Protective effects of *Lactobacillus plantarum* against epithelial barrier dysfunction of human colon cell line NCM460

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Abstract

AIM: To investigate the effects of *Lactobacillus plantarum* (*L. plantarum*) in the intestinal permeability and expression of tight junction (TJ) using the normal human colon cell line NCM460.

METHODS: Paracellular permeability of NCM460 monolayers was determined by transepithelial electrical resistance and dextran permeability. Expression of TJ proteins in NCM460 cell monolayers was detected by Western blotting and quantitative real-time polymerase chain reaction.

RESULTS: *L. plantarum* played an important role in increasing transepithelial electrical resistance and decreasing the permeability to macromolecules of NCM460 monolayers against the disruption caused by enteropathogenic *Escherichia coli* (*E. coli*) or enteroinvasive *E. coli*. *L. plantarum* also prevented the decrease in the expression of TJ proteins and F-actin in NCM460 cells.

CONCLUSION: *L. plantarum* can protect against dysfunction of NCM460 intestinal epithelial barrier caused by enteropathogenic *E. coli* or enteroinvasive *E. coli*, and thus can be a potential candidate of therapeutic agents for the treatment of intestinal diseases.

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Key words: *Lactobacillus plantarum*; NCM460; Tight junction; Intestinal barrier

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INTRODUCTION

The human intestinal system included a group of viable microorganisms, which exceed the total number of somatic and germ cells[1-3]. Therefore, the human colon is confronted with the highest bacterial load in the digestive tract with enormous bacteria per gram of feces. Growing evidence showed that bacteria closely adherent to the mucosa are more relevant to human body, compared with those evacuated in the feces[4-6]. There is a homeostasis between probiotics and pathogens in the intestinal
systems of healthy individuals\textsuperscript{[6,7]}, which, if broken, may lead to an imbalanced ecological microenvironment and subsequent intestinal barrier dysfunction\textsuperscript{[8-10]}. Thereafter, the accumulation of pathogens and secretory products, such as the exotoxins and secretory antigens, can also directly or indirectly initiate and amplify the local and systemic inflammatory responses\textsuperscript{[10,11]}. Probiotics of the genus lactobacillus that reside in the human intestine play an important part in maintaining the homeostasis of gut flora by adhering and colonizing to the intestinal mucosa and competing with pathogenic bacteria, such as pathogenic Escherichia coli (E. coli)\textsuperscript{[12,13]}. Enteric diseases with flora disequilibrium have been treated with lactobacillus over the past decades\textsuperscript{[14-16]}. There is evidence indicating that the modulation of the gut flora by lactobacillus can improve the intestinal epithelial barrier function\textsuperscript{[17]}

Adhesion of lactobacillus to the intestinal epithelium initially involves the activation of specific binding between bacterial ligands and their corresponding surface receptors on the intestinal cells of the host, following the nonspecific physical interactions\textsuperscript{[16,18]}. Generally, these ligands are adhesive molecules either existing on the surface layer of the bacteria or secreting from the mycelium of the bacteria. Furthermore, these ligands could interact with the corresponding receptors on the surface of the intestinal epithelial cells. Thereafter, these adhesins activate specific signal transduction pathways in both the bacteria and host cells. The interaction between lactobacillus and the intestinal epithelial cells can also block the adhesion of other pathogenic bacteria to the receptors of the intestinal epithelial cells, such as enteropathogenic E. coli (EPEC) and enteroinvasive E. coli (EIEC). As a widespread member of the genus lactobacillus, Lactobacillus plantarum (L. plantarum) is commonly found in many fermented food and anaerobic plant products. Our previous studies demonstrated that L. plantarum was able to prevent colonic damage caused by EIEC or inflammation in vitro, in vivo and in patients with acute pancreatitis\textsuperscript{[19,20]}. The normal human colon cell line NCM460, which is derived from the normal human colon mucosal epithelium and expresses colonic epithelial cell-associated antigens such as cytokeratins and villin, has been applied exclusively in various intestinal research areas, including the infectious diseases\textsuperscript{[20-22]}

Our previous studies indicated that L. plantarum exerted its therapeutic effects by adhering to the intestinal epithelial cells, restoring tight junction (TJ) structure and function, and reducing paracellular permeability. However, studies about the interaction between lactobacillus and the human intestine were limited in the cancer cell line and the animal models, and further researches based on the normal human intestinal cells are still needed. Therefore, our study aims to investigate the protective effects of L. plantarum against epithelial barrier dysfunction of the normal human colon cell line NCM460 caused by EIEC and EPEC.

**MATERIALS AND METHODS**

**Bacterial strains and culture conditions**

*L. plantarum* CGMCC 1258 (generously provided by Dr. Xiao-Min Hang, the Onlly Institute of Life Science, Shanghai Jiao Tong University, Shanghai, China) was inoculated in 5% fresh De Man, Rogosa and Sharpe broth at 37℃ for 24 h, harvested by centrifugation (3500 × g at 4℃ for 20 min, and washed with 50 mL 0.01 mol/L phosphate buffered saline (PBS) (pH 7.4). The EIEC strain ATCC 43893 (O124:NM) and EPEC strain ATCC 43887 (O111:NM) (both from Shanghai Municipal Center for Disease Control and Prevention, Shanghai, China) were grown in static Dulbecco's modified eagle media (DMEM) at 37℃ for 24 h. Quantification of bacterial density was measured at 600 nm (Beckman DU-50 spectrophotometer) with the colony forming units.

NCM460 cells were purchased from INCELL Corporation (San Antonio, TX, USA) and cultured in M3 media supplemented with 10% FBS, 100 U/mL penicillin and 100 g/mL streptomycin at 37℃ in a 95% humidified atmosphere with 5% CO\textsubscript{2}, as previously described\textsuperscript{[23]}

**Measurement of transepithelial electrical resistance in NCM460 cell monolayers**

NCM460 cells were grown on filters (Millicell culture plate inserts; 0.4 μm pore size; 0.6 cm\textsuperscript{2} surface area) at 37℃ in a 95% humidified atmosphere, with 5% CO\textsubscript{2}. At full confluence (10-14 d) (i.e. a monolayer was formed), a transepithelial electrical resistance (TER) of > 450 Ω cm\textsuperscript{2} monolayer was achieved as measured using a voltmeter (Millicell-ERS; Millipore, MA, USA). The intestinal epithelial monolayers were treated with EIEC or EPEC in the presence or absence of L. plantarum. In infection groups, 100 μL EIEC ATCC43893 (O124:NM) and EPEC ATCC43887 (O111:NM) at 1.0 × 10\textsuperscript{8}/mL were, respectively, added to the apical side of the cell culture insert for rapid infection of the monolayer, with an inoculation ratio of EIEC/EPEC to NCM460 cells of 100:1, and the insert was placed in a 50-mL tube and centrifuged at 200 × g for 4 min. In L. plantarum groups, L. plantarum (100 μL of 1.0 × 10\textsuperscript{8}/mL) was added onto the monolayer of NCM460 cells simultaneously with the EIEC/EPEC infection. NCM460 cells cultured under the same conditions but without the infection of EIEC/EPEC, and addition of L. plantarum served as the control group. Two experiments were performed separately for EIEC and EPEC.

The integrity of the confluent polarized monolayers was verified by measuring TER at different time intervals. TER (Ω cm\textsuperscript{2} monolayer) = (Total resistance - Blank resistance) (Ω) × Area (cm\textsuperscript{2} monolayer). Because TER values often vary among individual NCM460 cultures, the electrical resistance value was recorded for each monolayer before and after the treatment, and the percentage in the decrease of TER from the baseline (%TER) was calculated.
compartment were placed into a 96-well plate (Corning Costar Co., MA, USA) and analyzed to determine their fluorescent intensity using the Odyssey infrared imaging system (LI-COR Biosciences, NE, USA) at a wavelength of 700 nm. Relative intensity (RI; the integrated intensities of treated samples relative to the integrated intensity of untreated samples) was calculated to indicate the effect of the treatment.

**Western blotting for determining the distribution and expression of TJ proteins in NCM460 cell monolayers**

For Western blotting, NCM460 cells were cultured and the monolayers were treated as described above, and the protein samples from NCM460 cells was prepared for sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) according to the previous studies.[20] SDS-PAGE was performed using the standard laboratory techniques with a discontinuous gradient, 5% (w/v) stacking gel and a 10% (w/v) separating gel, in a Mini-PROTEAN II (Bio-Rad Laboratories, CA, USA). Briefly, samples were mixed with loading buffer containing SDS and mercaptoethanol, boiled for 3 min, centrifuged, and loaded onto the SDS-PAGE gel for separation. Molecular weights of samples were determined by comparing mobility with known marker proteins. Gel was then transferred to PVDF membrane (Millipore, MA, USA) in a semidy electroblotter (Bio-Rad Laboratories) for 120 min at 100 V. The membrane was washed three times (20 min each) with PBS containing 0.1% Tween-20 (PBS-T buffer). After blocking overnight in Tris-buffered saline containing 0.05% Tween-20 (TBS-T) and 5% dry powdered milk, membranes were washed three times for 5 min each with TBS-T and incubated with corresponding primary antibodies against TJ proteins (claudin-1, occludin, JAM-1, and ZO-1) and a cell cytoskeleton element F-actin (all from Abcam, MA, USA) for 2 h at room temperature. After three washes with TBS-T, the membranes were incubated for 1 h with corresponding HRP-conjugated secondary antibodies. The membrane was washed three times (60 min each) with PBS-T buffer. The TJ proteins were tested using enhanced chemiluminescence (ECL kit; Pierce, IL, USA) according to the manufacturer’s instructions.

**Quantitative real-time polymerase chain reaction for detecting the mRNA expression of TJ proteins in NCM460 cells**

mRNA expression of TJ proteins, including claudin-1, occludin, JAM-1, and ZO-1, was determined by quantitative real-time polymerase chain reaction (RT-PCR). After the treatment as described above, total RNA was isolated from NCM460 cells using the Trizol reagent (Gibco Brl, USA)[28], followed by DNase I treatment. The quantity and quality of RNA were verified with the ratio of absorbance values at 260 and 280 nm, and by visualization of the bands on agarose gels. For each sample, 600 ng mRNA was used in reverse transcription reaction (iScript kit from BioRad Laboratories) according to the manufacturer’s specifications. Further analysis of mRNA levels of each group was performed by RT-PCR with a light-cycling system (LightCycler; Roche Diagnostics GmbH, Mannheim, Germany). Sequences of the primers used are listed in Table 1. The mRNA expression level was described as the ratio of the mean reading of the experimental group over that of the control group for NCM460 cells.

**RESULTS**

*L. plantarum prevented EIEC/EPEC-induced decrease of TER in NCM460 cells*

TER in the NCM460 cell monolayers was decreased significantly in response to infection with EIEC/EPEC compared with uninfected control cells. However, decrease of TER induced by EIEC/EPEC was prevented by the simultaneous treatment of *L. plantarum* (Figure 1A and B).

*L. plantarum inhibited increased macromolecular permeability of NCM460 cell monolayers in response to EIEC/EPEC*

EIEC/EPEC had an obvious enhancing effect on permeability of NCM460 cell monolayers, as compared with the uninfected control cells. However, this effect was inhibited by the co-treatment of *L. plantarum* (Figure 2A and B).

*L. plantarum prevented the decreased expression of TJ proteins and a cell cytoskeleton element F-actin detected by Western blotting*

The expression of TJ proteins, including claudin-1, occludin, JAM-1 and ZO-1, and the cytoskeleton element F-actin, was decreased in NCM460 cells infected with EIEC or EPEC (Figure 3A and B) compared with the control cells, as detected by Western blotting of epithelial whole cell protein extracts of NCM460 cells (P < 0.001). However, after the pre-treatment of *L. plantarum*, the expression of TJ proteins and F-actin remained at similar levels to the control cells.

*L. plantarum prevented the decreased expression of TJ proteins as detected by quantitative RT-PCR*

mRNA expression of TJ proteins, including claudin-1, oc-
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**Figure 1** *Lactobacillus plantarum* inhibited the decreased transepithelial electrical resistance through NCM460 cells induced by enteroinvasive *Escherichia coli* (A) or enteropathogenic *Escherichia coli* (B). A: After infection with enteroinvasive Escherichia coli (EIEC) for 24 h, transepithelial electrical resistance (TER) of NCM460 monolayers was decreased significantly compared with the value in the control group. However, TER of EIEC-infected NCM460 monolayers, when simultaneously co-incubated with *Lactobacillus plantarum* (*L. plantarum*), was significantly higher than those in the EIEC-infected NCM460 monolayers. *P < 0.05* vs control group; *c P < 0.05* vs corresponding EIEC or EPEC group.

**Figure 2** *Lactobacillus plantarum* inhibited the increased macromolecular permeability through NCM460 cells induced by enteroinvasive *Escherichia coli* (A) or enteropathogenic *Escherichia coli* (B). A: After infection with enteroinvasive Escherichia coli (EIEC) for 120 min, the relative intensity (RI) was significantly increased in the EIEC group compared with the control group. However, the RI was decreased significantly in *Lactobacillus plantarum* (*L. plantarum*) groups compared with the EIEC group; B: Similar results were obtained in the experiments with enteropathogenic Escherichia coli (EPEC). The data at each time point represent the mean ± SD obtained from four individual NCM460 monolayers. *P < 0.05* vs control group; *c P < 0.05* vs corresponding EIEC or EPEC group.

**Figure 3** *Lactobacillus plantarum* prevented the decrease in the expression of tight junction proteins in NCM460 cells induced by enteroinvasive *Escherichia coli* or enteropathogenic *Escherichia coli* detected by Western blotting. A: The expression level of tight junction (TJ) proteins was high, including claudin-1, occludin, JAM-1 and ZO-1, in the control group. However, in the enteroinvasive Escherichia coli (EIEC) or enteropathogenic Escherichia coli (EPEC) group, TJ proteins were significantly decreased compared with the control group, which was not observed in the *Lactobacillus plantarum* (*L. plantarum*) group; B: Semi-quantitative analysis of Western blotting showed similar results. The data at each time point represent the mean ± SD obtained from four individual NCM460 monolayers. *P < 0.05* vs control group; *c P < 0.05* vs corresponding EIEC or EPEC group.

claudin, JAM-1 and ZO-1, was significantly decreased in the NCM460 cells infected with EIEC or EPEC, as compared with the uninfected control cells (Figure 4A and B). However, treatment with *L. plantarum* raised mRNA expres-
It has been reported that probiotics, such as *L. plantarum*, have beneficial effects on the human intestinal barrier function in patients with intestinal diseases [13,31]. Our previous studies also found that *L. plantarum* adhered to the intestinal epithelial cells, restored TJ structure and function, reduced paracellular permeability, and then showed the therapeutic effects [20,23]. However, the studies about the interaction between lactobacillus and the human intestine were only limited in the cancer cell line and the animal models. The present study investigated the protective effects of *L. plantarum* against epithelial barrier dysfunction of the normal human colon cell line NCM460.

*E. coli* and *L. plantarum* are able to attenuate the deleterious effects of the intestinal epithelial barrier, while p38, ERK1, 2, or PI3K had no effects. After treated by lactobacillus, epithelial cells exposed to cytokines, LPS, and interferon-γ in epithelial function, and interferon-γ in epithelial function [32,33]. Furthermore, special signal transduction pathway is involved in the protective effects of *L. plantarum* on the intestinal epithelial barrier. Janus kinase inhibitor synergistically potentiated the effects of lactobacillus acidophilus on TER and permeability, while p38, ERK1, 2, or PI3K had no effects. After treated by lactobacillus, epithelial cells exposed to cytokines reduced the activation of SOCS3 and STAT1, 3.

**DISCUSSION**

**Figure 4 Protective effects of *Lactobacillus plantarum* and MIMP in mRNA expression of tight junction proteins in NCM460 cells detected by real-time polymerase chain reaction.** A: The mRNA expression of tight junction (TJ) proteins, including occluding, claudin-1, JAM-1 and ZO-1, was decreased in enteroinvasive *Escherichia coli* (EIEC) group compared with the control group. However, in *Lactobacillus plantarum* (*L. plantarum*) group, the mRNA expression levels of the TJ proteins were similar to those in the control group. B: Similar results were obtained with enteropathogenic *Escherichia coli* (EPEC). The data represent the mean ± SD obtained from three individual NCM460 monolayers. *P* < 0.05 vs control group; *P* < 0.05 vs corresponding EIEC or EPEC group.
We believe that our study broadens our knowledge of effects of \textit{L. plantarum} in intestinal epithelial function and its therapeutic effects in the cellular and molecular mechanisms of intestinal barrier dysfunction and intestinal inflammation and justifies the use in inflammatory disorders, which is significant to both biotechnical and clinical fields. \textit{L. plantarum} can protect against intestinal epithelial barrier dysfunction of NCM460 caused by EIEC or EPEC. However, the bacterial protein and its exact mechanisms of action remain unknown. We are conducting a study in an attempt to identify the protein and the smallest active domain within the protein from \textit{L. plantarum} strain CGMCC1258 that is responsible for the adhesion of the bacterium to the intestinal epithelium. And further functional characterization by determining the effects of smallest active domain on the intestinal barrier function and immune responses is also in progress.

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**COMMENTS**

**Background**

\textit{Lactobacillus plantarum} (\textit{L. plantarum}) that resides in the human intestine plays an important part in maintaining the homeostasis of gut flora by adhering and colonizing to the intestinal mucosa and competing with pathogenic bacteria, which contributes to the protection of the human intestinal barrier function.

**Research frontiers**

Although \textit{L. plantarum} exerted its therapeutic effects by adhering to the intestinal epithelial cells, restoring tight junction structure and function, and reducing paracellular permeability, the studies about interaction between lactobacillus and the human intestine were just limited in the cancer cell line and the animal models, and further studies based on the normal human intestinal cell had been unavailable.

**Innovations and breakthroughs**

Using the normal human colon cell line NCM460, this study investigated the protective effects of \textit{L. plantarum} against epithelial barrier dysfunction caused by enteropathogenic \textit{Escherichia coli} (E.coli) or enteroinvasive E. coli.

**Applications**

\textit{L. plantarum} can be a potential candidate of therapeutic agents for the treatment of intestinal diseases.

**Terminology**

NCM460 cell line is a normal human colon cell line, which is derived from the normal human colon mucosal epithelium and expresses colonic epithelial cell-associated antigens such as cytokeratins and villin.

**Peer review**

This is a straightforward study extending previous work of the authors showing that \textit{L. plantarum} maintains a high resistance to permeability to enteropathogenic and enteroinvasive E. coli. The work is extended to NCM460 colon cell cells in culture. Transepithelial electrical resistance was maintained high, dextran permeability was low and TJ protein expression was normal.

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