



# Protective effect of *Bombyx mori* gloverin on intestinal epithelial cells exposure to enterotoxigenic *E. coli*

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## Abstract

*Bombyx mori* gloverin A2 (BMGlvA2) is an induced antimicrobial insect protein isolated from *Bombyx mori*. This study was conducted to explore the effect and potential mechanisms of BMGlvA2 on inflammatory responses and cellular functions in intestinal epithelial cells (IPEC-J2) exposure to enterotoxigenic *E. coli* (ETEC). IPEC-J2 cells pretreated with or without BMGlvA2 (12.5 µg/mL) were challenged by ETEC K88 (1×10<sup>6</sup> CFU/well) or culture medium. We show that BMGlvA2 pretreatment increased the cell viability and improved the distribution and abundance of tight junction protein ZO-1 in IPEC-J2 cells exposure to ETEC ( $P < 0.05$ ). Interestingly, BMGlvA2 not only decreased the expression levels of inflammatory cytokines such as the tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) and interleukin-1 $\beta$  (IL-1 $\beta$ ), but also decreased the expression level of Caspase3 and the apoptosis rate in the ETEC-challenged cells ( $P < 0.05$ ). Importantly, BMGlvA2 decreased the protein abundances of two critical inflammation-associated signaling proteins, phosphorylated nuclear factor-kappa-B inhibitor alpha (p-I $\kappa$ B $\alpha$ ) and phosphorylated nuclear factor-kappa B (p-NF- $\kappa$ B), in the ETEC-challenged cells. These results indicate that BMGlvA2 attenuates ETEC-induced inflammation in the IPEC-J2 cells by regulating the NF- $\kappa$ B signaling pathway, resulting in decreased secretion of inflammatory cytokine and reduced cell apoptosis.

**Keywords** Antimicrobial peptide · BMGlvA2 · ETEC · Inflammation · IPEC-J2 cells

## Introduction

Microbial infection increases diarrhea rate and mortality of neonatal animals. A variety of antibiotics have been used to treat most bacteria-induced infections; however, the emergence of antibiotic resistance not only decreases the effectiveness of traditionally used antibiotics, but also leads to their residues in animal products and causes various toxic

effects, including allergy, immunopathological effects, carcinogenicity, mutagenicity, nephropathy (gentamicin) and bone marrow toxicity (chloramphenicol) and even anaphylactic shock in humans [1–5]. Thus, there is an urgent need to develop novel substitutes for antibiotics [6].

Antimicrobial peptides (AMPs), also known as host defense peptides, are a large class of small cationic polypeptide molecules, which widely exist in various organisms to protect themselves from the infection of bacteria, fungi, viruses, etc. [7, 8]. Previous study indicated that AMPs exert their roles through destroying bacterial cell membrane, modulating immune response, and regulating inflammation [9]. For instance, amphoteric AMPs destroy membrane integrity through hydrophobic and electrostatic interactions, resulting in membrane leakage and death of cancer cells [10, 11]. Cecropin B has potential anticancer activity against leukemia cells, gastric cancer, and lung cancer [12]. A variety of amphibian AMPs, including magainins, aureins, tryptophyllins, demaseptin, phylloseptin, bombinins (BLP-7), and bombinin (H-BO), showed antiproliferative activities [13]. BMGlvA2, an induced antimicrobial insect protein isolated from *Bombyx mori* (*B.*

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*mori*), is a small cationic linear  $\alpha$ -helical peptide belonging to cecropin family [14]. In previous study, BMGlvA2 was found to inhibit the growth of bacteria and increase the permeability of bacterial membrane by binding to the lipopolysaccharide (LPS) components on the surface of bacterial membrane [15]. However, it has no toxic effects on mammalian cells [14]. Although increasing evidences have shown that the AMPs can serve as a critical regulator for diverse biological events including immune responses [16–19], the mechanism behind the BMGlvA2-regulated immunity is just beginning to be explored.

In our previous study, the BMGlvA2 was successfully expressed and purified by using a heterologous system, and the recombinant BMGlvA2 was found to show a moderate antibacterial activity against various gram-negative and gram-positive bacteria [20]. In this study, we explored its protective effect and potential mechanisms of BMGlvA2 on inflammatory responses and cellular functions in intestinal epithelial cells (IPEC-J2) exposure to ETEC K88.

## Materials and methods

### Expression and purification of BMGlvA2

The engineered strain L Orgami B (DE3)-harboring the recombinant plasmid (pet32a-gloverin A2) was previously constructed in our laboratory [20]. The protein expression was induced by 1.0 mM isopropyl  $\beta$ -D-1 thiogalactoside (IPTG). After incubation for 8 h at 28 °C, bacterial cells were harvested by centrifugation at 8000 r/min for 15 min at 4 °C, 20 mL phosphate-buffered solution (PBS) was added to washing the precipitation, centrifuged at 8000 r/min for 15 min at 4 °C, and the supernatant was discarded. Then, 20 mL lysis buffer was added and incubated at 4 °C overnight. Then, schizolytic cells were sonicated (4-s pulse and 8-s interval; 30 cycles; Sonics-Vibra cell, USA). The supernatant was harvested by centrifugation at 8000 r/min for 20 min at 4 °C. The supernatant obtained above was filtered by 0.22- $\mu$ m filter, and then applied to Ni<sup>2+</sup>-IDA column (Sangon Biotech, China) and purified according specification. Protein concentration was quantified by the BCA assay (Beyotime, China).

### Cell culture

Intestinal porcine epithelial cells (IPEC-J2) were cultured in a 75-cm<sup>2</sup> cell culture flask in DMEM-F12 with 10% FBS, 100 U/mL penicillin, and 100  $\mu$ g/mL streptomycin.  $1 \times 10^5$  cells/well were seeded in 12-well plates and grown to ~60% confluence at 37 °C in a CO<sub>2</sub> incubator (5% v/v). On reaching 80–90% confluence, the cells were routinely passaged at a 1:3 split ratio (approximately every 2–3 days).

### E. coli K88 challenge

IPEC-J2 cells were seeded in 12-well cell culture plates ( $2 \times 10^5$  cells/well), cultured overnight to make cells adhere and then incubated with different concentrations of *E. coli* K88 (0,  $10^5$ ,  $10^6$ ,  $10^7$ , and  $10^8$  CFU/well) for 0.5, 1, 2, and 3 h to choose the suitable *E. coli* K88 treatment dose and duration for inducing inflammatory injury. Finally, the cells were collected for gene expression analysis.

### BMGlvA2 pretreatment

IPEC-J2 cells were seeded in 12-well cell culture plates at a density of  $2 \times 10^5$  cells/well and then maintained at 37 °C in a CO<sub>2</sub> incubator (5% v/v). After reaching sub-confluence, the cells were pretreated with different concentrations of BMGlvA2 (0, 12.5, 25, 50, and 100  $\mu$ g/mL) for 24 h. Finally, the cells were collected for gene expression analysis.

### Cell treatment

IPEC-J2 cells were cultured in a 75-cm<sup>2</sup> cell culture flask in DMEM-F12 with 10% FBS, 100 U/mL penicillin, and 100  $\mu$ g/mL streptomycin.  $1 \times 10^5$  cells/well were seeded in 12-well plates and grown to ~60% confluence at 37 °C in a CO<sub>2</sub> incubator (5% v/v). After incubated with antimicrobial peptides BMGlvA2 (12.5  $\mu$ g/mL) for 24 h or BAY11-7081 (1  $\mu$ M) for 2 h, then cells were challenged with  $1 \times 10^6$  CFU/well ETEC K88 for 1 h or 2.5 h (only for assessment of apoptosis). It is worth noting that when challenged with ETEC K88, the cells were cultured in DMEM F12 medium supplemented with 2% FBS (without any antibiotics).

### Cell viability assay

Cytotoxic effects on cell viability were measured by the MTT [3-(4, 5-dimethylthiazol-2-yl)-2, 5-diphenyltetrazolium] assay as previously described (Beyotime, Shanghai, China) after treatment as designed. After treatment, the supernatants were removed and cells were gently washed three times with phosphate-buffered saline (PBS, pH 7.4) to remove non-adherent bacteria. Then, 10  $\mu$ L of MTT solution (5 mg/mL) was added to each well and plates were incubated for 4 h at 37 °C. After incubation, 150  $\mu$ L of DMSO (dimethyl sulfoxide) was added into each well and incubated in a cell incubator for 10 min. The absorbance of the plate was determined at 570 nm by a UV-1100

spectrophotometer (ShangHai, China). All experiments were performed in triplicate.

### RNA extraction and RT-PCR

Total RNA was isolated from IPEC-J2 cells using RNAiso Plus reagent (TaKaRa, Dalian, China). The quantity and quality of the isolated RNA were determined by absorbance at 260 and 280 nm [21]. And then, cDNA was synthesized using a reverse transcriptase kit (Takara, Dalian, China). Briefly, quantitative PCR was performed by QuanStudio 6 Flex Real-Time PCR detection system (Applied Biosystems, Foster City, CA, USA) with a total of 10  $\mu$ L of assay solution containing 5  $\mu$ L SYBR Green mix (Takara), 0.2  $\mu$ L Rox, 3  $\mu$ L deionized H<sub>2</sub>O, 1  $\mu$ L cDNA template, and 0.4  $\mu$ L each of forward and reverse primers (Qingke, China). The relative gene expressions compared with the housekeeping gene  $\beta$ -actin were calculated by  $2^{-\Delta\Delta Ct}$  [22].

### Total protein extraction and western blot analysis

Protein samples were extracted using lysis buffer (Beyotime, Shanghai, China) according to the manufacturer's recommendations. Western blot analysis was performed as before [23]. Briefly, proteins (25  $\mu$ g/lane) were resolved by electrophoresis on 12% SDS-poly-acrylamide gels and then transferred to polyvinylidene fluoride (PVDF) membranes (Bio-Rad Laboratories, Inc., Richmond, CA, USA). The PVDF membranes were blocked (1 h at room temperature) in 5% non-fat dry milk in Tris-buffered saline containing 0.1% Tween-20 (TBS-T) before incubation with primary antibody. After thoroughly rinsing with TBS-T, the membranes were respectively incubated with rabbit anti-Tight Junction Protrin 1 (ZO1) (at 1:500 dilution, Novus Biologicals), rabbit anti-phospho-nuclear factor- $\kappa$ B (pNF- $\kappa$ B) p65 (at 1:500 dilution, Cell Signalling Technology, Inc., Danvers, MA, USA), mouse anti-NF- $\kappa$ B (at 1:500 dilution, Cell Signalling Technology), mouse anti-phospho-I $\kappa$ b $\alpha$  (pI $\kappa$ b $\alpha$ ) (at 1:500 dilution, Invitrogen), mouse anti-I $\kappa$ b $\alpha$  (at 1:500 dilution, Cell Signalling Technology), and rabbit anti-GAPDH (at 1:15,000 dilution, Abcam plc) antibodies, with gentle agitation overnight at 4°C. Subsequently, the membranes were rinsed several times with TBS-T and then incubated with HRP-conjugated goat anti-rabbit IgG or anti-mouse IgG secondary antibody, at room temperature for 1 h (at 1:5000 dilution; Abcam plc). Finally, the membranes were rinsed several times with the TBS/T buffer at room temperature for 10 min each time. Blots were developed using a Clarity™ Western ECL Substrate (Bio-Rad Laboratories, Inc). The bands were visualized by exposure to X-OMAT BT films (Beyotime Institute of Biotechnology, Shanghai, China) for 1 min, and were quantified by using Quantity One software (Bio-Rad Laboratories, Inc). The relative abundance of each

target protein was expressed as the ratio of targeted protein to GAPDH protein.

### Immunofluorescence

Preparation of cell climbing slice: add 50  $\mu$ L culture medium into the 6-well plate, put the treated with concentrated sulfuric acid coverslips, and press the coverslips with the gun head to remove the bubbles. Then, 500  $\mu$ L of cell suspension was added to the coverslips. After 6 h, 1.5 mL medium was added to each well.

The cells were washed with PBS (pH7.4) three times after treatment according to the experimental design. Add 1 mL formaldehyde into each well and stand for 20 min at room temperature. Wash three times with PBS (pH 7.4) for 2 min each time, and then seal with 3% BSA (Beyotime Institute of Biotechnology, Shanghai, China) at room temperature for 30 min. Wash three times with PBS (pH 7.4) for 2 min each time, and then, samples were incubated overnight with the primary antibody at 4°C and kept dark from light (rabbit anti-ZO-1; Novus). The samples were washed three times with PBS (pH 7.4) for 2 min each time and incubated for 2 h at room temperature with the appropriate secondary antibody (Alexa Fluor 488 conjugated goat anti-rabbit immunoglobulin (CST)). Samples with the biotinylated secondary antibody were incubated with Avidin Alexa Fluor 488 (CST) for visualization of the secondary antibody. Then, the nuclei were counterstained with DAPI (Sigma-Aldrich). Samples were imaged using a confocal scanning microscope (NIKON ECLIPSE TI-SR).

### Assessment of apoptosis by flow cytometry

The proportion of apoptotic cells in IPEC-J2 cells was determined by flow cytometry (CytoFlex, Beckman Coulter, Inc., Brea, CA, USA) using PE Annexin V Apoptosis Detection Kit I (Becton, Dickinson and Company, BD Biosciences, San Jose, CA, USA). When IPEC-J2 cells were grown to ~60% confluence at 37 °C in a CO<sub>2</sub> incubator (5% v/v), they were incubated with BMG1vA2 for 24 h (BMG1vA2, BMG1vA2 + *E. coli* K88). When IPEC-J2 cells were grown to ~90% confluence at 37 °C in a CO<sub>2</sub> incubator (5% v/v), they were incubated with Bay for 2 h (Bay, Bay + *E. coli* K88). Cells were challenged with  $1 \times 10^6$  CFU/well *E. coli* K88 for 2.5 h (BMG1vA2 + *E. coli* K88, *E. coli* K88, Bay + *E. coli* K88) before sample collection; control cells were cultured in a culture medium without any treatment. Treated cells were harvested and labeled with an anti-Annexin V-FITC Apoptosis Detection Kit (BD Biosciences, USA). Floating cells were collected; then, attached cells were washed with 0.01 M PBS and digested with trypsin for 2 min. Finally, the digested cells and floating cells were added together to centrifugate at 350 g for 10 min; then, the

cells were stained with 2  $\mu\text{L}$  of Annexin V-FITC fluorescent dye at 4  $^{\circ}\text{C}$  in the dark. After 10 min, add 2  $\mu\text{L}$  of PI staining for 5 min at 4  $^{\circ}\text{C}$  in the dark. Finally, detection of apoptotic cells was completed within 1 h after the addition of 400  $\mu\text{L}$  Annexin V binding buffer ( $1\times$ ).

### Statistics analysis

All statistical analysis was performed using SPSS 21.0 software. Data were expressed as the mean  $\pm$  standard error of the mean (SEM). Statistical analysis of treatment of IPEC-J2 was carried out using two-way ANOVA followed by Duncan's multiple comparisons test. Image production was carried using GraphPad Prism software (Version 7. GraphPad Software Inc., CA, USA).

## Results

### Influences of enterotoxigenic *Escherichia coli* challenge on viability and inflammatory response of IPEC-J2 cells

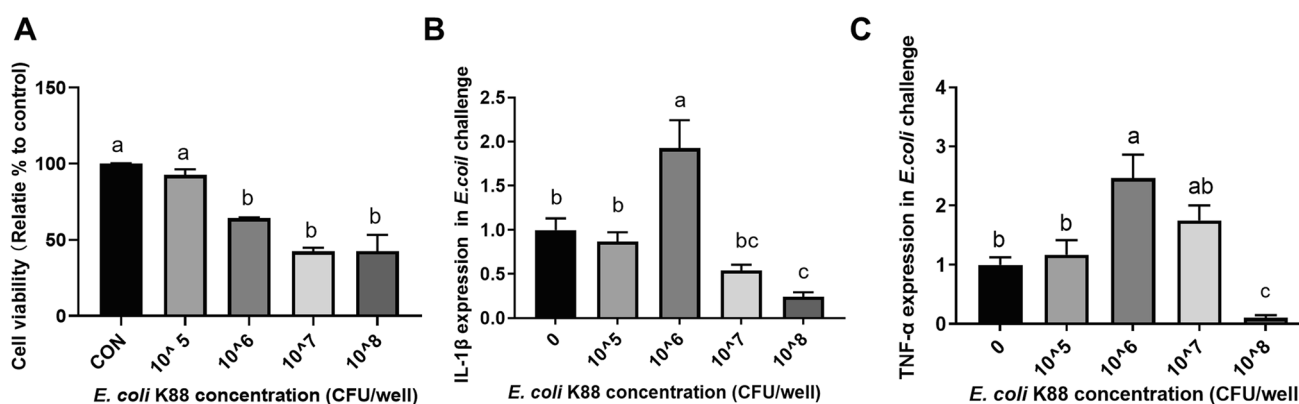
To explore the influence of enterotoxigenic *Escherichia coli* (ETEC) challenge on cell viability, IPEC-J2 cells were treated by ETEC at different final concentrations ( $1\times 10^5$ ,  $1\times 10^6$ ,  $1\times 10^7$ , and  $1\times 10^8$  CFU/well) for 3 h. As shown in Fig. 1A, the cell viability was significantly decreased upon ETEC challenge at a dose higher than  $1\times 10^5$  CFU/well. Moreover, the expression levels of IL-1 $\beta$  and TNF- $\alpha$  were significantly increased in the IPEC-J2 cells exposure to ETEC at a dose of  $1\times 10^6$  CFU/well. But a higher dose ( $1\times 10^7$  and  $1\times 10^8$ ) led to downregulation of the IL-1 $\beta$  and TNF- $\alpha$  expression in the IPEC-J2 cells (Fig. 1B, C). Hence, the ETEC treatment dose used in further experiments was  $1\times 10^6$  CFU/well.

### Effect of BMGlvA2 on cell viability and inflammatory responses of IPEC-J2 cells upon ETEC challenge

In this study, the BMGlvA2 has been successfully expressed and purified (Fig. S1). To explore the influence of BMGlvA2 on cell viability, the IPEC-J2 cells were treated with BMGlvA2 at different concentrations (0, 4, 20, and 100  $\mu\text{g}/\text{mL}$ ) for 24 h. As shown in Fig. 2, BMGlvA2 had no negative influence (toxic effect) on cell viability in the range from 4 to 100  $\mu\text{g}/\text{mL}$  ( $P > 0.05$ ). Therefore, a moderate dose of 12.5  $\mu\text{g}/\text{mL}$  was used for further studies. As shown in Fig. 3A, ETEC challenge decreased the viability in the IPEC-J2 cells. However, pretreatment of the cells with 12.5  $\mu\text{g}/\text{mL}$  BMGlvA2 significantly increased the cell viability upon ETEC challenge ( $P < 0.05$ ). Moreover, ETEC challenge significantly elevated the expression levels of inflammatory cytokines such as the IL-1 $\beta$ , IL-6, and TNF- $\alpha$  in the IPEC-J2 cells ( $P < 0.05$ ). However, BMGlvA2 pretreatment significantly downregulated the expression levels of IL-1 $\beta$  and TNF- $\alpha$  in the IPEC-J2 cells exposure to ETEC (Fig. 3B–D).

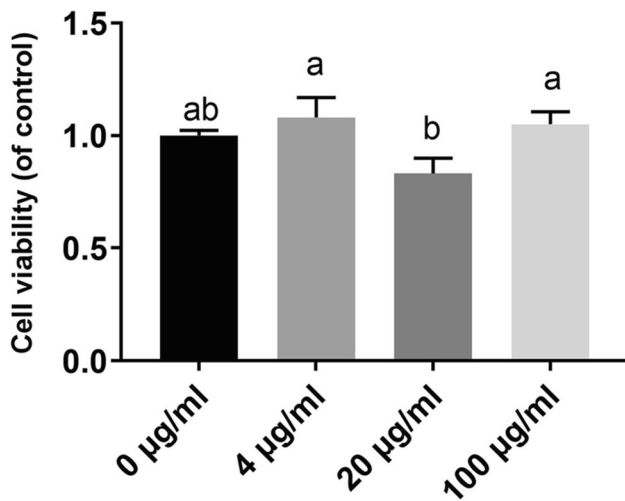
### Effect of BMGlvA2 on tight junction protein distribution and abundance in IPEC-J2 cells upon ETEC challenge

We explored the distribution and abundance of zonula occludens-1 (ZO-1), one of the major tight junction-related proteins located in the IPEC-J2 cells by immunofluorescence analysis. We found that the ZO-1 staining in IPEC-J2 cells was diffuse due to ETEC challenge, with less staining in the tight intercellular junction area, indicating disruption of the tight junction upon ETEC infection. However, BMGlvA2 pretreatment attenuated the ETEC-induced disruption by improving the localization



**Fig. 1** ETEC challenge affects cell viability and inflammatory gene expression in IPEC-J2 cells. IPEC-J2 cells ( $1\times 10^5$  cells/well) were treated with *E. coli* K88 at different final concentrations ( $1\times 10^5$ ,  $1\times 10^6$ ,  $1\times 10^7$ , and  $1\times 10^8$  CFU/well) for 3 h. **A**, cell viability; **B**, the

expression of IL-1 $\beta$ ; **C**, the expression of TNF- $\alpha$ . Data are presented as mean  $\pm$  SEM. <sup>a–b</sup>Values within a column differ if they do not share a common superscript ( $P < 0.05$ )



**Fig. 2** The influence of BMGlvA2 on cell viability in IPEC-J2 cells. IPEC-J2 cells were pretreated with various concentrations of BMGlvA2 (0, 4, 20, and 100 µg/mL) for 24 h. Data are presented as mean  $\pm$  SEM. <sup>a–b</sup>Values within a column differ if they do not share a common superscript ( $P < 0.05$ )

and abundance of the ZO-1 proteins in IPEC-J2 cells (Fig. 4A). The abundance of ZO-1 protein was validated by using western blot. We found that ETEC challenge significantly decreased the protein abundance of ZO-1 in the IPEC-J2 cells (Fig. 4B). However, BMGlvA2 pretreatment significantly elevated the protein abundance of ZO-1 in the ETEC-challenged cells ( $P < 0.05$ ). Moreover, BMGlvA2 pretreatment significantly increased the expression levels of major tight junction-related proteins ZO-1 and Occludin in the ETEC-challenged cells (Fig. 4C) ( $P < 0.05$ ).

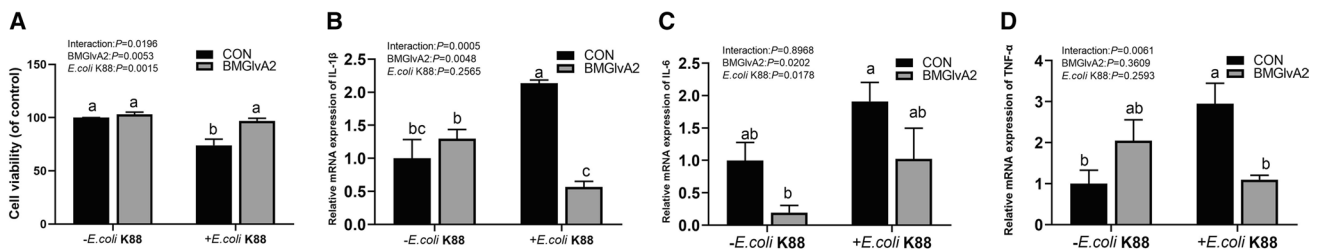
### Effect of BMGlvA2 on apoptosis of IPEC-J2 cells upon ETEC challenge

As shown in Fig. 5A and B, as compared to the control group, ETEC challenge significantly elevated the early, late and total apoptosis rates in the IPEC-J2 cells ( $P < 0.05$ ). However, BMGlvA2 pretreatment significantly decreased

the late apoptosis and total apoptosis rates in the ETEC-challenged cells ( $P < 0.05$ ). Additionally, ETEC challenge significantly elevated the expression levels of critical apoptosis-related genes such as the caspase 3, caspase 8, and caspase 9 in the cells (Fig. 5C). BMGlvA2 pretreatment significantly downregulated the expression level of apoptosis-related gene caspase3 in the ETEC-challenged cells ( $P < 0.05$ ).

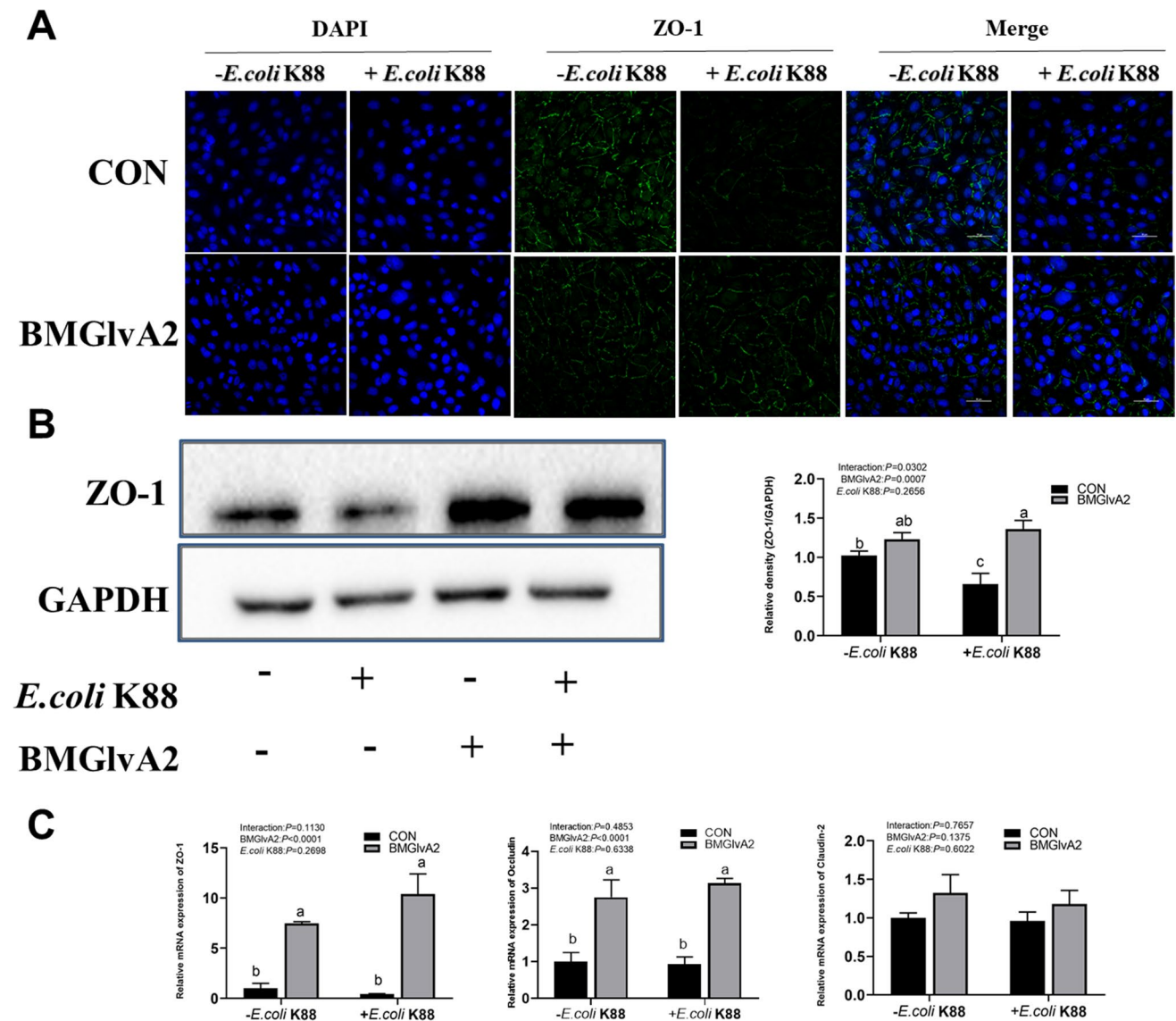
### BMGlvA2 suppressed ETEC-induced cell apoptosis and inflammatory response via suppressing the I $\kappa$ B $\alpha$ /NF- $\kappa$ B signaling

NF- $\kappa$ B transcription factor plays an important role in cell survival and apoptosis, and participates in a variety of inflammatory signal transduction. To further explore if the BMGlvA2-modulated inflammatory responses in the intestinal epithelial cells are associated with the NF- $\kappa$ B signaling pathway; the IPEC-J2 cells were pretreated with BMGlvA2 and NF- $\kappa$ B inhibitor and subsequently challenged by ETEC. To further explore whether the inflammatory response of intestinal epithelial cells regulated by BMGlvA2 is associated with NF- $\kappa$ B signaling pathway, IPEC-J2 cells were pretreated with BMGlvA2 and NF- $\kappa$ B inhibitor (Bay 11–7082) and subsequently challenged by ETEC. Both significantly inhibit the ETEC-induced inflammatory response, which is manifested as decreased apoptosis (such as early apoptosis, late apoptosis, and total apoptosis) and secretion of critical apoptosis-related genes (Caspase 3 and Caspase 9) and inflammatory cytokines (such as IL-1 $\beta$ , IL-6, and TNF- $\alpha$ ) (Fig. 6). Meanwhile, BMGlvA2 and Bay pretreatment improved distribution and abundance of ZO-1 in the IPEC-J2 cells exposure to ETEC (Fig. 7A). Moreover, we detected the expression levels of key proteins in the signaling pathway by western blot analysis (Fig. 7B). The results showed that ETEC significantly elevated the abundance of phosphorylated I $\kappa$ B $\alpha$  (p-I $\kappa$ B $\alpha$ ) and phosphorylated NF- $\kappa$ B (p-NF- $\kappa$ B) in the IPEC-J2 cells ( $P < 0.05$ ). However, BMGlvA2 and Bay pretreatment significantly decreased their expression levels in the ETEC-challenged cells ( $P < 0.05$ ).



**Fig. 3** The influence of BMGlvA2 on cell viability and inflammatory responses in IPEC-J2 cells induced by ETEC. IPEC-J2 cells were pretreated with 12.5 µg/mL BMGlvA2 for 24 h, followed by co-treatment with *E. coli* K88 ( $1 \times 10^6$  CFU/well) for another 1 h. **A**, cell

viability; **B**, the expression of IL-1 $\beta$ ; **C**, the expression of IL-6; **D**, the expression of TNF- $\alpha$ . Data are presented as mean  $\pm$  SEM. <sup>a–b</sup>Values within a column differ if they do not share a common superscript ( $P < 0.05$ )



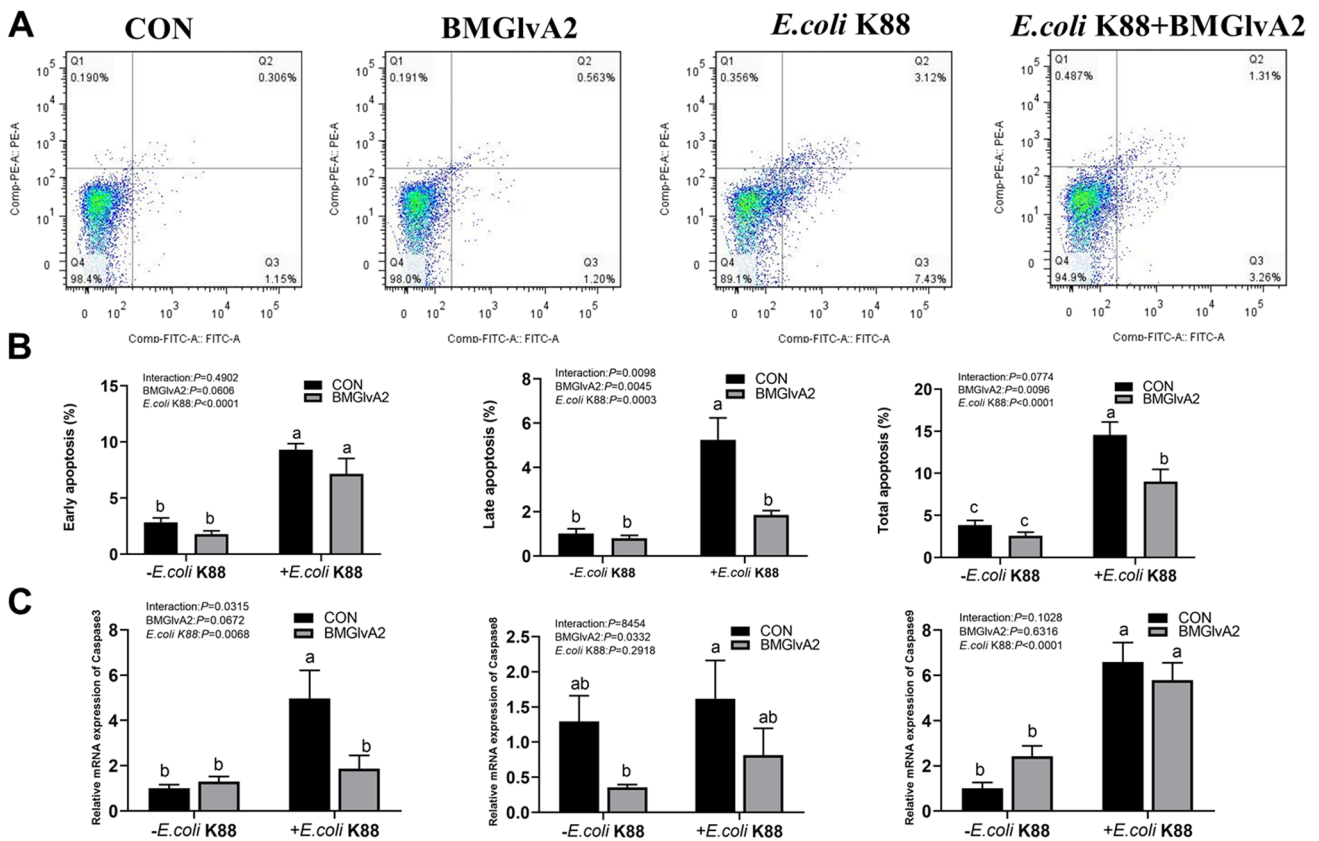
**Fig. 4** The influence of BMGlvA2 on tight junction protein distribution and abundance in IPEC-J2 cells induced by ETEC. IPEC-J2 cells were pretreated with 12.5  $\mu\text{g}/\text{mL}$  BMGlvA2 for 24 h, followed by co-treatment with *E. coli* K88 ( $1 \times 10^6$  CFU/well) for another 1 h. **A**, immunofluorescence of ZO-1; **B**, western blot analysis of protein

expression of ZO-1; **C**, tight junction-related proteins. Representative immunofluorescent images for detection of ZO-1 (green) and DAPI (blue). Scale bar = 50  $\mu\text{m}$ . Data are presented as mean  $\pm$  SEM. <sup>a-c</sup>Values within a column differ if they do not share a common superscript ( $P < 0.05$ )

## Discussion

The integrity of intestinal epithelial barrier not only is the key to maintain the health and disease of the body [24], but also plays an important role in maintaining homeostasis of host intestinal epithelium [25]. Destruction of intestinal barrier function may lead to chronic immune activation, leading to local and systemic infections or diseases, including coeliac disease, colorectal cancer, inflammatory bowel disease (IBD), and metabolic disorders [26–28]. ETEC is a kind of common

pathogenic bacteria, which produces heterogeneous fimbriae or non-fimbriae adhesins, attaches bacteria to the host receptor, and colonizes in the small intestine of animals [29]. Meanwhile, they transfer heat-labile and heat-stable enterotoxins to intestinal epithelial cells, destroy homeostasis, disrupt homeostasis, and cause excessive fluid hyper secretion and watery diarrhea [30]. Previous studies have shown that Cecropins (e.g., Cecropin A, Cecropin AD, and Mdc) alleviates intestinal inflammation in *E. coli*-induced animals by modulating the MyD88-NF $\kappa$ B signaling pathway [31–33].



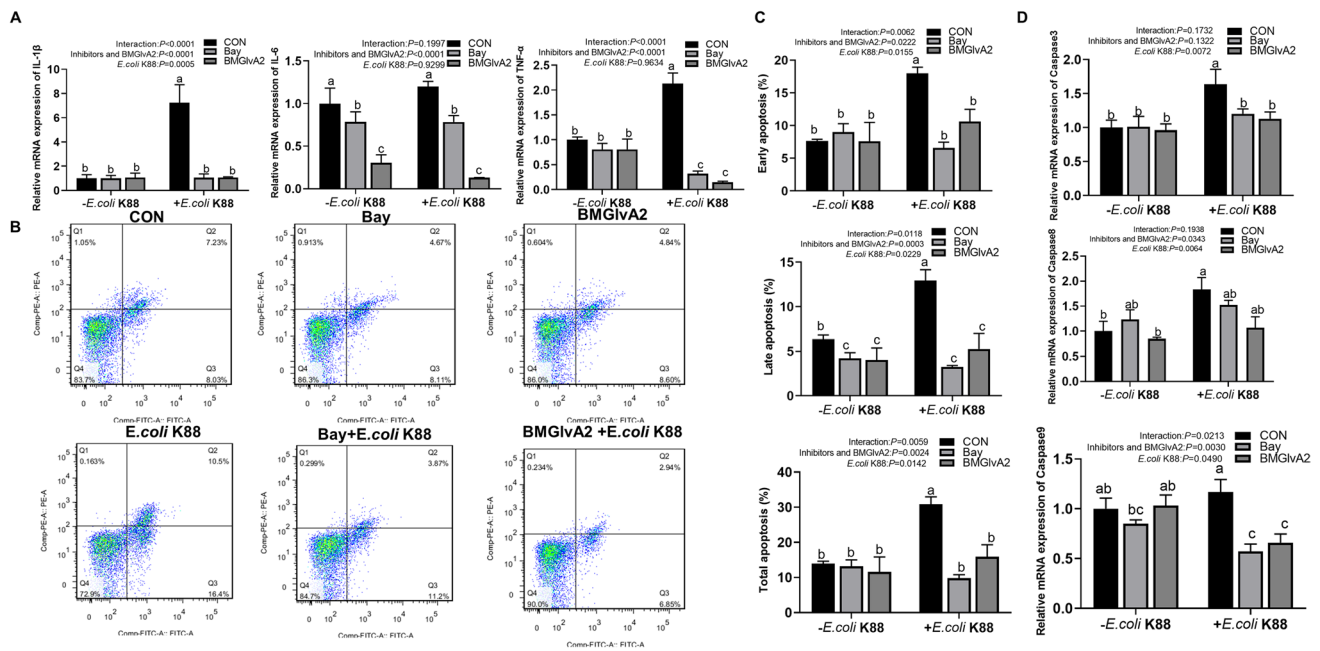
**Fig. 5** The influence of BMGIvA2 on cell apoptosis in IPEC-J2 cells induced by ETEC. IPEC-J2 cells were pretreated with 12.5  $\mu$ g/mL BMGIvA2 for 24 h, followed by co-treatment with *E. coli* K88 ( $1 \times 10^6$  CFU/well) for another 2.5 h. **A**, apoptosis of flow cytometry;

**B**, cell apoptosis rate; **C**, apoptosis-related genes. Data are presented as mean  $\pm$  SEM. <sup>a-c</sup>Values within a column differ if they do not share a common superscript ( $P < 0.05$ )

In the present study, ETEC challenge significantly elevated the expression levels of inflammatory cytokines such as the IL-1 $\beta$  and TNF- $\alpha$  in the IPEC-J2 cells, which was consistent with previous studies using the same cell line [34, 35]. ETEC activates the innate immune system of host cells, producing pro-inflammatory and chemokines [36, 37]. However, BMGIvA2 pretreatment significantly decreased the expression levels of IL-1 $\beta$  and TNF- $\alpha$  in ETEC-induced cells. Intestinal epithelial integrity is regulated by the tight junction [38, 39]. Substantial evidence has shown that the pro-inflammatory cytokine TNF- $\alpha$  can reduce the expression of ZO-1 and Claudin-2, and destroy the structural integrity of intestinal epithelial cells [40–42]. In this study, we found that BMGIvA2 increased the expression of tight junction proteins and inhibited the decrease of ZO-1 abundance in IPEC-J2 cells induced by ETEC. These results showed that an imbalance in pro- and anti-inflammatory cytokine levels has been correlated with ETEC-induced intestinal epithelial cell injury. This

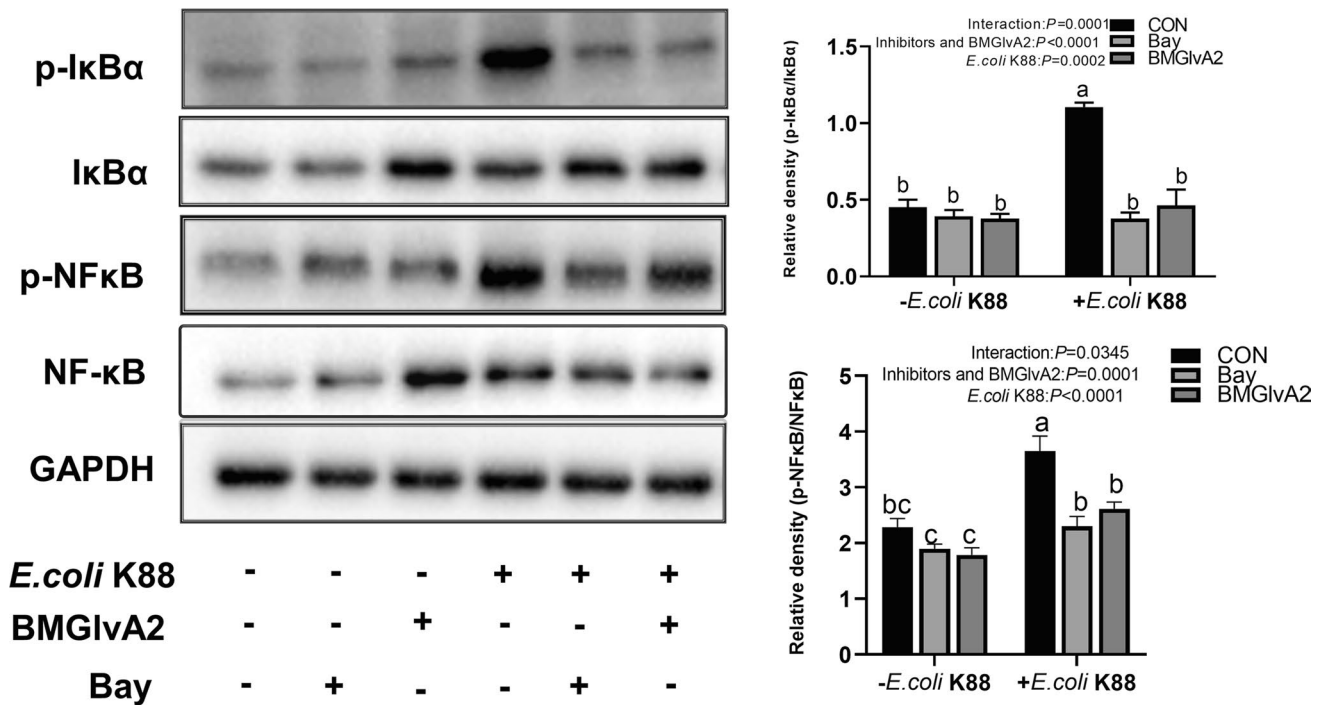
is similar to previous studies [43, 44]. BMGIvA2 can increase the expression of tight junction protein in intestinal epithelial cells and improve the structural integrity of intestinal epithelial cells by inhibiting the expression of pro-inflammatory factors.

Apoptosis is the process of programmed cell death. Two pathways (intrinsic pathway and extrinsic pathway) are associated with apoptosis [45, 46]. In cells, apoptosis can be regulated by Bax and Bcl-2, which plays a role through p53 signal pathway and Caspase protein [29]. Caspase 3 directly promotes apoptosis through Caspase 8 of death receptor pathway and Caspase 9 of mitochondrial pathway [47]. In the present study, we detected the apoptosis of IPEC-J2 cells by flow cytometry. These results showed that the expression of Caspase 3, 8, and 9 were significantly upregulated in ETEC-induced cells, and their expressions were significantly decreased by BMGIvA2, which was consistent with previous research [48]. Moreover, ETEC significantly increased the early, late, and total apoptosis,



**Fig. 6** BMGIvA2 inhibits ETEC-induced cell apoptosis and inflammation response by inhibiting the IκB-α/NF-κB signaling. IPEC-J2 cells were pretreated with 12.5 μg/mL BMGIvA2 for 24 h or Bay for 2 h, followed by co-treatment with *E. coli* K88 (1 × 10<sup>6</sup> CFU/well) for

another 1 or 2.5 h. **A**, inflammatory mediator; **B**, apoptosis of flow cytometry; **C**, cell apoptosis rate; **D**, apoptosis-related genes. Data are presented as mean ± SEM. <sup>a-c</sup>Values within a column differ if they do not share a common superscript (*P* < 0.05)



**Fig. 7** BMGIvA2 improved ETEC-induced tight junction protein distribution and abundance by inhibiting the IκB-α/NF-κB signaling. IPEC-J2 cells were pretreated with 12.5 μg/mL BMGIvA2 for 24 h or Bay for 2 h, followed by co-treatment with *E. coli* K88 (1 × 10<sup>6</sup> CFU/

well) for another 1 h. **A** Western blot analysis of protein expression of NF-κB signaling pathway. Data are presented as mean ± SEM. <sup>a-b</sup>Values within a column differ if they do not share a common superscript (*P* < 0.05)



while BMGlvA2 pretreatment significantly decreased the early, late, and total apoptosis. These results indicate that BMGlvA2 can inhibit the expression of Caspase gene and reduce apoptosis.

To further explore the anti-inflammatory mechanism of BMGlvA2, we used Bay11-7082 (specific inhibitors of NF- $\kappa$ B signaling pathway) to investigate whether NF- $\kappa$ B signaling pathway is involved. The NF- $\kappa$ B transcription factor can be activated by a series of stimuli (e.g., pathogenic derived substances, inflammatory cytokines, and a variety of enzymes) [49, 50], which play an important role in cell survival and apoptosis (Girard et al. 2009). Activation of NF- $\kappa$ B upregulated Bcl-Xs in rat hippocampal studies, and these effects were inhibited by I $\kappa$ B $\alpha$  [51]. Under physiological conditions, I $\kappa$ B proteins present in the cytoplasm inhibit NF- $\kappa$ B [52]. Phosphorylation of I $\kappa$ B leads to its degradation by proteasome, followed by release of NF- $\kappa$ B for nuclear translocation and gene transcription activation [53]. This pathway regulates the production of pro-inflammatory cytokines and recruitment of inflammatory cells, which contribute to the inflammatory response [54, 55]. In agreement with previous studies [56], elevated level of phosphorylated NF- $\kappa$ B and I $\kappa$ B $\alpha$  protein expression was documented in the IPEC-J2 cells induced by ETEC. However, BMGlvA2 inhibits the NF- $\kappa$ B and I $\kappa$ B $\alpha$  phosphorylation. The inhibition of phosphorylation of I $\kappa$ B $\alpha$  and NF- $\kappa$ B by BMGlvA2 was similar to that of inhibitor. Thus, we speculated that BMGlvA2 play a potential regulatory role by inhibiting I $\kappa$ B $\alpha$ /NF- $\kappa$ B pathway. Additionally, we also found that Bay11-7082 and BMGlvA2 had similar effects on inflammatory factors, tight junction proteins, and apoptosis, which further indicate that BMGlvA2 protects IPEC-J2 cells by inhibiting NF- $\kappa$ B signaling pathway.

In conclusion, BMGlvA2 can attenuate ETEC-induced inflammation and injury in the IPEC-J2 cells which was associated with decreased secretion of inflammatory cytokine, reduced cell apoptosis, and improved intestinal barrier functions. Moreover, the BMGlvA2 may act as a novel inhibitor of the NF- $\kappa$ B signaling pathway that can be tentatively used for the treatment of various inflammatory bowel diseases.

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**Author contribution** All authors contributed significantly to the work. JH conceived the study. QL designed and performed experiments and analyzed results with the help of QQF, GQS, DWC, BY, YHL, PZ, and XBM. QL wrote the manuscript with help of ZQH, JY, JQL, and HY.

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**Data availability** The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

## Declarations

**Competing interest** The authors declare no competing interests.

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