Basic Study

Herbs-partitioned moxibustion alleviates aberrant intestinal epithelial cell apoptosis by upregulating A20 expression in a mouse model of Crohn’s disease

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Abstract

BACKGROUND
A20 inhibits intestinal epithelial cell apoptosis in Crohn’s disease, and herbs-partitioned moxibustion (HPM) has been demonstrated to be an effective treatment for Crohn’s disease. However, the mechanism by which HPM reduces intestinal epithelial cell apoptosis in Crohn’s disease has not been thoroughly elucidated to date.

AIM
To elucidate whether HPM exerts its effects by upregulating A20 to affect intestinal epithelial cell apoptosis in a Crohn’s disease mouse model.

METHODS
In this study, mice with A20 deletion in intestinal epithelial cells (A20IEC-KO) were utilized to establish a Crohn’s disease mouse model with 2,4,6-trinitrobenzene.
INTRODUCTION

Crohn's disease is a chronic inflammatory bowel disease with symptoms of abdominal pain, diarrhea, weight loss, perianal lesions, and anemia[1]. In the past ten years, Crohn's disease has increased significantly as a worldwide health problem[2]. The highest reported prevalence areas are Europe and North America with 322 cases per 100000[3,4]. The high prevalence and long-term nature of the disease pose a great burden on patients and the society, due to health-related reduction in quality of life and decreased economic productivity, which calls for high-quality and cost-efficient care for patients.

Crohn's disease most likely results from complex interactions between genetics, environment, and gut microbiota, which lead to dysfunction of the epithelial barrier.
with consequent deregulation of the mucosal immune system and responses to gut microbiota[9]. The intestinal epithelium is a notably large mucosal surface with a strong capacity for self-renewal[9]. Proliferation of progenitor cells and their differentiation into mature epithelial cells continuously recompense cell losses[9]. However, in Crohn’s disease, this homeostasis is disrupted by increased apoptosis of epithelial cells due to increased stimulation of the immune system, leading to loss of epithelial integrity and local inflammation[9]. In many patients with Crohn’s disease, epithelial injury and inflammation due to increased cell apoptosis depend on tumor necrosis factor (TNF)-α, TNF-alpha (TNF-α), a transmembrane protein, binds to its ligand, TNF receptor (TNFR) 1, recruiting TNFR1-associated death domain (TRADD) and receptor-interacting protein (RIP). In turn, TRADD and RIP associate with FAS-associated death domain protein (FADD) to activate caspase 8, leading to apoptosis[10]. Studies have shown that lower expression of A20 correlates with improved anti-TNF-α drug responses[11], and mice with A20 deletion in intestinal epithelial cells (A20IEC-KO) are hypersensitive to TNF-α-induced intestinal epithelial apoptosis[12]. These studies have revealed a negative regulatory role of A20 in the TNF-α-induced apoptosis pathway.

A20 is a cytoplasmic protein that was originally identified as a primary TNF-α-induced responsive molecule in endothelial cells and negatively regulates NF-kB-dependent gene expression in response to stimuli, such as TNF-α and interleukin-1 (IL-1)[15,16]. A20 contains an N-terminal ovarian tumor deubiquitinating and E3 ubiquitin ligase activity toward the death-domain-containing protein kinase RIP1 and adaptor proteins in the TNF-α/TNFRI pathway, and is the most important anti-apoptotic protein involved in diseases such as Crohn's disease[17,18] and glioblastoma[19]. A20 inhibits TNF-α-induced apoptosis by disrupting recruitment of TRADD and RIP1 to the TNFR1 complex[19]. It has been reported that A20IEC-KO mice are normal, but they are more likely to suffer from intestinal injury induced by intraperitoneal TNF-α injection[19,20]. Furthermore, clinical studies show that the mucosal expression of A20 was significantly lower in Crohn's disease patients compared to healthy controls[21,22]. Taken together, these findings indicate that A20 expression levels are critical in maintaining epithelial barrier function, which may provide a molecular mechanism for illustrating apoptosis in the development of Crohn's disease.

Despite the effectiveness of Western medications, such as aminosalicylates and thiopurines, in inducing and maintaining remission in Crohn's disease, they are limited by their serious side effects, such as nausea, bone marrow suppression, hepatitis, allergic reaction, pancreatitis, and infections and opportunistic infections[20-24]. Approximately 10%-26% of patients withdraw from these treatments because of the adverse effects[21]. Thus, a safer therapy is necessary for managing the disease. Moxibustion, a traditional Chinese medicine, has been demonstrated to be an effective and safe method in treating mild and moderate active-phase Crohn's disease with long-term clinical efficacy[25-28]. After 12 wk of herbs-partitioned moxibustion (HPM) therapy, 74% of 46 mild and moderate Crohn’s disease patients [Crohn’s disease Activity Index (CDAI) from 151 to 350] entered remission periods (CDAI scores < 150). In addition, moxibustion can effectively relieve symptoms such as abdominal pain and diarrhea and can increase hemoglobin counts and decrease C-reactive protein levels[29]. No adverse events were reported in these studies. We previously demonstrated that HPM improved intestinal epithelial barrier function by downregulating the apoptosis of intestinal epithelial cells[29]. However, whether HPM regulates A20 expression to downregulate intestinal epithelial cell apoptosis in Crohn’s disease is obscure. Thus, we used HPM and the most frequently prescribed medication, mesalazine (an aminosalicylate)[30], in both A20IEC-KO and wild-type (WT) Crohn’s disease mice to evaluate the efficacy of HPM in upregulating A20 expression in an apoptotic pathway induced by TNF-α/TNFRI.

MATERIALS AND METHODS

Animals
Eight-week-old A20IEC-KO and WT C57BL/6 mice were obtained from the Shanghai Model Organisms Center (Shanghai, China). The mice were housed at the animal care center of the Shanghai University of Traditional Chinese Medicine and were provided with humane care in a temperature-controlled room (temperature of 22 ± 1 °C and humidity 50% ± 70%) under a 12-h light–dark cycle with free access to food and water. All animal experiments in this study were performed under guidelines approved by the Animal Ethics Committee of the Shanghai University of Traditional Chinese Medicine (No. 2013025).

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Establishment of a mouse model of Crohn's disease

Forty-eight C57BL/6 WT and A20IEC-KO mice each were randomly divided into normal control (NC, \( n = 12 \)), model control (MC, \( n = 12 \)), mesalazine (MESA, \( n = 12 \)), and HPM (\( n = 12 \)) groups. The MC, MESA, and HPM groups were administered with 2,4,6-trinitrobenzene sulfonic acid (TNBS) enemas to establish an experimental Crohn's disease model. The enema solution was prepared with absolute ethanol and 5% TNBS at a 1:1 proportion. The solution was stored away from light. Mice were provided access to water only for 24 h prior to TNBS administration and were weighed. Mice were then anesthetized with 0.05 mL/10 g of 1% pentobarbital sodium via intraperitoneal injection. All mice apart from NC group mice were administered TNBS/ethanol (0.06 mL/10 g of TNBS + 50% ethanol 0.25 mL) intra-anally via a rubber tube, and the solution was retained in the gut cavity at a depth of 3-4 cm. NC mice were administered with physiological saline at 0.05 mL/10 g. All mice were fixed in a handstand posture for 2 min after the rubber tube was withdrawn to prevent outflow of solution. This procedure was performed once. Two mice were randomly selected from each group and sacrificed to confirm the presence of Crohn's disease-like intestinal pathology by hematoxylin and eosin stain (H&E) staining after 4 d.

Treatment methods

Upon confirmation of the model establishment, HPM group mice were treated with HPM. Moxa cones (0.5 cm in diameter and 0.3 cm high) made of refined mugwort floss were placed on herbal cakes [e.g., medicinal formula dispensing (radix) Aconiti praeparata, (cortex) Cinnamomi] at Tianshu (ST25) and Qihai (CV6) acupuncture points (which regulate intestinal functions) and ignited. Two moxa cones were used per treatment, which was administered once daily for 10 d. MESA group mice were fed MESA, which was prepared at a proportion of 1:0.0026, twice daily for 10 d. Mice in the MC and NC groups did not undergo any treatment.

Histological observation

All mice were sacrificed simultaneously at the conclusion of treatment. Approximately 4 cm of colon lesions were resected at a 3-4 cm distance from the anus. A 1 cm length of the dissected colon was removed, washed with iced saline, fixed in 10% natural buffered formalin solution, embedded in paraffin, cut into tissue sections, and stained with H&E. Obtained images were observed under a light microscope (Olympus, Tokyo, Japan).

Enzyme-linked immunosorbent assay

A 96-well commercial kit (MyBioSource, Inc. San Diego, CA, United States) was applied to evaluate serum endotoxin levels. Blood samples were centrifuged at 3000 \( \times \) g for 10 min. Diluted serum sample (1:10), endotoxin test water, and TAL were added to the plate and incubated for 10 min at 37 °C. Then, chromogenic matrix solution was added to the plate and incubated for 6 min at 37 °C. After that, azo reagent was added to the plate and incubated for 5 min at 37 °C. Optical densities were detected with a plate reader (BioTek Instruments, Winooski, VT, United States).

Western blot analysis

Protein (60 µg) extracted from isolated rat intestinal epithelial cell samples was separated by SDS-PAGE and transferred to polyvinylidene difluoride (PVDF) membranes. The membranes were then blocked with 5% skimmed milk in TBS-T for 1 h at room temperature and incubated with the following primary antibodies overnight at 4 °C: A20 (1:1000; ab13597, Abcam), TNF-α (1:1000; ab6671, Abcam), TNFR1 (1:5000; ab19139, Abcam), TRADD (1:500; ab110644, Abcam), RIP1 (1:500; ab72139, Abcam), FADD (1:400; ab24533, Abcam), and GAPDH (1:1500; 5174, CST). Following several sequential washes, the membranes were incubated with the corresponding secondary anti-mouse antibody (A0208, Beyotime) for 1 h at room temperature. Blots were then washed four times with TBS-T (10 min each time). The membranes were stained with ECL enhanced chemiluminescence solution and visualized using a visualizer.

Terminal dUTP nick-end labeling assay

A terminal dUTP nick-end labeling (TUNEL) kit purchased from MyBioSource was utilized to evaluate apoptosis. Paraffin sections were first fully deparaffinized and hydrated, treated with protease K solution (20 µg/mL) for 15 min at room temperature, and subsequently washed and immersed in 3% hydrogen peroxide for 15 min. The sections were then immersed in an equilibrium buffer for 20 min at room temperature and afterward incubated with TdTase for 60 min at 37 °C. After that, the sections were stained with 3,3′-diaminobenzidine (DAB, Shanghai Long Island Biotec).
Co., Ltd, China) and counterstained with hematoxylin. Apoptotic cells as well as the total number of cells were calculated.

**Double immunofluorescence staining**

Paraffin sections were fully deparaffinized and hydrated, and then washed and heated to 92-98 °C for antigen retrieval. Samples were incubated with primary antibodies against A20 (ab13597, Abcam), RIP1 (ab72139, Abcam), and TRADD (ab110644, Abcam) in blocking buffer overnight at 4 °C. Samples were subsequently incubated with the corresponding secondary antibody in blocking buffer at room temperature, and finally incubated with DAPI staining solution for 10 min. Images were obtained under a fluorescence microscope (Nikon, Japan).

**Statistical analysis**

Experimental data were analyzed with SPSS 21.0 software (SPSS Inc., Wacker Drive, Chicago, IL, United States) and GraphPad Prism 5 (GraphPad Software, San Diego, CA, United States). All data are presented as the mean ± standard deviation (SD). Statistics among each experimental group were analyzed using one-way analysis of variance (ANOVA) followed by the least significant difference test. The level of significance was set at \( p < 0.05 \) and \( p < 0.01 \). Variation in epithelial permeability across groups

**RESULTS**

**Intestinal morphological observations in each group**

Previous studies have shown that the Crohn’s disease model of A20IEC-KO mice showed a more severely damaged mucosa than WT mice[12]. In this study, by histopathological evaluation, we found that in WT NC mice, intact mucosal epithelial cells and normal morphological changes affecting the submucosa and muscularis were observed (Figure 1A). In WT MC mice, sparse goblet cells, fibrous hyperplasia, as well as damage to mucosal glands, vasodilation, and hyperemia were noted in the mucosa, and hyperemia and edema were observed under the submucosa (Figure 1B). In WT MESA mice, sparse goblet cells, infiltration of inflammatory cells into the mucosa and submucosa, as well as hyperemia and edema under the submucosa were noted. No obvious abnormal changes were observed in the structural morphology of mucosal, submucosal, and muscularis layers (Figure 1C). In WT HPM mice, sparse goblet cells, mild infiltration of inflammatory cells into the colonic mucosa and submucosa, sparse fibroblast hyperplasia, and healing ulcers were observed. No obvious abnormal changes were observed in the structural morphology of the mucosal, submucosal, or muscularis layer (Figure 1D).

In A20IEC-KO NC mice, intact mucosal epithelial cells and normal submucosal and muscularis structural morphology were observed (Figure 1E). In A20IEC-KO MC mice, partial epithelial cells loss, mucosal goblet cell depletion, inflammatory cell infiltration, glandular damage, and proliferation of fibrous tissue were observed in the mucosa in addition to erosion and necrosis of the mucosal surface. Hyperemia and edema were observed in both mucosal and submucosal layers (Figure 1F). In A20IEC-KO MESA mice, sparse goblet cells, small healing ulcers with associated glandular destruction, and massive inflammatory cell infiltration were observed in the mucosa. No obviously abnormal structural changes in the mucosal, submucosal, or muscularis layer were observed (Figure 1G). In A20IEC-KO HPM mice, sparse goblet cells, inflammatory cell infiltration, and proliferation of fibrous tissue were observed in mucosal tissue. No obviously abnormal structural changes in the mucosal, submucosal, or muscularis layer were noted (Figure 1H).

**Variations in epithelial permeability across groups**

Next, we observed the intestinal epithelial barrier permeability by detecting serum endotoxin levels(Figure 2). In WT groups, as compared with NC mice, serum endotoxin levels were upregulated in the MC, MESA, and HPM groups (\( P_{\text{MC}} < 0.01, P_{\text{MESA}} < 0.01, P_{\text{HPM}} < 0.01 \)). Serum endotoxin levels were downregulated in HPM (\( P < 0.01 \)) and MESA (\( P < 0.05 \)) mice as compared to MC mice. Serum endotoxin levels were significantly decreased in HPM mice (\( P < 0.01 \)) as compared to MESA mice.

In mice from A20IEC-KO groups, as compared with NC mice, serum endotoxin levels were upregulated in the MC, MESA, and HPM groups (\( P_{\text{MC}} < 0.01, P_{\text{MESA}} < 0.01, P_{\text{HPM}} < 0.01 \)). They were downregulated in MESA and HPM mice as compared to MC mice (\( P_{\text{MESA}} < 0.01, P_{\text{HPM}} < 0.01 \)). Serum endotoxin levels were significantly decreased in HPM mice (\( P < 0.01 \)) as compared to those of the MESA group.

Compared with WT NC mice, no significant difference in serum endotoxin levels were noted in NC A20IEC-KO mice (\( P > 0.05 \)). Compared with WT MC mice, endotoxin
levels in MC A20\(^{IEC-KO}\) mice were upregulated (\(P < 0.05\)). Compared with WT HPM mice, serum endotoxin levels were upregulated in HPM A20\(^{IEC-KO}\) mice (\(P < 0.01\)).

**Observation of percentage of apoptotic intestinal epithelial cells in each group**

Since A20 protein is involved in epithelial barrier function by its anti-apoptotic role in Crohn’s disease, we observed cell apoptosis percentage in different groups by TUNEL method. In WT groups, as compared with NC mice, apoptosis percentages were significantly increased in the MC, MESA, and HPM groups (\(P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01\)). Apoptosis percentages were significantly decreased in MESA and HPM mice (\(P_{MESA} < 0.01, P_{HPM} < 0.01\)) as compared to those of the MC group. No significant difference in apoptosis percentages was noted in HPM mice (\(P > 0.05\)) (Figure 3A-D, I) as compared to those of the MESA group.

In A20\(^{IEC-KO}\) groups, as compared with NC mice, apoptosis percentages were significantly increased in the MC, MESA, and HPM groups (\(P_{NC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01\)). Apoptosis percentages were significantly decreased in the MESA and HPM groups (\(P_{MESA} < 0.01, P_{HPM} < 0.01\)) as compared to the MC group. There was no difference between the HPM and MESA groups (\(P > 0.05\)) (Figure 3E-I).

Compared with each corresponding group of WT mice, apoptosis percentages were significantly increased in the NC, MC, MESA, and HPM groups of A20\(^{IEC-KO}\) mice (\(P_{NC} < 0.01, P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01\)) (Figure 3A-I).

**Expression of members of the TNF-α/TNFR1-TRADD-FADD apoptotic pathway in the intestinal epithelium across groups**

A20 expression across groups: In WT groups, as compared with the NC group, A20 levels were decreased in MC, MESA (\(P_{MC} < 0.01, P_{MESA} < 0.01\)), and HPM mice (\(P_{HPM} < 0.05\)). Compared with MC mice, A20 expression was significantly increased in HPM mice (\(P < 0.01\)) and increased in MESA mice (\(P < 0.05\)). Compared with MESA mice, no difference in A20 levels was noted in HPM mice (\(P > 0.05\)) (Figure 4A). No significant differences in A20 expression was noted among A20\(^{IEC-KO}\) groups (\(P_{NC} > 0.05, P_{MC} > 0.05, P_{MESA} > 0.05, P_{HPM} > 0.05\)) (Figure 4A).

Compared with each corresponding group of WT mice, expression levels were significantly increased in all A20\(^{IEC-KO}\) groups (\(P_{NC} < 0.01, P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01\)) (Figure 4A).

TNF-α expression across groups: In WT groups, as compared with NC mice, TNF-α levels were significantly increased in MC, MESA, and HPM mice (\(P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01\)). Compared with MC mice, they were significantly decreased in MESA and HPM mice (\(P_{MESA} < 0.01, P_{HPM} < 0.01\)). No differences in TNF-α expression in HPM mice were found as compared to those of the MESA group (\(P > 0.05\)) (Figure 4B).

In A20\(^{IEC-KO}\) groups, TNF-α levels were significantly increased in the MC, MESA, and HPM groups as compared to those in NC mice (\(P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01\)). TNF-α levels were significantly decreased in MESA and HPM mice (\(P_{MESA} < 0.01, P_{HPM} < 0.01\)) as compared to MC mice. TNF-α levels were decreased in HPM mice as
RIP1 expression across groups: Compared with WT NC mice, no difference in RIP1 expression in those of the HPM group (P > 0.05). No difference in RIP1 expression in those of the HPM group was noted compared to those of the MESA group (P = 0.05) (Figure 4E). Compared with each corresponding group of WT mice, TNF-α expression was significantly increased in A20IEC-KO mice (P < 0.05); TNF-α expression was increased in MC mice (P < 0.05) and significantly increased in MESA mice (P < 0.01) (Figure 4B).

TRADD expression across groups: In WT groups, as compared with NC mice, TRADD expression was significantly increased in MC, MESA, and HPM mice (P<0.01, P<0.01, P<0.05). Compared with MC mice, TRADD expression was significantly decreased in mice of the MESA and HPM groups (P<0.01, P<0.01). No difference in TRADD expression was found in HPM mice as compared to those of the MESA group (P > 0.05) (Figure 4C). In the A20IEC-KO groups, as compared with NC mice, TRADD expression was significantly increased in MC and HPM mice (P<0.01, P<0.01). No difference in TRADD expression in mice of the HPM group was noted as compared to that in the MESA group (P > 0.05) (Figure 4C). Compared with the corresponding WT MC and WT HPM groups, TRADD expression was increased in A20IEC-KO mice (PMC < 0.05, PHPM < 0.05). No difference in TRADD expression among corresponding WT and A20IEC-KO NC and MESA groups was noted (PNC > 0.05, PMESA > 0.05) (Figure 4C).

RIP1 expression across groups: In WT groups, compared with NC group mice, RIP1 levels were significantly increased in the MC, MESA, and HPM groups (P<0.01, P<0.01, P<0.01). Compared with MC group mice, RIP1 levels were decreased in those of the HPM group (P < 0.05); no difference in RIP1 levels (P > 0.05) was noted in MESA group mice. Compared with MESA group mice, no difference in RIP1 levels was noted in those of the HPM group (P > 0.05) (Figure 4E). In mice of A20IEC-KO groups, compared with those of the NC group, RIP1 levels were significantly increased in MC, MESA, and HPM mice (P<0.01, P<0.01, P<0.01). No differences in RIP1 levels among the MC, MESA, and HPM groups were noted (P > 0.05) (Figure 4E). Compared with WT NC mice, no difference in RIP1 expression in compared to those of the MESA group (P < 0.05) (Figure 4B). Compared with each corresponding group of WT mice, TNF-α levels in the A20IEC-KO NC and HPM groups were different (P > 0.05); TNF-α expression was increased in MC mice (P < 0.05) and significantly increased in MESA mice (P < 0.01) (Figure 4B).

TNFR1 expression across groups: In WT groups, as compared with NC mice, TNFR1 expression was significantly increased in MC, MESA, and HPM mice (P<0.01, P<0.01, P<0.05). Compared with MC mice, TNFR1 expression was significantly decreased in mice of the MESA and HPM groups (P<0.01, P<0.01). No difference in TNFR1 expression was found in HPM mice as compared to those of the MESA group (P > 0.05) (Figure 4C). In the A20IEC-KO groups, as compared with NC mice, TNFR1 expression was significantly increased in MC and HPM mice (P<0.01, P<0.01). It was increased in MESA mice as well (P<0.05). Compared with MC mice, TNFR1 expression was significantly decreased in mice of the MESA and HPM groups (P<0.01, P<0.01). No difference in TNFR1 expression in mice of the HPM group was noted as compared to that in the MESA group (P > 0.05) (Figure 4C). Compared with the corresponding WT MC and WT HPM groups, TNFR1 expression was increased in A20IEC-KO mice (PMC < 0.05, PHPM < 0.05). No difference in TNFR1 expression among corresponding WT and A20IEC-KO NC and MESA groups was noted (PNC > 0.05, PMESA > 0.05) (Figure 4C).
Figure 3: Apoptosis percentages of intestinal epithelial cells across groups (magnification, ×200). A: Wild-type mice in normal control group; B: Wild-type mice in model control group; C: Wild-type mice in mesalazine group; D: Wild-type mice in herbs-partitioned moxibustion; E: A20IEC-KO mice in normal control group; F: A20IEC-KO mice in model control group; G: A20IEC-KO mice in mesalazine group; H: A20IEC-KO mice in herbs-partitioned moxibustion group; I: Percentage of apoptotic cells. Data are presented as the mean ± standard deviation (n = 10). Data were evaluated for statistical significance by one-way analysis of variance and are represented as follows: *P < 0.05, **P < 0.01 as compared to normal control; *P < 0.05, **P < 0.01 as compared to model control; *P < 0.05, **P < 0.01 as compared to mesalazine; *P < 0.05, **P < 0.01 as compared to wild type. WT: Wild type; NC: Normal control; MC: Model control; MESA: Mesalazine; HPM: Herbs-partitioned moxibustion.

A20IEC-KO NC mice (P > 0.05) was found. Compared with WT HPM mice, expression of RIP1 was increased in A20IEC-KO HPM mice (P < 0.01). Compared with corresponding WT MC and MESA mice, RIP1 expression was increased in A20IEC-KO mice (P_{MC} < 0.05, P_{MESA} < 0.05) (Figure 4E).

FADD expression across groups: In WT groups, compared with NC mice, FADD levels were significantly increased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). Compared with MC mice, FADD expression was decreased in MESA and HPM mice (P_{MESA} < 0.01, P_{HPM} < 0.01). Compared with mice of the MESA group, no difference in FADD expression levels was found in HPM mice (P > 0.05) (Figure 4F). In the A20IEC-KO groups, compared with NC mice, FADD levels were significantly increased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). No difference in FADD expression was noted among MC, MESA, and HPM mice (P > 0.05) (Figure 4F). Compared with WT NC and MC mice, no difference in FADD levels was found between A20IEC-KO NC and MC mice (P_{NC} > 0.05, P_{MC} > 0.05). Compared with WT HPM mice, FADD levels were significantly increased in A20IEC-KO HPM mice (P < 0.01). Compared with WT MESA mice, FADD levels were increased in A20IEC-KO MESA mice (P < 0.05) (Figure 4F).

Co-expression of A20/TRADD and A20/RIP1 in intestinal epithelial tissues across groups

Co-expression of A20/TRADD: Cell nuclei stained blue. Regions rich in A20 expression stained green while those rich in TRADD stained red. Co-expression of A20 and TRADD stained yellow (red and green fluorescence). In WT NC mice, green fluorescence predominated (Figure 5A). In WT MC mice, red fluorescence predominated along with sparse yellow staining (Figure 5B). In WT MESA mice,
Figure 4 Expression levels of A20, tumor necrosis factor alpha, tumor necrosis factor receptor 1, tumor necrosis factor receptor 1-associated death domain, receptor-interacting protein 1, and FAS-associated death domain protein across groups. Data are presented as the mean ± standard deviation (n = 10). Data were evaluated for statistical significance using one-way analysis of variance and are represented as follows: *P < 0.05, **P < 0.01 as compared to normal control; †P < 0.05, ‡P < 0.01 as compared to model control; §P < 0.05, ¶P < 0.01 as compared to mesalazine; ‡‡P < 0.05, ‡‡‡P < 0.01 as compared to wild type. TNFR1: Tumor necrosis factor receptor 1; RIP1: Receptor-interacting protein 1; TNF-α: Tumor necrosis factor alpha; TRADD: Tumor necrosis factor receptor 1-associated death domain; FADD: FAS-associated death domain; WT: Wild type; NC: Normal control; MC: Model control; MESA: Mesalazine; HPM: Herbs-partitioned moxibustion.

Sparse yellow fluorescence predominated (Figure 5C). In WT HPM mice, immunofluorescence mainly revealed yellow fluorescence (Figure 5D). In A20IEC-KO NC, MC, MESA, and HPM mice, red fluorescence predominated (Figure 5E-H). In WT groups, compared with NC mice, A20 levels were significantly decreased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). Compared with MC mice, expression of A20 was significantly increased in HPM mice (P < 0.01) and
Figure 5 Co-expression of A20/tumor necrosis factor receptor 1-associated death domain in the intestinal epithelium of mice across groups. A: Wild-type mice in normal control group; B: Wild-type mice in model control group; C: Wild-type mice in mesalazine group; D: Wild-type mice in herbs-partitioned moxibustion group; E: A20$^{IEC-KO}$ mice in normal control group; F: A20$^{IEC-KO}$ mice in model control group; G: A20$^{IEC-KO}$ mice in mesalazine group; H: A20$^{IEC-KO}$ mice in herbs-partitioned moxibustion group. Data are presented as the mean ± standard deviation (n = 10). Data were evaluated for statistical significance using one-way analysis of variance and are represented as follows: *P < 0.05, **P < 0.01 as compared to normal control; *P < 0.05, **P < 0.01 as compared to model control; *P < 0.05, **P < 0.01 as compared to mesalazine; *P < 0.05, **P < 0.01 as compared to wild type. TRADD: Tumor necrosis factor receptor 1-associated death domain; WT: Wild type; NC: Normal control; MC: Model control; MESA: Mesalazine; HPM: Herbs-partitioned moxibustion.

Increased in those of the MESA group (P < 0.05). Compared with MESA mice, no difference in A20 levels in those of HPM mice was noted (P > 0.05) (Figure 5I). In A20$^{IEC-KO}$ mice, no difference of A20 levels was noted among each group (P > 0.05) (Figure 5I). Compared with each corresponding group of WT mice, expression of A20 was significantly decreased in A20$^{IEC-KO}$ mice (P < 0.01) (Figure 5I).

In WT groups, compared with NC mice, TRADD levels were significantly increased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). Compared with MC mice, TRADD levels were decreased in mice of the MESA and HPM groups (P_{MESA} < 0.05, P_{HPM} < 0.05). Compared with MESA mice, no difference in TRADD levels was found in HPM mice (P > 0.05) (Figure 5I). In A20$^{IEC-KO}$ groups, compared with NC mice, TRADD levels were significantly increased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). No difference in TRADD levels were noted among MC, MESA, and HPM mice (P > 0.05) (Figure 5I). Compared with WT MC mice, TRADD levels were increased in A20$^{IEC-KO}$ MC mice (P < 0.05). Compared with WT HPM and MESA mice, TRADD levels were significantly increased in mice of corresponding A20$^{IEC-KO}$ groups (P_{HPM} < 0.01, P_{MESA} < 0.01) (Figure 5I).

Co-expression of A20/RIP1: Figure 6A-H shows that the nuclei exhibited blue fluorescence while regions rich in A20 and RIP1 stained green and red, respectively. Regions co-expressing A20 and RIP1 mainly stained yellow. In WT NC mice, imaged regions mainly stained green, occasionally intermixed with yellow fluorescence (Figure 6A). In WT MC mice, imaged regions stained mainly red and yellow (Figure 6B). In WT MESA mice, imaged regions mainly stained green (Figure 6C). In WT HPM mice, imaged regions mainly stained yellow (Figure 6D). In A20$^{IEC-KO}$ NC mice, sparse red and green fluorescence was apparent (Figure 6E). In A20$^{IEC-KO}$ MC, MESA, and HPM group mice, red fluorescence predominated (Figure 6F-H).
Figure 6 Co-expression of A20/receptor-interacting protein 1 in the intestinal epithelium of mice across groups. A: Wild-type mice in normal control group; B: Wild-type mice in model control group; C: Wild-type mice in mesalazine group; D: Wild-type mice in herbs-partitioned moxibustion group; E: A20IEC-KO mice in normal control group; F: A20IEC-KO mice in model control group; G: A20IEC-KO mice in mesalazine group; H: A20IEC-KO mice in herbs-partitioned moxibustion group. Data are presented as the mean ± standard deviation (n = 10). Data were evaluated for statistical significance using one-way analysis of variance and are represented as follows: *P < 0.05, **P < 0.01 as compared to normal control; †P < 0.05, ††P < 0.01 as compared to model control; ‡P < 0.05, ‡‡P < 0.01 as compared to mesalazine; §P < 0.05, §§P < 0.01 as compared to wild type. RIP1: Receptor-interacting protein 1; WT: Wild type; NC: Normal control; MC: Model control; MESA: Mesalazine; HPM: Herbs-partitioned moxibustion.

In WT groups, compared with NC mice, A20 expression was significantly decreased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). A20 levels were significantly increased in HPM and MESA mice (P_{HPM} < 0.01, P_{MESA} < 0.01) as compared to MC mice. Compared with MESA mice, no difference in levels of A20 were found in HPM mice (P > 0.05). In A20IEC-KO groups, no differences in A20 levels were found among groups (P > 0.05). Compared with each corresponding group of WT mice, A20 levels were significantly decreased in A20IEC-KO NC, MESA, and HPM mice (P_{NC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01) and decreased in MC mice (P_{MC} < 0.05) (Figure 6I).

In WT groups, compared with NC mice, RIP1 levels were significantly increased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). Compared with MC mice, RIP1 levels were significantly increased in MESA mice (P < 0.01) and decreased in HPM mice (P < 0.05). Compared with MESA mice, no difference in RIP1 levels were noted in HPM mice (P > 0.05). In A20IEC-KO groups, compared with NC mice, RIP1 levels were significantly increased in MC, MESA, and HPM mice (P_{MC} < 0.01, P_{MESA} < 0.01, P_{HPM} < 0.01). No difference in RIP1 levels was found among MC, MESA, and HPM mice (P > 0.05) (Figure 6I). Compared with WT NC mice, no difference in RIP1 levels was found in A20IEC-KO NC mice (P > 0.05). Compared with WT MC mice, RIP1 expression was increased in A20IEC-KO MC mice (P < 0.05). Compared with WT MESA mice, RIP1 expression was significantly increased in mice of the corresponding A20IEC-KO group (P_{MESA} < 0.01). Compared with WT HPM mice, RIP1 levels were increased in A20IEC-KO HPM mice (P_{HPM} < 0.05) (Figure 6I).

DISCUSSION
Although the etiology of Crohn’s disease is still unknown, excessive apoptosis of intestinal epithelial cells leads to villus atrophy and epithelial destruction, which plays a central role in the pathogenesis of the disease[29,30]. A20, as an intestinal epithelial protector, is widely known for its functions in maintaining the epithelial barrier stability in inflammatory conditions. In Crohn’s disease patients, there is an excessive inflammatory response and insufficient upregulation of A20 expression[31]. Our previous studies have indicated that HPM plays a beneficial role in Crohn’s disease by decreasing intestinal epithelium apoptosis[32]. However, whether the effect of HPM is through A20 has not been determined. In the present study, we examined the anti-apoptotic properties of A20, confirmed its protective role in the intestinal epithelial barrier, and explored whether the effect of HPM in reducing intestinal epithelium apoptosis is through upregulating A20 levels in apoptotic signaling in a TNBS-induced Crohn’s disease mouse model in WT and A20IEC-KO lineages.

Aberrant apoptosis of intestinal epithelial cells lacking the A20 gene leads to increased intestinal epithelial permeability in Crohn’s disease patients[33-35]. In the present study, we found that the intestinal epithelial cell apoptosis percentage was significantly increased in the Crohn’s disease model of A20IEC-KO mice (P < 0.01); moreover, serum endotoxin level was upregulated in this lineage (P < 0.05). Accordingly, H&E staining analysis showed that the intestinal epithelial barrier was severely damaged in the Crohn’s disease model of A20IEC-KO mice, consistent with the Lars Vereecke’s report[36]. A previous study showed that HPM reduces intestinal epithelial apoptosis in a Crohn’s disease mouse model[19]. The present study indicated that after treatment with HPM, despite improved intestinal morphological changes with decreased intestinal epithelial cell apoptosis percentages and endotoxin levels in A20IEC-KO mice, a more significant improvement was detected in WT mice compared with A20IEC-KO mice (P < 0.01). These findings suggested that upregulated A20 can protect intestinal epithelial barrier function and initially confirmed that HPM can upregulate A20 expression to protect the intestinal epithelial barrier in Crohn’s disease.

The anti-apoptotic function of A20 has been found to inhibit the sequential signaling complexes of the TNF-α/TNFFR1 apoptotic pathway upstream of caspase 8[38]. When TNFR1 binds to TNF-α, the death domain of TNFR1 enables the recruitment of TRADD and RIP1 proteins and their assembly with FADD to activate caspase-8 and induce apoptosis[39-41]. Therefore, we measured the protein expression levels of TNF-α, TNFR1, TRADD, RIP1, and FADD in the apoptotic pathway. The present study found no differences in TNF-α, TNFR1, TRADD, RIP1, or FADD levels among A20IEC-KO NC mice, revealing that A20IEC-KO NC intestinal epithelial cells do not spontaneously undergo apoptosis, in agreement with prior research by Lars Vereecke et al[36]. We previously reported that HPM and mild moxibustion downregulate TNF-α and TNFR1 expression levels and decrease intestinal epithelial cell apoptosis in Crohn’s disease[19]. Here, we found that when A20 was upregulated in WT mice, TNF-α expression was decreased after HPM treatment in both WT and A20IEC-KO Crohn’s disease mice. Interestingly, TNF-α expression levels were not different in A20IEC-KO and WT HPM mice. These results may indicate that HPM could downregulate TNF-α, which is consistent with our previous study[19], but the mechanism enacted by HPM in regulating TNF-α is not specifically associated with A20. TNFR1, TRADD, and RIP1 levels in A20IEC-KO MC mice were found to be increased compared with those in WT mice, suggesting that A20IEC-KO mice are hypersensitive to TNF-α-induced intestinal epithelial apoptosis, consistent with Vereecke’s research[19]. Western blot analysis revealed that HPM significantly decreased the expression levels of TNFR1, TRADD, RIP1, and FADD in WT mice but had no effect on the levels of those proteins in A20IEC-KO mice. All of these results indicated that although HPM can downregulate the expression levels of TNFR1, TRADD, RIP1, and FADD, the upregulated expression of A20 by HPM can downregulate TNFR1, TRADD, and RIP1 signaling molecules in the apoptotic pathway.

As TRADD and RIP1 play critical roles in the TNFR1-related signaling apoptotic pathway, we further observed the role of A20 in affecting TRADD and RIP1 expression[32]. A study showed that ligand-dependent association of RIP1 with TNFR1 was significantly reduced in A20-expressing cells in TNF-α-induced apoptosis. Furthermore, the recruitment of TRADD to the TNFR1 complex was also inhibited by A20[39]. In the present study, immunofluorescence showed a predominantly green color with a few yellow areas in intestinal epithelial tissue of the WT HPM group. The imaging data revealed co-expression of A20/TRADD and A20/RIP1 in WT HPM mice and decreased expression of TRADD and RIP1 along with increased expression of A20 in WT HPM mice. In A20IEC-KO HPM mice, immunofluorescence showed a predominantly red color in intestinal epithelial tissue, suggesting a significant amount of TRADD and RIP1 expression without A20 expression. These results may identify a close association of A20, TRADD, and RIP1 within the TNF-α/TNFFR1 apoptotic
pathway in a Crohn’s disease mouse model. Our findings indicated that HPM upregulates the A20 level, which may affect the expression levels of TRADD and RIP1 in the apoptotic pathway.

In conclusion, HPM in treating Crohn’s disease functions possibly via upregulation of the A20 expression level, resulting in downregulation of TNFR1, TRADD, and RIP1 to alleviate increased cell apoptosis in the intestinal epithelial barrier in Crohn’s disease.

**ARTICLE HIGHLIGHTS**

**Research background**
A20, as an intestinal epithelium protector, plays a critical role in anti-apoptosis in Crohn’s disease. Previous studies have indicated a beneficial role of herbs-partitioned moxibustion (HPM) in Crohn’s disease by decreasing intestinal epithelial apoptosis. However, whether the effect of HPM is through A20 is unclear.

**Research motivation**
Our findings will suggest a role of HPM in regulating A20 level in anti-apoptotic pathway in the intestinal epithelium of mice with Crohn’s disease.

**Research objectives**
To explore whether HPM alleviates cell apoptosis in the intestinal epithelium by upregulating A20 level in Crohn’s disease.

**Research methods**
Two types of mice were included in this study, namely, mice with A20 deletion in intestinal epithelial cells (A20IEC-KO) and wild-type mice. Both of them were randomly divided into normal control (NC), model control (MC), mesalazine (MESA), and HPM groups. 2,4,6-trinitrobenzene sulfonic acid (TNBS) was administered to establish a Crohn’s disease model in the two types. The morphology of the colonic mucosa, serum endotoxin, apoptosis of epithelial cells, protein levels of A20 and tumor necrosis factor receptor (TNFR) 1-related signaling molecules, co-expression of A20 and TNFR1-associated death domain (TRADD), and co-expression of A20 and receptor-interacting protein (RIP) 1 were observed. All data are presented as the mean ± standard deviation.

**Research results**
Compared with A20IEC-KO mice, wild-type mice in the HPM group showed that damage of intestinal epithelial barrier was improved, serum endotoxin levels were significantly downregulated (P < 0.01), apoptosis percentages were significantly decreased (P < 0.01), A20 level was significantly upregulated (P < 0.01), and TNFR1, TRADD, and RIP1 levels were downregulated (P_{TNF-α} < 0.01, P_{TNFR1} < 0.05, P_{TRADD} < 0.05, P_{RIP1} < 0.05). In addition, the co-expression of A20/TRADD and A20/RIP1 showed a predominant yellow fluorescence in WT HPM mice, while a predominantly red fluorescence was noted in A20IEC-KO HPM mice.

**Research conclusions**
HPM can upregulate A20 level, resulting in decreased expression of TNFR1, TRADD, and RIP1 to alleviate aberrant cell apoptosis in the intestinal epithelial barrier in Crohn’s disease.

**Research perspectives**
Effect of HPM in decreasing cell apoptosis of intestinal epithelial cells is through upregulating A20 level in Crohn’s disease.

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