

Redistribution of Tight Junction Proteins During EPEC Infection *In Vivo*

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Abstract—Enteropathogenic *Escherichia coli* (EPEC) is a leading cause of diarrhea among infants. Tight junction plays a vital role in intestinal paracellular permeability by forming physical intercellular barriers in epithelial cells. However, the impact of this enteric pathogen on tight junctions *in vivo* has not been fully investigated. In the present study, the alterations in tight junctions following EPEC infection *in vivo* were investigated. Western blot analysis revealed that the tight junction proteins, occludin and claudin-1, were displaced from tight junction membrane microdomains to Triton X-100 soluble fractions after EPEC infection. Changes in intestinal paracellular permeability were determined using the molecular tracer biotin, which was observed to penetrate the epithelia and extended into the lamina propria, indicating disruption in tight junction barrier function. Our results suggested that redistribution of tight junction proteins plays an important role in the disruption of epithelial barrier function induced by EPEC infection, which may provide new insight into the pathogenesis of diarrhea caused by EPEC.

KEY WORDS: enteropathogenic *E. coli*; tight junction; claudins; occludin; barrier function.

INTRODUCTION

Enteropathogenic *Escherichia coli* (EPEC) infection triggers severe watery diarrhea which can lead to death in infants [1]. However, the pathological mechanisms leading to diarrhea during EPEC infection remain largely unknown. EPEC infection induces a typical intestinal lesion characterized by intimate adherence of bacteria to the epithelium, which is associated with microvilli destruction and reorganization of cytoskeletal actin [2]. The formation of attaching and effacing (A/E) lesions requires type III secretion system, which is encoded on a pathogenicity island called the locus of enterocyte effacement [3]. The bacterial proteins and effector molecules are injected into the host cell

cytoplasm through the avenue formed by the secretion system apparatus [4].

The effectors delivered into the cells can function together to subvert host cellular processes [5, 6]. A great number of cellular responses occur following translocation of effectors into host cells, including induction of actin-rich pedestals underneath adherent bacteria, mitochondrial dysfunction, a weak inflammatory response, and tight junction (TJ) disruption [7–9]. They also induce changes in host cell signaling pathways that act in mediating the diarrheal response to EPEC infection [10].

TJs are the apical-most adhesive junctional complexes in mammalian epithelial cells, which function as permeability barriers by segregating luminal from adluminal compartments [11]. TJs are formed of continuous strands of TJ proteins, consisting of occludin, claudins, and junctional adhesion molecule, all of which encircle the most apical aspect of the lateral plasma membrane [12]. There has been increasing recognition of an association between disrupted intestinal barrier function and the development of diseases including Crohn's disease, ulcerative colitis, and microbial infections [13–15]. It is well established that tight junctions are altered by EPEC *in vitro*. A

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decrease in transepithelial electrical resistance was observed during formation of attaching and effacing lesions, which was accompanied by an appearance of aberrant strands throughout the lateral membrane [16]. In addition to these morphological and physiological changes, EPEC has been shown to change the localization of occludin, zonula occludens-1, and the barrier forming claudin proteins [16, 17]. In addition to TER decline, molecular tracers have also indicated that the barrier is functionally compromised during these infections [18].

Although much work has investigated the effects of EPEC infection on tight junctions in tissue culture systems, there have been few investigations in relevant infection models. To investigate the mechanisms underlying whole animal disease, experiments in relevant infections are essential [19]. Until now, the most frequently used animal model was the murine infection model using the A/E mouse pathogen *Citrobacter rodentium* which could induce watery diarrhea. Similar results as the *in vitro* studies have been shown during animal infections with *C. rodentium*. In these animal models, alteration in TJ proteins localization and disruption of epithelial barrier integrity were revealed [19, 20]. A recent study has claimed that infection of EPEC in C57/BL6 mice caused destruction of enterocyte tight junctions [21].

To gain insight into the mechanism underlying the pathogenesis of EPEC infection, we used the *in vivo* A/E mouse infection model to investigate the effects of EPEC infection on tight junctions. We found that following wild-type EPEC infection, occludin and claudin-1 were displaced from the tight junction membrane microdomains to Triton X-100 soluble fractions. Moreover, tight junctions were functionally impaired.

MATERIALS AND METHODS

Animals

C57BL/6J mice were purchased from the Department of Experimental Animals, Medical College of Beijing University, China. Four- to 6-week-old male mice were used in this study. The animals were acclimatized to the new environment for 1 week. Mice were treated in accordance with the local animal protection legislation and were provided food and water *ad libitum*. Experimental protocols were

approved by the Institutional Animal Care and Use Committee of Nanjing University, and the Principles of Laboratory Animal Care (NIH publication no. 86-23, revised in 1985) were followed.

Mouse Infections

The wild-type EPEC 2348/69 was used in the experiment. Mice were orally infected with 2×10^8 EPEC suspended in 200 μ l of sterile phosphate-buffered saline (PBS). Mice that received only sterile PBS served as control. Infections persisted for 1, 3, or 5 days at which point the mice were sacrificed by cervical dislocation.

Histological Analysis

The colon tissues were fixed in 10% neutrally buffered formalin, stained in hematoxylin and eosin (H&E), sectioned, and examined by light microscopy for the changes induced by EPEC infection.

Immunoblot Analysis

The previously reported method was followed to isolate tight junction membrane microdomains [22]. Briefly, tissue samples were homogenized in lysis buffer (50 mM Tris, 25 mM KCl, 5 mM $MgCl_2 \cdot 6H_2O$, 2 mM EDTA, 40 mM NaF, 4 mM Na_3VO_4 , pH 7.4) containing 1% Triton X-100 and protease inhibitor mixture solution. The resulted lysate was subjected to SDS-PAGE. Proteins were transferred electrophoretically onto PVDF membranes (Bio-Rad, Hercules, CA, USA). The membranes were blocked with 3% BSA in Tris-buffered saline containing 0.1% Tween-20 (TBS-T) for 20 min at room temperature. Membranes were then incubated overnight at 4°C with primary antibodies diluted in TBS-T buffer. Following three 5-min washes with TBS-T buffer, the membranes were incubated with appropriate secondary antibodies for 1 h at room temperature. The hybridized band was detected by ECL kit (Amersham Biosciences, Chalfont St Giles, UK) according to the manufacturer's instructions. Immunoblots were scanned using ChemiDOC XRS instrument (Bio-Rad), and densitometry was performed by Quantity One 1-d analysis software (Bio-Rad).

Assessment of Intestinal Mucosal Permeability

The previously reported method was adopted to assess the intestinal mucosal permeability [19]. In this method, EZ-link Sulfo-NHS-Biotin was used as a

molecular tracer. In brief, tissues were fixed for 3 h in 3% paraformaldehyde and washed three times with PBS prior to cryoembedding and sectioning. Subsequently, tissues were embedded in optimal cutting temperature compound (Sakura Finetech, USA) and serial sections were cut at a thickness of 6 μm . Thereafter, tissues were permeabilized with 0.2% Triton X-100 in PBS for 20 min, and blocking was achieved by incubation with 5% goat serum in PBS containing 0.05%

Tween-20 and 0.1% bovine serum albumin. Incubation with primary antibodies claudin-3 and 5 (Zymed Laboratories Inc., San Francisco, CA, USA) was performed overnight at 4°C. The primary antibodies were used at a dilution of 1:200. After three washes with PBS, the sections were incubated with Alexa 488-conjugated streptavidin (1:500) and Alexa 635-conjugated secondary antibodies (Molecular Probes, Eugene, OR, USA) (at 1:100 dilution) at room temperature for 1 h in the dark.

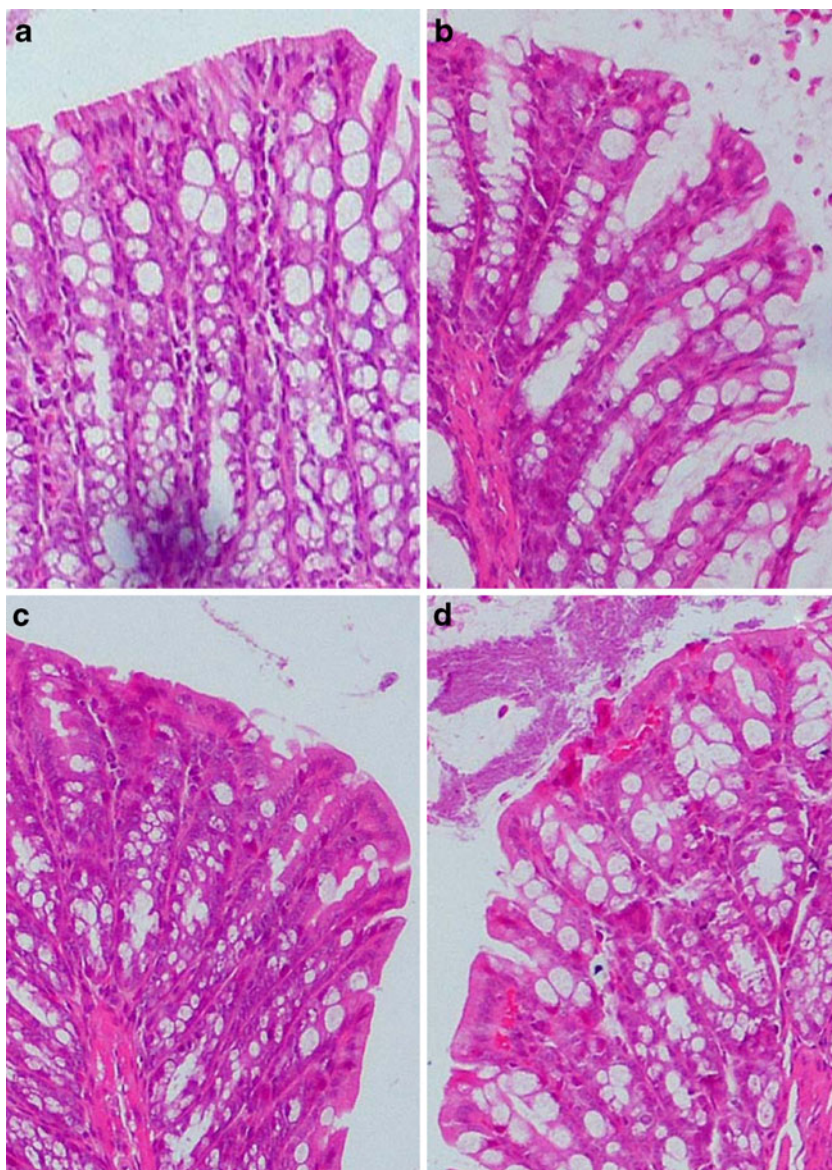


Fig. 1. Histological findings of the colon from control and EPEC-infected mice. **a** Control, showing normal colonic architecture ($\times 100$). **b-d** Mice infected for 1, 3, and 5 days ($\times 100$). Note the distortion of colon epithelium and infiltration of inflammatory cells.

Cell DNA was visualized by 4', 6'-diamidino-2-phenylindole (DAPI; Molecular Probes). The stained sections were visualized and photographed using a Leica TCS SP2 laser confocal scanning microscope (Leica Microsystems, Heidelberg GmbH, Mannheim, Germany).

Statistical Analysis

Data were analyzed using paired Student's *t* test. Data were presented as mean ± SEM and were considered significant if *P* was <0.05.

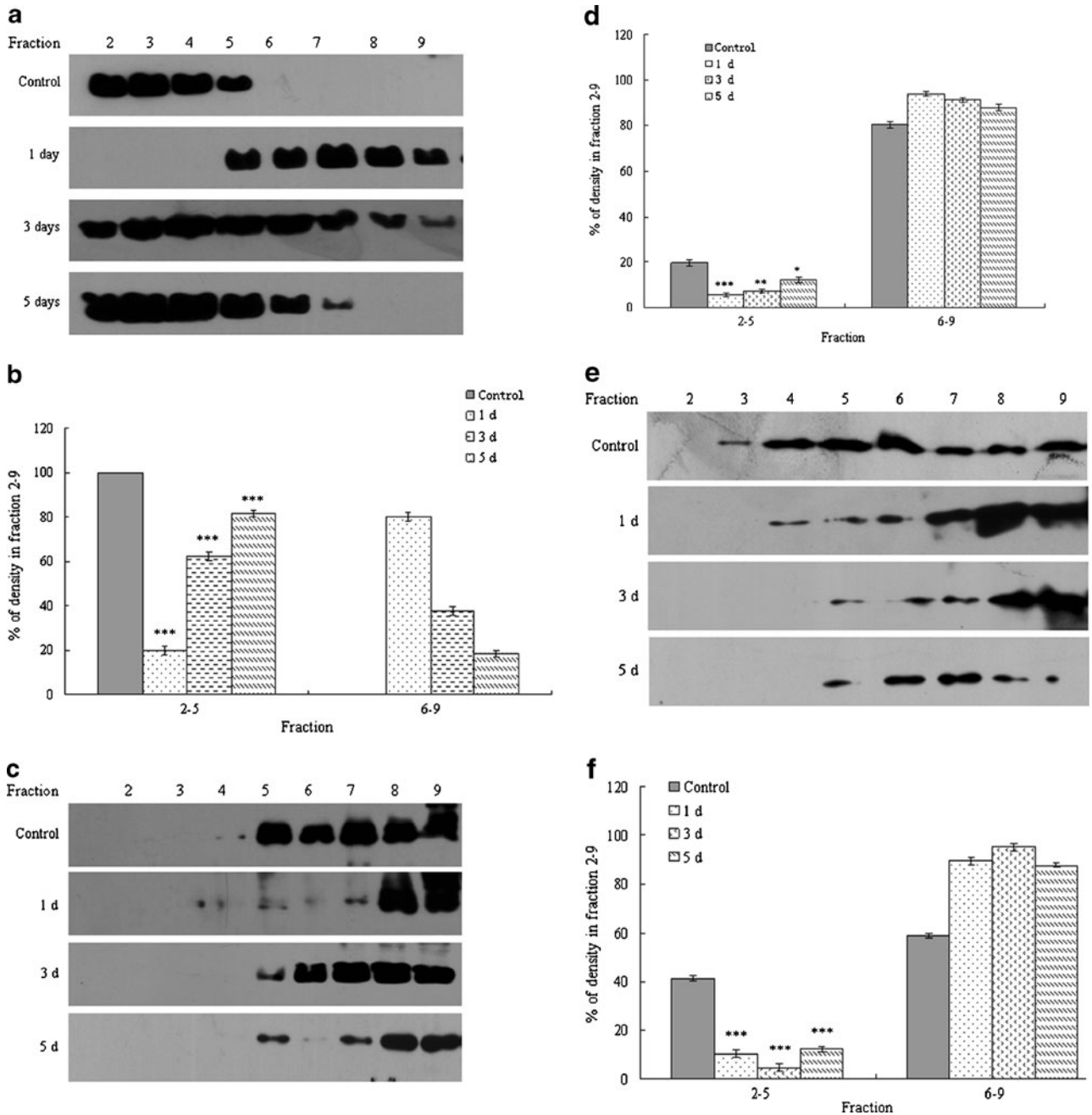


Fig. 2. Effect of EPEC infection on the distribution of occludin and claudin-1 in tight junction membrane microdomains. Tight junction membrane microdomains were isolated by sucrose density gradient centrifugation and analyzed by immunoblotting, probing with antibodies against caveolin-1 (a, b), occludin (c, d), or claudin-1 (e, f). The results of densitometric analysis were shown in b, d, and f. Asterisks indicated significant difference as compared with control (**P*<0.05; ***P*<0.01; ****P*<0.001).

RESULTS

Histological Assessment

We first examined the influence of EPEC infection on colonic mucosal architecture by H&E staining. The epithelial lining of the colon from control mice was intact and showed well-organized colonic mucosa (Fig. 1a). However, when mice were treated with EPEC for a 5-day period, loss of epithelial cells was seen and the surface epithelium had an irregular appearance (Fig. 1b–d). This infection also caused a mucosal hyperplasia in the colon accompanied by mild infiltration of inflammatory cells. Thus, this

study revealed considerable tissue damages during EPEC infection *in vivo*.

Redistribution of Tight Junction Proteins Induced by EPEC Infection

To investigate whether EPEC infection affected tight junction, we examined the expression of tight junction proteins in membrane microdomains of tight junction. A previous report has suggested that tight junctions are specific lipid raft [23]. The detergent-resistant tight junction membrane microdomains were isolated using detergent extraction and sucrose density gradient centrifugation, and the extracted tight junction

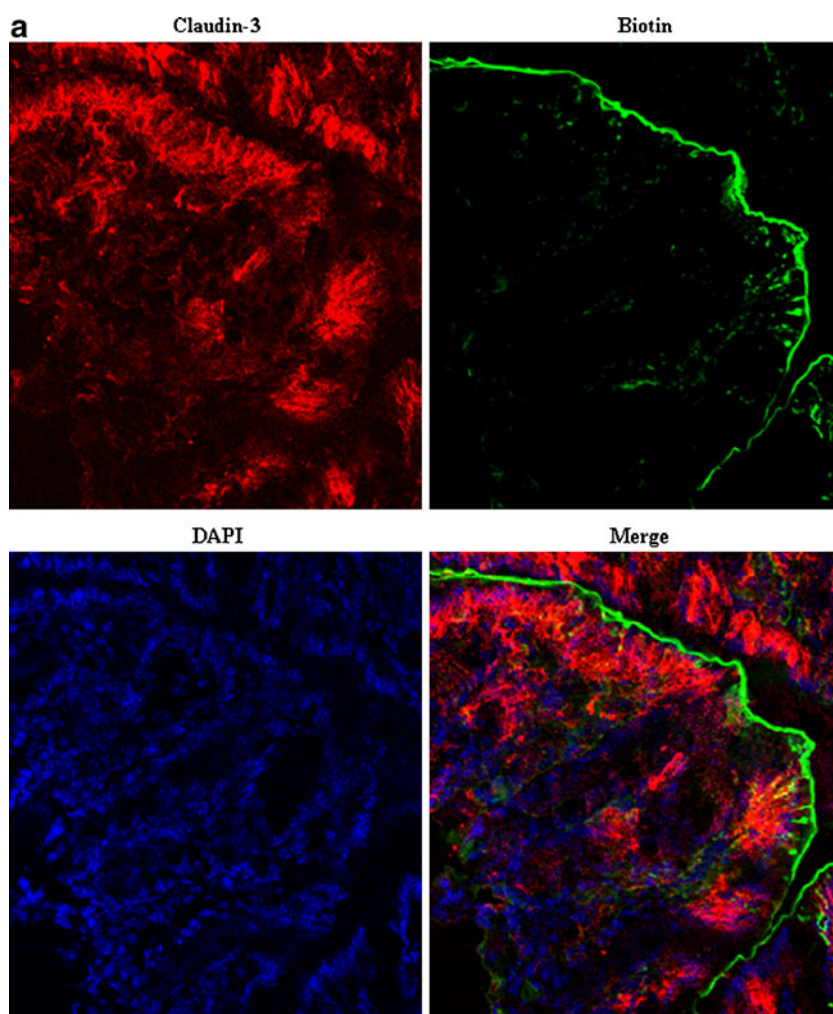


Fig. 3. Tight junction barrier were disrupted by EPEC infection. The control (a, c) and EPEC-infected (b, d) colon tissues were stained with claudin antibodies and the cell tracer biotin to examine barrier permeability. DAPI was used to label the nucleus. a, b Claudin-3 and biotin staining. c, d Claudin-5 and biotin staining.

proteins were analyzed by Western blot analysis. Using this method, we found that the lipid raft marker protein caveolin-1 was localized in fractions 2–5 in control, corresponding to tight junction membrane microdomains (Fig. 2a). EPEC infection affected the distribution of caveolin-1 in tight junction membrane microdomains. There was an 80.1% decrease in caveolin-1 localization in tight junction membrane microdomains after 1 day of infection with EPEC, and the expression of caveolin-1 declined by 37.8% and 18.4% following 3 and 5 days of infection, respectively (Fig. 2b). Moreover, Western blot analysis revealed that EPEC infection induced a movement of occludin and claudin-1 out of tight junction membrane microdomains. In control mice, 19.6% occludin was found in tight junction membrane microdomains (Fig. 2c, d). However, after EPEC infection, occludin was displaced

from the low-density fractions to the bottom of the gradient. Another tight junction protein claudin-1 was also redistributed after EPEC infection (Fig. 2e, f).

EPEC Infection Induced Mucosal Barrier Dysfunction

To determine whether the altered distribution of tight junction proteins induced functional disruption of tight junctions, we utilized biotin as a molecular tracer to assess the integrity of the epithelial barrier, and claudin-3 and 5 were double-labeled with the biotin tracer. Biotin was found to be restricted to the luminal boundary of the colon epithelium in control (Fig. 3a, c), which is in consistent with our previous study [24]. After EPEC infection, biotin was no longer held to the

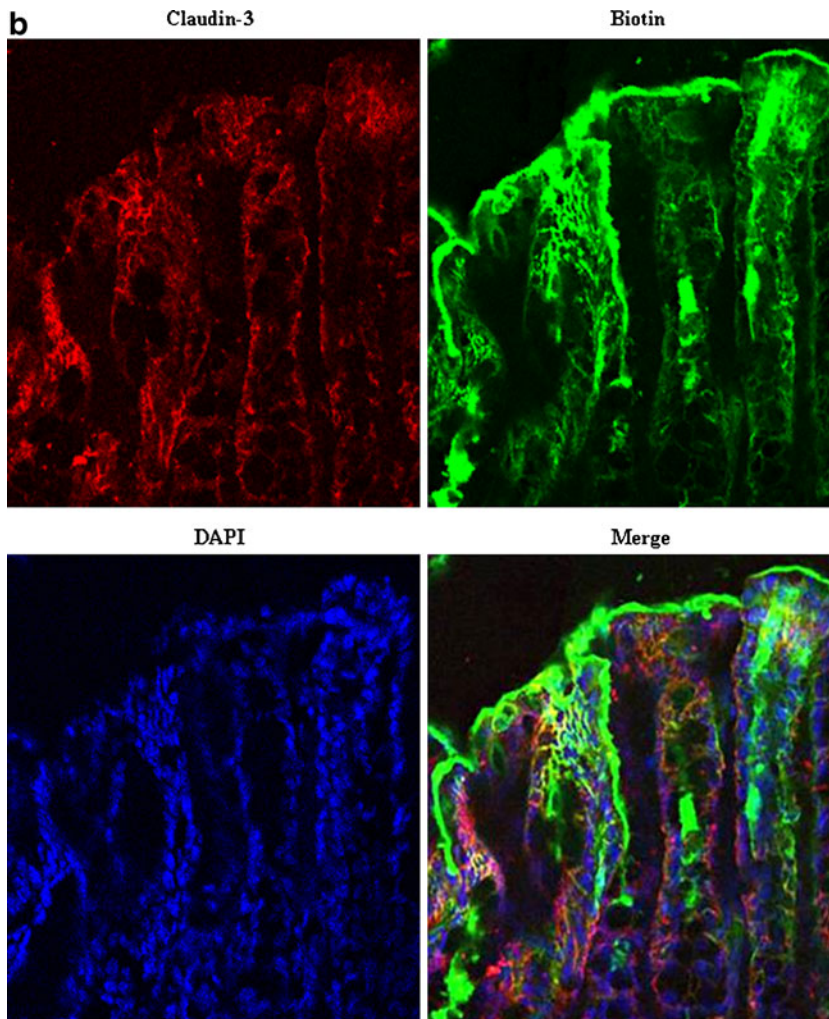


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luminal bounder, and it permeated the epithelium and extended into the lamina propria. The biotin was observed to pass into the epithelium through areas of altered claudin location (Fig. 3b, d). It indicated that the tight junction was functionally altered by EPEC infection.

DISCUSSION

Enteropathogenic *E. coli* infection is a main cause of infantile diarrhea in developing countries, which leads to several hundred thousand infant deaths each year [25]. Until now, the underlying mechanisms by which EPEC triggers the severe watery diarrhea are unclear. Previous

studies have suggested that this disease is in part dependent on EPEC interaction with enterocytes [26, 27]. This intimate attachment of bacteria to the host cell membrane results in loss of absorptive microvilli, which is called an A/E lesion. Thus, understanding how A/E bacterial pathogens colonize their hosts is of great importance in combating these infections. Our previous *in vitro* study demonstrated that EPEC infection caused disruption of tight junction structure and function [15]. Furthermore, our results indicated that the tight junction membrane microdomains were key sites of EPEC entry. To further our study, in the present report, we infected mice with a wild-type EPEC 2348/69 and investigated changes in the distribution of occludin and claudins in tight junction membrane microdomains and alteration in intestinal barrier function. We found that EPEC induced

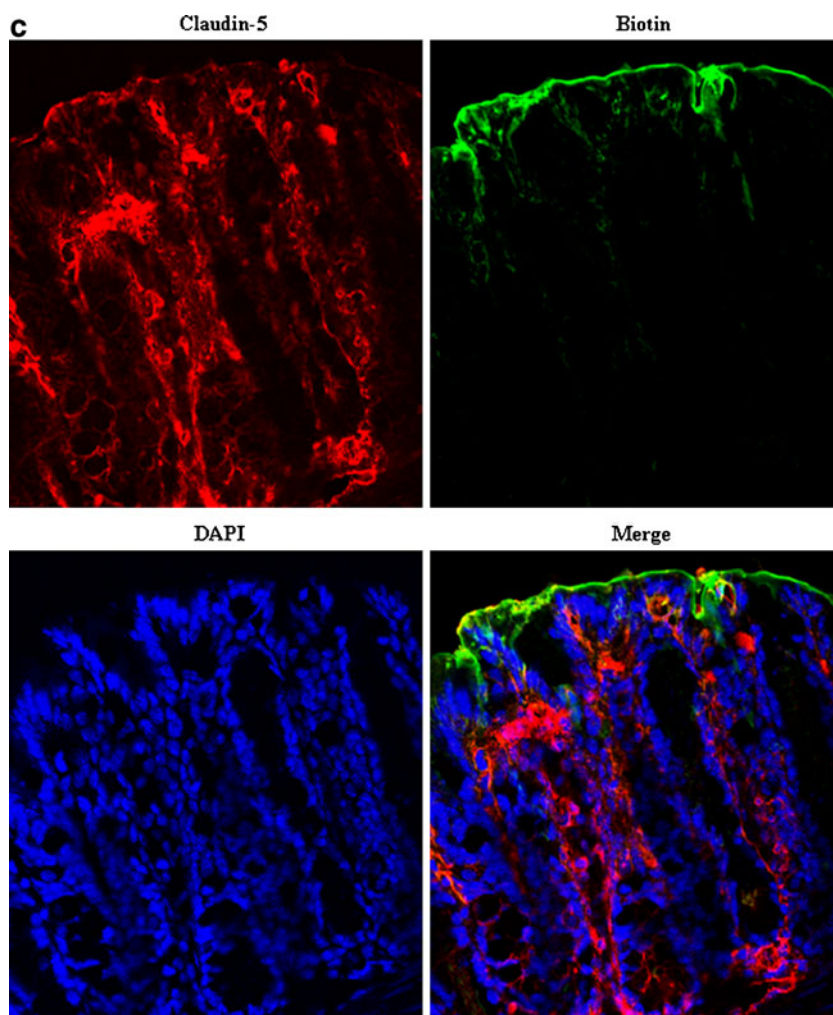


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tight junction barrier dysfunction which was accompanied by redistribution of occludin and claudins in tight junction membrane microdomains.

Tight junctions form a regulatable semipermeable barrier at the boundary between the apical and the basolateral membrane domains [28, 29]. The key role of tight junction is acting as a regulatory barrier to the movement of molecules. Tight junctions are constituted of an array of tight junction proteins. The regulation of tight junction barrier function is a very intricate process and many transmembrane and intracellular proteins are involved [30]. It is increasingly recognized that tight junctions are altered both physiologically and pathophysiologically. Perturbation of tight junction was found to be involved in several disease states including enteric infection [16], Crohn's disease [29], and experimental

colitis [22]. The impaired integrity of the intestinal barrier has been observed in patients with Crohn's disease [31]. A recent study reported that the expression pattern of claudins was altered in Crohn's disease [32].

The diarrhea induced by EPEC infection was once thought to be caused by the loss of absorptive microvilli [33]. Recently, a growing number of *in vitro* studies have demonstrated that EPEC infection affected tight junction barrier function. Infection of the intestinal cell monolayers induced a time-dependent decrease in transepithelial electrical resistance which was correlated with the dissociation of occludin from intestinal epithelial tight junctions [17]. Despite extensive studies on the effect of EPEC infection on tight junction function *in vitro*, little is known about the changes of tight junction during EPEC infection *in*

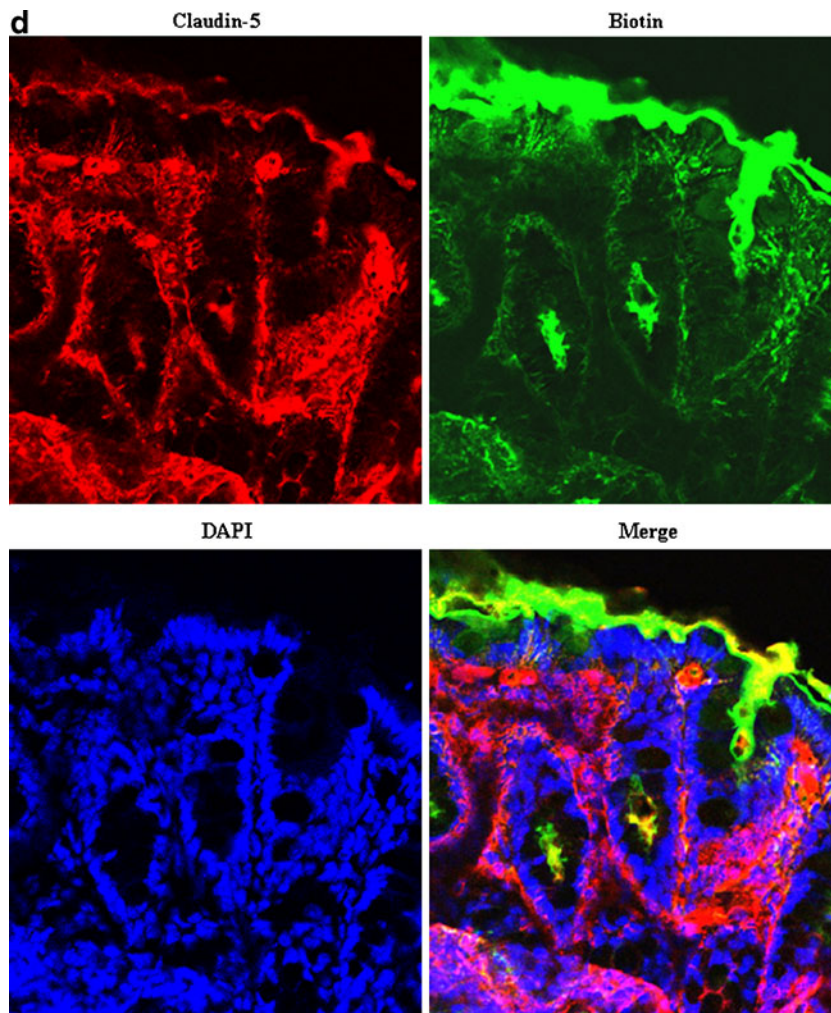


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in vivo. In the present study, we demonstrated that tight junction barrier function was altered during EPEC infection. Similar with what has been reported for *Campylobacter jejuni*, our findings showed that EPEC infection induced a translocation of occludin and claudin-1 from the TJ platform [34]. We also observed that the cell tracer could pass between cells through the tight junctions during EPEC infection. These findings indicated that EPEC disrupted multiple host tight junction proteins *in vivo* resulting in a functionally deficient epithelial barrier. In summary, our study demonstrated the underlying molecular mechanisms of tight junction barrier disruption triggered by EPEC infection. These results may provide critical information on the pathogenesis of EPEC infection.

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