli* O<sub>1</sub> and increased the expression of the tight junction proteins



Fig. 7 Effects of ASP treatment on calf immune markers. The levels of A serum IgA, B serum IgG, C serum SIgA, D serum IL-4, E serum IL-2, F serum IFN-γ, G serum IL-6, H serum TNF-α

ZO-1, Occludin, and Claudin-1. Currently, there is no relevant research on the effect of ASP powder on the intestinal morphology of calves infected with pathogenic *E. coli*  $O_1$ . Therefore, this study may be the first to report that ASP powder can significantly improve the intestinal tight junctions by increasing the expression of ZO-1, Occludin, and Claudin-1 in the intestine morphology. The ability of ASP powder to improve calf diarrhea by pathogenic *E. coli*  $O_1$  may be due to its effects of improving intestinal morphology and increasing tight junction proteins.

The intestine plays an important role in protecting the animal's body from bacteria and acting as an immune barrier [39]. To investigate ASP-induced immunity regulation in calves, our work analyzed changes in several T cell populations in the intestinal tissues,  $CD4^+T$ ,  $CD8^+T$ ,  $CD11c^+T$ , and several serum cytokines, IFN- $\gamma$ , IL-4, IL-6, IL-2, TNF- $\alpha$ , and IL-6, as well as serum IgG, IgA, and SIgA. An increase in CD11c<sup>+</sup>T and CD8<sup>+</sup>T cells on the intestinal mucosa has already been proven in animal models of virus infection [40]. The various functions exhibited by CD8+T are primarily determined by

changes in the microenvironment induced by inflammation [18]. Although high expression of CD8 + T reduced IFN-y levels, its mechanism was likely to occur in a cellcell contact-dependent fashion [41]. Compared with those in the MC group, the levels of 3 cytokines serum IL-2, IL-4 and IFN-y, and serum IgG, IgA, and SIgA, as well as tissue CD4<sup>+</sup>T of the ASP group had increased (p < 0.05). Thus, the increased CD8<sup>+</sup>T, CD11c<sup>+</sup>T cell populations, and the increased expression of IL-6 and TNF- $\alpha$  observed in the tissue of the MC group in the present study might be an indication of restricted immunity between the intestinal lumen and epithelium as a physiological response to diarrhea, but ASP opposes diarrhea. In the encephalitis models, the CD8<sup>+</sup>T inhibit the expansion of CD4<sup>+</sup>T cells via cell–cell contact [42]. Here, we observed that the concentration of CD4+T in the intestinal tissue of ASP-treated calves was significantly higher than that in the NC and MC groups. Immunoglobulin (Ig) has long been recognized as the first immune barrier against the invasion of pathogenic microorganisms at the mucosa [43]. It can prevent pathogenic microorganisms from adhering to the surface of intestinal epithelial cells and neutralize bacterial toxins [44]. IgG binds to gramnegative bacteria in the intestinal lumen to reduce the risks of bacterial translocation, intestinal damage and systemic infection [45]. Additionally, the levels of IL-6 and TNF- $\alpha$  were significantly reduced in the ASP and CIP groups, and SIgA was significantly higher in the ASP group than in the CIP group. Moreover, although the levels of intestinal tissue CD4<sup>+</sup>T and serum IL-2, IL-4, and IFN- $\gamma$ , and serum IgG, IgA, and SIgA in the MC group were lower than those in the NC group, their levels increased after ASP and CIP treatment, indicating that ASP and CIP could strengthen the immunity of calves. However, opposite observations were reported by Liu et al. [46].

Recent studies have shown that herbal medicines can directly lead to intestinal microflora shift and reduce inflammation, thereby improving the gut health of animals [15]. The gut microbiome has long been considered to be linked to immune system function and regulation [34]. The reduction of *Bacteroides* in the gut of obese humans and animals reduces the abundance of Firmi*cutes*, suggesting that these major phyla may play a role in obesity-related inflammation [47]. A lower ratio of *Firmicutes* to *Bacteroidetes* is a sign of intestinal health. Compared with the NC group, the Firmicutes to Bacteroidetes ratio in the cecal contents of calves in the MC group was increased, which is consistent with the high diarrhea rate in the MC group. We observed that ASP increased the abundance of Bacteroides and Firmicutes in the cecal contents of calves infected with pathogenic E. coli O1, reduced the abundance of Firmicutes and Proteobacteria in the cecal and colonic contents, and increased the abundance of Bacteroides in colonic contents. Intriguingly, any alteration in the physical or chemical colonic contents may affect the resident bacterial structure and potentially promote or inhibit the establishment of pathogenic bacteria [34]. In this study, ASP increased the relative abundance of the genera Faecalibacterium, Alloprevotella, and Bacteroides in the cecal and colonic contents and increased the relative abundance of the order Bacteroidales, while decreasing the abundance of Proteobacteria, Selenomonadales, Enterobacteriales, and *Clostridiales* in the cecal and colonic contents. The genera Faecalibacterium and Alloprevotella produce most of the amino acids that calves obtained through their intestines. These strains promote weight gain and reduce intestinal inflammation and diarrhea [48]. In addition, *Faecalibacterium* promotes the differentiation of T and B lymphocytes into plasma cells and increases the metabolism of T and B cells, thereby enhancing the production of host antibodies [49]. Therefore, ASP could improve the immune function of calves infected with pathogenic E. coli O<sub>1</sub> by increasing the relative abundance of *Faecalibacterium*. In addition, ASP may reduce the incidence of diarrhea caused by *E. coli*  $O_1$  by decreasing the abundance of *Proteobacteria*, *Selenomonadales*, and *Enterobacteriales*.

In the diarrhea of calves, a close link between intestinal flora and feces flora was observed, and the changes in the flora associated with diarrheal disease were similar between the two species, characterized by a decrease in Proteobacteria and Bacteroidetes. In the current study, ASP increased the relative abundance of Fusobacteria and Lactobacillus in the rectal feces of calves. Lactobacillus, a probiotic, promotes weight gain; therefore, ASP promotes the weight gain of calves by altering the rectal microbiota. In addition, a previous study showed that Lactobacillus is involved in regulating the immune system, preventing diarrhea, and improving animal health [50]. On the one hand, it is plausible that ASP could improve the immune function of calves infected with pathogenic *E. coli*  $O_1$  by increasing the relative abundance of Faecalibacterium. On the other hand, ASP may reduce the incidence of diarrhea caused by E. coli by decreasing the number of E. coli. Therefore, the restoration of intestinal ecosystem function can potentially be used as an effective treatment strategy for calves infected with pathogenic E. coli O1. This may be because ASP can reducing the occurrence of diarrhea by reducing the abundance of Proteobacteria and increasing the abundance of Faecalibacterium. This further illustrates that ASP completely blocks the release of pathogenic microorganisms in the calf intestine, protects the environment, and avoids infection of other animals.

### Conclusions

Overall, the results of this study suggest that milk supplementation with ASP powder promoted weight gain, ameliorated diarrhea and inflammation, and improved immunity in calves infected with pathogenic *E. coli*  $O_1$  in a 7-day trial. These results provide a basis for the further development of green products that maintain the functional integrity of the intestinal mechanical and biological barriers in calves infected with pathogenic *E. coli*  $O_1$ . These results provide a theoretical basis and experimental evidence for understanding the biological function of ASP powder and its potential for use in place of antibiotics.

### Acknowledgements

We thank Editage for their assistance in editing our article and for language revision. We thank X. Zhang and J. Xu for collecting the experimental samples for us.

### Authors' contributions

H. Chen, Z. F Jia and M. L He interpreted the results and wrote the manuscript. A. Chen and C. J Wang provided funding; Z. F Jia and C. J Wang designed the experiments and revised the manuscript. X. Zhang and J. Xu performed the experiments. All authors read and approved the final manuscript.

### Funding

This work was supported by the National Natural Science Foundation of China (No. 31760720), and they provided suggestions about the study design.

### Availability of data and materials

Raw 16S rRNA gene sequences for all samples used in this study have been deposited in the Sequence Read Archive (SRA) under project accession no. SUB6318504, SUB6321029, and SUB6321118.

### Declarations

### Ethics approval and consent to participate

The use of experimental newborn Holstein calf were approved by the College of Animal Science of Inner Mongolia Agricultural University of China IMAU (protocol 181010 for calf). Signed owner consent was obtained before enrollment of calves in the IMAU study. We obtained written informed consent to use the animals in our study from the owner(s) of the animals.

#### **Consent for publication**

Not applicable.

#### **Competing interests**

The authors declare no competing interests.

#### Author details

<sup>1</sup> College of Animal Science, Inner Mongolia Agricultural University, Hohhot 010018, People's Republic of China. <sup>2</sup>College of Veterinary Medicine, Inner Mongolia Agricultural University, Hohhot 010018, People's Republic of China. <sup>3</sup>Animal Disease Prevention and Control Center of Bazhou District, Bazhong, China. <sup>4</sup>College of Basic Medical, Inner Mongolia Medical University, Hohhot 010110, People's Republic of China. <sup>5</sup>Henan Houyi Bio-Engineering, Inc, He Nan, 451161 Zhengzhou, People's Republic of China.

### Received: 9 April 2022 Accepted: 7 July 2023 Published online: 05 August 2023

### References

- Mirhoseini A, Amani J, Nazarian S. Review on pathogenicity mechanism of enterotoxigenic *Escherichia coli* and vaccines against it. Microb Pathog. 2018;117:162–9.
- Baumgartner M, Bayer F, Pfrunder-Cardozo KR, Buckling A, Hall AR. Resident microbial communities inhibit growth and antibiotic-resistance evolution of *Escherichia coli* in human gut microbiome samples. PLoS Biol. 2020;18:e3000465.
- Liu LL, Wu W, Zhang JY, Lv P, Xu L, Yan YC. Progress of research on the toxicology of antibiotic pollution in aquatic organisms. Acta Ecol Sin. 2018;38(1):36–41.
- Li Q, Montalban-Lopez M, Kuipers OP. Increasing the antimicrobial activity of Nisin-based *Lantibiotics* against Gram-negative pathogens. Appl Environ Microbiol. 2018;84(12):e00052-e118.
- 5. Yang Y, Gang C, Qian Y, Ye J, Cai XT, Tsering P, et al. Gut microbiota drives the attenuation of dextran sulphate sodium-induced colitis by Huangqin decoction. Oncotarget. 2017;8(30):48863–74.
- Jia ZF, Chen ARGL, Bao FX, He ML, Gao SM, Zhang X, et al. Effect of nisin on microbiome-brain-gut axis neurochemicals by *Escherichia coli*induced diarrhea in mice. Microb Pathogenesis. 2018;119:65–71.
- Jia ZF, Wu AQM, He ML, Zhang L, Wang CJ, Chen ARGL. Metabolites of stable fly reduce diarrhea in mice by modulating the immune system, antioxidants, and composition of gut microbiota. Microb Pathog. 2019;134:103557.
- Li XQ, Zhu YH, Zhang HF, Yue Y, Cai ZX, Lu QP, et al. Risks associated with high-dose Lactobacillus rhamnosus in an Escherichia coli model of piglet Diarrhoea: intestinal microbiota and immune imbalances. PLoS ONE. 2012;7:e40666.
- Buffie CG, Bucci V, Stein RR, McKenney PT, Ling L, Gobourne A, et al. Precision microbiome reconstitution restores bile acid mediated resistance to Clostridium difficile. Nature. 2015;517:205–8.

- Rosa F, Busato S, Avaroma FC, Linville K, Trevisi E, Osorio JS, et al. Transcriptional changes detected in fecal RNA of neonatal dairy calves undergoing a mild diarrhea are associated with inflammatory biomarkers. PLoS ONE. 2018;13:e0191599.
- 11. Berchtold J. Treatment of calf diarrhea: intravenous fluid therapy. Vet Clin North Am Food Anim Pract. 2009;25(73–99):vi.
- 12. Culligan EP, Hill C, Sleator RD. Probiotics and gastrointestinal disease: successes, problems and future prospects. Gut Pathog. 2009;1:19.
- 13. WHO. General guidelines for methodologies on research and evaluation of traditional medicine. World Health Organization; 2000. p. 1–71.
- Zhang Y, Tang K, Deng Y, Chen R, Liang S, Xie H, et al. Effects of shenling baizhu powder herbal formula on intestinal microbiota in high-fat dietinduced NAFLD rats. Biomed Pharmacother. 2018;102:1025–36.
- Xu J, Chen HB, Li SL. Understanding the molecular mechanisms of the interplay between herbal medicines and gut microbiota. Med Res Rev. 2017;37:1140–85.
- Liu J, Jia ZF, Chen ARGL, Wang CJ. Effects of Mongolian veterinary drug Babu 7 active ingredient compound on growth and related immune factors in blood of calves with Escherichia coli diarrhea. Microb Pathog. 2021;17:51–5.
- 17. Gan Z, Wei W, Li Y, Wu J, Zhao Y, Zhang L, et al. Curcumin and resveratrol regulate intestinal Bacteria and alleviate intestinal inflammation in weaned piglets. Molecules. 2019;24:1220.
- Imhann F, Vich Vila A, Bonder MJ, Fu J, Gevers D, Visschedijk MC, et al. Interplay of host genetics and gut microbiota underlying the onset and clinical presentation of inflammatory bowel disease. Gut. 2018;67:108–19.
- Vinay DS, Kwon BS. CD11c<sup>+</sup>CD8<sup>+</sup>T cells: two-faced adaptive immune regulators. Cell Immunol. 2010;264:18–22.
- Jia ZF, Chen ARGL, Wang CJ, He ML, Xu J, Fu H, et al. Amelioration effects of Kaempferol on immune response following chronic intermittent coldstress. Res Vet Sci. 2019;125:390–6.
- Zhao J, Liu Y, Wu P, Jiang W, Li S, Tang L, et al. Effects of dietary isoleucine on the immune response, antioxidant status and gene expression in the head kidney of juvenile Jian carp (Cyprinus carpio var. Jian). Fish Shellfish Immunol. 2013;35:572–80.
- Hill TM, Aldrich JM, Schlotterbeck RL, et al. Amino acids, fatty acids, and fat sources for calf milk replacers. Professional Animal Scientist. 2007;23(4):401–8.
- 23. Caporaso JG, Lauber CL, Walters WA, Berg-Lyons D, Huntley J, Fierer N, et al. Ultra-high-throughput microbial community analysis on the Illumina HiSeq and MiSeq platforms. ISME J. 2012;6:1621–4.
- 24. Zeng B, Han S, Wang P, Wen B, Jian W, Guo W, et al. The bacterial communities associated with fecal types and body weight of rex rabbits. Sci Rep. 2015;5:9342.
- Hess M, Sczyrba A, Egan R, Kim TW, Chokhawala H, Schroth G, et al. Metagenomic discovery of biomass-degrading genes and genomes from cow rumen. Science. 2011;331:463–7.
- Caporaso JG, Kuczynski J, Stombaugh J, Bittinger K, Bushman FD, Costello EK, et al. QIIME allows analysis of high-throughput community sequencing data. Nat Methods. 2010;7:335–6.
- Kuczynski J, Stombaugh J, Walters W A, González A, Caporaso JG, Knight R. Using qiime to analyze 16s rRNA gene sequences from microbial communities. Curr Protoc Bioinformatics. 2011; Chapter 10: Unit 10 17.
- Wang S, Martins R, Sullivan MC, Friedman ES, Misic AM, El-Fahmawi A, et al. Diet-induced remission in chronic enteropathy is associated with altered microbial community structure and synthesis of secondary bile acids. Microbiome. 2019;7(1):126.
- Gao Y, Han F, Huang X, Rong Y, Yi H, Wang Y. Changes in gut microbial populations, intestinal morphology, expression of tight junction proteins, and cytokine production between two pig breeds after challenge with *Escherichia coli* K88: a comparative study. J Anim Sci. 2013;91:5614–25.
- Xiong Y, Chen L, Fan L, Wang L, Zhou Y, Qin D, et al. Free total rhubarb anthraquinones protect intestinal injury via regulation of the intestinal immune response in a rat model of severe acute pancreatitis. Front Pharmacol. 2018;9:75.
- Jin X, Zhang M, Cao GF, Yang YF. Saccharomyces cerevisiae mannan induces sheep beta-defensin-1 expression via Dectin-2-Syk-p38 pathways in ovine ruminal epithelial cells. Vet Res. 2019;50:8.
- Chen H,Wang C, Huasai S, Chen A. Metabolomics reveals the effects of high dietary energy density on the metabolism of transition angus cows. Animals. 2022:12(9).

- Wang TX, Yao WL, Li J, Shao YF, He QY, Xia J, et al. Dietary garcinol supplementation improves diarrhea and intestinal barrier function associated with its modulation of gut microbiota in weaned piglets. J Anim Sci Biotechnol. 2020;11:12.
- Chen Y, Zhao B, Wu Y, Hu S, Mu L, Zhu C, Pan Y, et al. Impacts of diarrhea on the immune system, intestinal environment, and expression of PGRPs in New Zealand rabbits. PeerJ. 2017;5:e4100.
- Huber D, Balda MS, Matter K. Occludin modulates transepithelial migration of neutrophils. J Biol Chem. 2000;275:5773.
- Van Itallie CM, Anderson JM. Architecture of tight junctions and principles of molecular composition. Semin Cell Dev Biol. 2014;36:157–65.
- Al-Sadi R, Khatib K, Guo S, Ye D, Youssef M, Ma T. Occludin regulates macromolecule flux across the intestinal epithelial tight junction barrier. Am J Physiol-Gastroin Liver Physiol. 2011;300:G1054–64.
- Muza-Moons MM, Schneeberger EE, Hecht GA. Enteropathogenic *Escherichia coli* infection leads to appearance of aberrant tight junctions strands in the lateral membrane of intestinal epithelial cells. Cell Microbiol. 2004;6(8):783–93.
- 39. Erturk-Hasdemir D, Kasper DL. Resident commensals shaping immunity. Curr Opin Immunol. 2013;25:450–5.
- Kim SK, Schluns KS, Lefrancois L. Induction and visualization of mucosal memory CD8 T cells following systemic virus infection. J Immunol. 1999;163:4125–32.
- Uss E, Rowshani AT, Hooibrink B, Lardy NM, van Lier RAW, ten Berge IJM. CD103 is a marker for alloantigen-induced regulatory CD8(+) T cells. J Immunol. 2006;177:2775–83.
- Najafian N, Chitnis T, Salama AD, Zhu B, Benou C, Yuan X, et al. Regulatory functions of CD8+CD28-T cells in an autoimmune disease model. J Clin Invest. 2003;112:1037–48.
- Dzunkova M, Moya A, Vazquez-Castellanos J F, Artacho A, Chen X, Kelly C, et al. Active and secretory IgA-Coated bacterial fractions elucidate dysbiosis in clostridium difficile infection. mSphere. 2016;1.
- McCoy KD, Ronchi F, Geuking MB. Host-microbiota interactions and adaptive immunity. Immunol Rev. 2017;279:63–9.
- Zeng MY, Cisalpino D, Varadarajan S, Hellman J, Warren HS, Cascalho M, et al. Gut microbiota-induced immunoglobulin G controls systemic infection by symbiotic bacteria and pathogens. Immunity. 2016;44:647–58.
- 46. Liu D, Jiang XY, Zhou LS, Song JH, Zhang X. Effects of probiotics on intestinal mucosa barrier in patients with colorectal cancer after operation: meta-analysis of randomized controlled trials. Medicine (Baltimore). 2016;95:e3342.
- Turnbaugh PJ, Hamady M, Yatsunenko T, Cantarel BL, Duncan A, Ley RE, et al. A core gut microbiome in obese and lean twins. Nature. 2009;457:480–4.
- Khan MT, Browne WR, van Dijl JM, Harmsen HJ. How can Faecalibacterium prausnitzii employ riboflavin for extracellular electron transfer? Antioxid Redox Signal. 2012;17(10):1433–40.
- 49. Kim M, Qie Y, Park J, Kim CH. Gut microbial metabolites fuel host antibody responses. Cell Host Microbe. 2016;20:202–14.
- Hou C, Zeng X, Yang F, Liu H, Qiao S. Study and use of the probiotic *Lacto-bacillus reuteri* in pigs: a review. J Animl Sci Biotechnol. 2015;6:14.

### **Publisher's Note**

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

### Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations

BMC

• maximum visibility for your research: over 100M website views per year

#### At BMC, research is always in progress.

Learn more biomedcentral.com/submissions



## Content courtesy of Springer Nature, terms of use apply. Rights reserved.

## Terms and Conditions

Springer Nature journal content, brought to you courtesy of Springer Nature Customer Service Center GmbH ("Springer Nature").

Springer Nature supports a reasonable amount of sharing of research papers by authors, subscribers and authorised users ("Users"), for smallscale personal, non-commercial use provided that all copyright, trade and service marks and other proprietary notices are maintained. By accessing, sharing, receiving or otherwise using the Springer Nature journal content you agree to these terms of use ("Terms"). For these purposes, Springer Nature considers academic use (by researchers and students) to be non-commercial.

These Terms are supplementary and will apply in addition to any applicable website terms and conditions, a relevant site licence or a personal subscription. These Terms will prevail over any conflict or ambiguity with regards to the relevant terms, a site licence or a personal subscription (to the extent of the conflict or ambiguity only). For Creative Commons-licensed articles, the terms of the Creative Commons license used will apply.

We collect and use personal data to provide access to the Springer Nature journal content. We may also use these personal data internally within ResearchGate and Springer Nature and as agreed share it, in an anonymised way, for purposes of tracking, analysis and reporting. We will not otherwise disclose your personal data outside the ResearchGate or the Springer Nature group of companies unless we have your permission as detailed in the Privacy Policy.

While Users may use the Springer Nature journal content for small scale, personal non-commercial use, it is important to note that Users may not:

- 1. use such content for the purpose of providing other users with access on a regular or large scale basis or as a means to circumvent access control;
- 2. use such content where to do so would be considered a criminal or statutory offence in any jurisdiction, or gives rise to civil liability, or is otherwise unlawful;
- 3. falsely or misleadingly imply or suggest endorsement, approval, sponsorship, or association unless explicitly agreed to by Springer Nature in writing;
- 4. use bots or other automated methods to access the content or redirect messages
- 5. override any security feature or exclusionary protocol; or
- 6. share the content in order to create substitute for Springer Nature products or services or a systematic database of Springer Nature journal content.

In line with the restriction against commercial use, Springer Nature does not permit the creation of a product or service that creates revenue, royalties, rent or income from our content or its inclusion as part of a paid for service or for other commercial gain. Springer Nature journal content cannot be used for inter-library loans and librarians may not upload Springer Nature journal content on a large scale into their, or any other, institutional repository.

These terms of use are reviewed regularly and may be amended at any time. Springer Nature is not obligated to publish any information or content on this website and may remove it or features or functionality at our sole discretion, at any time with or without notice. Springer Nature may revoke this licence to you at any time and remove access to any copies of the Springer Nature journal content which have been saved.

To the fullest extent permitted by law, Springer Nature makes no warranties, representations or guarantees to Users, either express or implied with respect to the Springer nature journal content and all parties disclaim and waive any implied warranties or warranties imposed by law, including merchantability or fitness for any particular purpose.

Please note that these rights do not automatically extend to content, data or other material published by Springer Nature that may be licensed from third parties.

If you would like to use or distribute our Springer Nature journal content to a wider audience or on a regular basis or in any other manner not expressly permitted by these Terms, please contact Springer Nature at

onlineservice@springernature.com