Impact of the alterations in the interstitial cells of Cajal on intestinal motility in post-infection irritable bowel syndrome

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Abstract. The interstitial cells of Cajal (ICC) are basic components of gastrointestinal motility. However, changes in ICC and their role in post-infection irritable bowel syndrome (PI-IBS) remain to be elucidated. To observe the impact of alterations in the ICC on intestinal motility in a PI-IBS mouse model, female C57BL/6 mice were infected by the oral administration of 400 Trichinella spiralis larvae. The abdominal withdrawal reflex, intestine transportation time (ITT), grain numbers, Bristol scores, wet/dry weights and the percentage water content of the mice feces every 2 h were used to assess changes in the intestinal motor function. The intestines were excised and sectioned for pathological and histochemical examination. These intestines were also used to quantify the protein and mRNA expression of c‑kit. The C57BL/6 mouse can act as a PI-IBS model at day 56 post-infection. Compared with the control mice, the ITT was shorter, the grain numbers, Bristol scores, wet and dry weights, and the percentage water content of the mice feces were higher and the dry weights were unchanged in the PI-IBS mice. The protein and mRNA expression levels of c‑kit were upregulated in the entire PI-IBS mouse intestines. Following immunohistochemical staining, the increased number of c-kit-positive cells were detected predominantly in the submucosa and myenteron. These results suggested that the alterations of the ICC resulted in the changes of the intestinal motility patterns in the PI-IBS mouse models induced by Trichinella spiralis infection, which may be the main mechanism underlying intestinal motility disorders in PI-IBS.

Introduction

Irritable bowel syndrome (IBS) is a highly prevalent functional gastrointestinal disorder characterized by abdominal pain and alterations in bowel habits (1). IBS caused by viral, bacterial and parasitic infection, observed in 3.7-36% of patients, has been commonly referred to as PI-IBS (2). The pathophysiology of PI-IBS remains to be elucidated, however, low-grade inflammation and chronic alteration of the immune system at the molecular level have been shown to target the mucosal secretory function, smooth muscle and enteric nervous fibers (3-5). At present motility disorder is an established typical character in PI-IBS patients, however, alterations in the motility patterns in PI-IBS and the underlying mechanisms remain to be elucidated.

The interstitial cells of Cajal (ICC) are present throughout the gastrointestinal tract, from the esophagus to the anus, with three main functions, including acting as mechanoreceptors, transmitting enteric neuronal signals to the smooth muscle cells and placing slow waves and regulating their propagation (6). According to their location, the ICC are divided into the myenteric plexus (MP) ICC, intramuscular (IM) ICC, deep muscular plexus (DMP) ICC and the submucosa (SM) ICC (6). All the subtypes are important components of gastrointestinal motility. Although certain studies have reported that ICC changed in PI-IBS patients and animal models (7,8), the correlation between the patterns of gastrointestinal motor dysfunction and changes of the ICC remain to be elucidated. Therefore, the present study measured the feces and the c-kit protein in the different intestinal segments in PI-IBS model mice to elucidate the types of motility disorders and the association with changes in the ICC.

Materials and methods

Animals. The experimental procedure was approved by the Animal Welfare Committee of Chongqing Medical University (Chongqing, China). Specifically, pathogen-free female C57BL/6 mice (4-6 weeks old) were purchased from the Animal Center of the Third Military Medical University (Chongqing, China). A total of 34 mice were randomly assigned to a control group (n=17) and a model group (n=17) and maintained in normal 12 h light/dark cycles in a regulated environment (20-25°C). The study was approved by the Ethics Committee of the First Affiliated Hospital of Chongqing Medical University.

Trichinella spiralis infection. The infective muscle larvae were obtained from mice infected with Trichinella spiralis at 30 days. The infected mouse was sacrificed, skinned and...
DM6000B, Germany). The immunohistochemical analysis was performed by boiling the sections in 95% ethanol and antigen retrieval was performed by boiling the sections in citrate buffer (0.01 M; pH 6.0) for 20 min. Following cooling at room temperature for 1 h, the specimens were treated with 5% bovine serum albumin for 30 min at 37°C followed by overnight incubation with 1:200 diluted rabbit anti-mouse c-kit polyclonal antibody (Boster Biotechnology, Ltd., Wuhan, China) at 4°C. Following washing in phosphate-buffered saline, the slides were incubated for 60 min at 37°C with the corresponding secondary biotinylated goat anti-rabbit antibody (Boster Biotechnology, Ltd.) at a 1:300 dilution. Following washing, the slides were incubated with Diaminobenzidine chromogen (Zsbio, Beijing China) for 5 min. A nuclei counterstain was performed using Mayer's hematoxylin (Boster Biotechnology, Ltd., Wuhan, China). The slides were then washed with water and immersed in 75, 85, 90 and 100% ethanol and 100% xylene. The sealed slides were recorded digitally using a Leica Microsystem (DM6000B, Leica, Mannheim, Germany). The immunohistochemical analysis was scored as staining intensity and the proportion of positive cells (12). The scale of staining intensity was as follows: 0, not detected; 1, minimal; 2, mild; 3, moderate and 4, marked. The proportion of positive cells scale was: 0, no staining, 1, c-kit-positive cells in <1% population; 2, c-kit-positive cells in 1-10% population; 3, c-kit-positive cells in 10-50% and 4, c-kit-positive cells in >50% population.

Reverse transcription quantitative polymerase chain reaction (RT-qPCR) mRNA assay. The total RNA in the intestinal mucosa was extracted using TRIzol solution (Takara Bio, Inc., Shiga, Japan). The gene expression of c-kit was assayed by RT-qPCR. The β-actin mRNA level was determined as an internal reference and their levels of expression were quantitated by optical densitometry following electrophoresis on an agarose gel. The primers were designed based on the following complementary DNA sequences: β-actin (470 bp), forward 5'-AGGCTGTGCTGTCCCTGTATG-3' and reverse 5'-GAGGTCTTTTACGGATGTCAACG-3' and c-kit (214 bp), forward 5'-CGACGCACTTTCTTATGAT-3' and reverse 5'-AGGACCTCTAGTCCAGAT-3'. The RT was performed at 37°C for 15 min and 85°C for 5 sec. The PCR cycling condition was 36 cycles at 94°C for 30 sec, 59°C (β-actin) or 56°C (c-kit) for 30 sec and 72°C for 35 sec. The PCR end products were electrophoresed on 4% agarose gel and stained with ethidium bromide. The gray values of the bands were calculated using Quantity One software (Bio-Rad). The relative mRNA expression levels of target genes were normalized to the corresponding internal standard.

Western blotting for protein assay. The frozen sample was fractured using an ultrasonic disrupter (HD3100, Bandelin Electronic, Berlin, Germany), suspended in radioimmunoprecipitation buffer (Takara Bio Inc.) for 20 min and the protein levels in the supernatants were obtained following centrifugation at 12,000 x g for 20 min and quantified using a bicinchoninic acid protein assay kit (Beyotime Institute of Biotechnology, Shanghai, China). Aliquots (60 µg) of the protein were subjected to electrophoresis on a polyacrylamide gel for the c-kit protein. The primary antibodies were goat anti-mouse c-kit polyclonal antibody (Boster Biotechnology, Ltd.) at a dilution of 1:500 and goat anti-mouse β-actin polyclonal antibody (Boster Biotechnology, Ltd.) at a dilution of 1:500.
The \( \beta \)-actin protein was used as an internal standard. The signal was detected using an enhanced chemiluminescence detection system (Bio-Rad). The densities of the bands were assessed using Quantity One software (Bio-Rad). The relative protein levels of the target protein were obtained by correction with the corresponding internal standard.

**Statistical analysis.** Statistical analysis was performed using SPSS 19.0 software (IBM SPSS, Armonk, NY, USA). Data are expressed as the mean ± standard deviation. The independent sample t-test was used to compare the results between the two groups. \( P<0.05 \) was considered to indicate a statistically significant difference.

**Results**

**Pathological examination.** H&E staining (Fig. 1) revealed a marked infiltration by neutrophils in the lamina propria and interstitial edema on day 14 PI. Infiltration and edema gradually reduced between days 14 and 28 PI. At 56 days PI, no obvious inflammatory infiltrate was observed and from this day, the mice were deemed suitable as PI-IBS model mice.

**AWR.** At distention volumes of 0.35 and 0.5 ml, the AWR scores in the model mice were higher compared with those in the controls, which indicated that the visceral sensitivity to colorectal distension was increased in the model mice. No significant differences were observed at volumes of 0.25 or 0.65 ml (Table I).

**Measurement of intestinal motility.** The ITT of the PI-IBS group was markedly shorter compared with the control group. Compared with those of the control group, the grain numbers, wet weights, Bristol scores and percentage liquid contents were significantly higher, however, the dry weights were not different between the two groups (Fig. 2).

<table>
<thead>
<tr>
<th>Group</th>
<th>n</th>
<th>0.25 ml</th>
<th>0.35 ml</th>
<th>0.5 ml</th>
<th>0.65 ml</th>
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<tr>
<td>Model</td>
<td>8</td>
<td>0.50±0.25</td>
<td>2.42±0.24(^a)</td>
<td>3.54±0.31(^a)</td>
<td>3.88±0.35</td>
</tr>
<tr>
<td>Control</td>
<td>8</td>
<td>0.33±0.31</td>
<td>1.83±0.44</td>
<td>2.92±0.43</td>
<td>3.75±0.46</td>
</tr>
</tbody>
</table>

Data are presented as the mean ± standard deviation. \(^a\)\( P<0.01 \) compared with the control group.

Figure 1. Histological analysis of the intestinal tissues. Sections of duodenum, jejunum, ileum and colon from the control and model mice were stained with hematoxylin and eosin on different days (original magnification, x200; \( n=3 \)).
Immunohistochemical staining. The immunolabeling revealed that the c-kit signals were detected predominantly in the submucosa and myenteron (Fig. 3). The staining scores of the entire intestines in the model group were notably higher compared with those of the control group (Fig. 4).

RT-PCR and western blotting assays. The results of the RT-PCR and western blotting assays demonstrated that the mRNA and protein levels of c-kit were significantly higher in the duodenum, jejunum, ileum and colon of the PI-IBS mice compared with the control mice (Figs. 5 and 6).

Discussion

PI-IBS is defined as the acute onset of IBS symptoms (meeting Rome III criteria) that develop when an individual, who has not previously met the Rome criteria, experiences a gastrointestinal infection with two or more of the following characteristics: fever, vomiting, diarrhea or a stool culture positive for an infectious agent (13). The Rome III criteria...
are as follows: Recurrent abdominal pain or discomfort should be present for at least three days per month over the past three months and be associated with ≥2 of: involvement with defecation, onset associated with a change in frequency of stool, onset associated with a change in form of stool. These criteria must be fulfilled over the past three months with symptom onset occurring at least six months previously. Although the mechanism remains to be elucidated, visceral hypersensitivity and intestinal motor dysfunction are reported to be involved (14). The present study used a previously described method to establish a PI-IBS mouse model (15). Through the dynamic observation of intestinal histology and AWR scores, the infected mice were used as PI-IBS mice when the intestine returned to normal. According to the results of intestinal motility, the present study suggested that the frequency and rhythm of intestinal peristalsis altered in the model mice. The frequency and rhythm of bowel movement is completed by ICC, smooth muscle cells and the enteric nervous system in combination and ICC provides a pacemaker signal for coordinated motility patterns. As ICC is important in bowel movement (16), the present study hypothesized that ICC were changed in the PI-IBS model mice infected by *Trichinella spiralis*. To confirm this hypothesis, the c-kit protein was used to determine the number and function of the intestinal ICC, which is an important biological symbol and is expressed in the ICC of the intestinal muscular layer and lamina propria (6). The mucosal inflammation of the gut induced by *Trichinella spiralis* infection resulted in loss of ICC function due to structural injury and loss of c-kit positivity (17). In inflammation subsidence, the rough endoplasmic reticulum and Golgi apparatus of the injured ICC are required to synthesize proteins, including c-kit, to form a new plasma membrane and to recover structure and function (17). The present study found that the protein and mRNA expression of c-kit were significantly higher in the PI-IBS mice compared with the control mice. Overexpressed c-kit is required for development and maintenance of ICC, was also necessary for development of coordinated motility patterns and for the survival of ICC (18,19). In several diseases, an association between macrophages and ICC injury has been identified (20,21). However, in vivo, 5-hydroxytryptamine (5-HT) released from macrophages and an increase in PI-IBS can combine with 5-HT2B receptors, which are expressed on the plasma membrane of ICC and this promotes proliferation of the ICC (22,23). The present study found that the c-kit-positive cells increased in the intestine of the PI-IBS mice.

On resolution of submucosal and myenteric inflammation, the network of ICC, particularly one that involved in connecting ICC to each other and to smooth muscle cells, return to a normal physiological state, whereas the increased ICC in PI-IBS mice may induce intestinal motor disturbances similar to that induced by inflammation (24,25). The present study observed that the signals of c-kit were predominantly detected in the submucosa and myenteron, where the ICC mainly pace slow waves (26). Increased ICC has the potential to place more slow waves and this alteration can accelerate the intestinal muscular contraction and peristaltic propagation (27). In addition, increased cholecystokinin in PI-IBS can bind to cholecystokinin A receptors expressed on ICC and promote gastrointestinal emptying (28,29). Furthermore, an increase in the number of IM-ICC can shorten the intervals between the migratory motor complex (MMC) cycles and decrease the bowel transmitting time (30). These mechanisms mean that the reduced time of the whole gastrointestinal transmitting was closely associated with the ICC changes.

The small intestine is the primary site of nutrient absorption. As the frequency and rhythm of bowel movement increase, the higher osmotic pressure of food residue entering the colon is not conducive to water absorption. However, the reduced contact time between food waste and intestinal wall results in a markedly higher water content and higher Bristol scores of the feces (31). In the present study, the observed increase in feces grains and similar dry weights between the two groups indicated
that the MMC frequency increased and the underlying mechanisms may be associated with an increase in the number of slow waves and short time intervals in the MMC cycles (30). However, an increase in the number of IM-ICC leads to increased nitric oxide transmission and subsequently decreased MMC frequency (15). Whether the association between ICC and nitric oxide transmission was abnormal in PI-IBS requires further investigation.

In conclusion, a PI-IBS model was successfully established in the present study using C57BL/6 mice induced by *Trichinella spiralis*. Alterations in the ICC numbers and c-kit protein were closely associated with changes in the intestinal motor patterns in PI-IBS. However, further studies are required to confirm the association between the ICC and intestinal motility disorders in PI-IBS.

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References