Role of Urea in Intestinal Barrier Dysfunction and Disruption of Epithelial Tight Junction in Chronic Kidney Disease

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Key Words

Endotoxin · Inflammation · Gastrointestinal pathology · End-stage renal disease · Uremia · Cardiovascular disease

Abstract

Background: Chronic kidney disease (CKD) impairs intestinal barrier function which leads to endotoxemia and systemic inflammation. We have found depletion of intestinal epithelial tight junction (TJ) proteins in animals with CKD. We further showed that addition of end-stage renal disease patients' plasma to the culture medium provokes a marked drop in transepithelial electrical resistance (TER) and depletion of TJ proteins in cultured human enterocytes. These effects were less severe with post- than with prehemodialysis plasma, suggesting a role of dialyzable agent(s). This study tested the hypothesis that intestinal barrier dysfunction in uremia may be due to diffusion of urea into the gut and its conversion to ammonia by microbial urease. **Methods:** Human enterocytes (T84 cells) were seeded on Transwell plates and utilized when TER exceeded $1,000 \text{ m}\Omega \cdot \text{cm}^2$ to ensure full polarization and TJ formation. Confluent cells were then incubated for 24 h in media containing 0, 42 or 74 mg/dl urea or urea plus urease to simulate presence of microbial flora. **Results:** At clinically relevant concentrations, urea caused a concentration-dependent fall in TER and the key TJ proteins

claudin-1, occludin and zonula occludens 1. The effects of urea were dramatically amplified by urease causing cell detachment, dissipation of TER, and massive loss of TJ proteins. **Conclusions:** Uremia-induced disruption of intestinal TJ and barrier function is, in part, mediated by urea which is generally considered to be a nontoxic retained metabolite. These findings reveal a novel mechanism for a salutary effect of urea-lowering strategies, e.g. low-protein diet and longer and more frequent dialysis regimens in advanced CKD.

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Introduction

There is emerging evidence pointing to intestinal barrier dysfunction and its role in the pathogenesis of systemic inflammation in humans and animals with advanced chronic kidney disease (CKD) [1, 2]. In an earlier study, we found marked depletion of the key transcellular (claudin-1 and occludin) and intracellular [zonula occludens 1 (ZO-1)] constituents of colonic epithelial tight junction (TJ) in the rats with CKD induced by either 5/6 nephrectomy or adenine-induced tubulointerstitial nephropathy. The reduction in the abundance of the TJ proteins was accompanied by their normal or even elevated mRNA abundance [3], suggesting a posttranscriptional or

posttranslational nature of the downregulation of these proteins. In a recent study [4], we asked whether uremia-induced intestinal epithelial barrier dysfunction and the TJ protein depletion are mediated by the retained uremic toxins/metabolites and if so, whether they can be removed by dialysis. To this end, we tested the effects of addition to the culture media of pre- and posthemodialysis plasma samples from end-stage renal disease (ESRD) patients and plasma from healthy controls on the polarized T84 human colonic epithelial cells. When seeded on Transwell plates, T84 cells form a polarized, impermeable monolayer that exhibits many of the functional characteristics of the intestinal epithelial cells *in vivo*, including vectorial solute transport and barrier function [5, 6]. The study showed that compared with the control plasma, incubation in media containing predialysis plasma from ESRD patients caused a marked drop in transepithelial electrical resistance (TER) denoting increased epithelial permeability. This was accompanied by significant reductions in transcellular, *i.e.* claudin-1 and occludin, and intracellular, *i.e.* ZO-1, protein constituents of the epithelial TJ apparatus. The severity of TJ damage and dysfunction was significantly less in cells exposed to the postdialysis than predialysis plasma. These findings pointed to the presence of as yet unidentified, but partially dialyzable, product(s) in the uremic plasma capable of degrading epithelial TJ and impairing intestinal epithelial barrier function.

The reduction in urinary excretion of nitrogenous waste products in advanced CKD results in their accumulation in the body fluids. The most abundant among these metabolites is urea, which has been considered to be a nontoxic and inert retained uremic metabolite. The rise in urea concentration in the intracellular and extracellular fluid compartments results in its heavy influx into the gastrointestinal tract [7]. Within the intestinal lumen, urea is hydrolyzed by microbial urease forming large quantities of ammonia [$\text{CO}(\text{NH}_2)_2 + \text{H}_2\text{O} \rightarrow \text{CO}_2 + 2\text{NH}_3$] which is readily converted to ammonium hydroxide [$\text{NH}_3 + \text{H}_2\text{O} \rightarrow \text{NH}_4\text{OH}$] [8, 9]. Ammonium hydroxide, in turn, leads to a modest rise in the luminal fluid's pH, causes mucosal irritation and promotes enterocolitis [10, 11]. Based on these observations, we hypothesize that ammonium hydroxide generated from hydrolysis of urea by microbial urease in the intestinal lumen of uremic patients may contribute to the epithelial barrier dysfunction and erosion of the TJ protein constituents. The present study was designed to test this hypothesis by exploring the effect of clinically relevant concentrations of urea alone and urea plus urease on intestinal epithelial barrier function and structure *in vitro*.

Fig. 1. Bar graphs depicting the TER in intestinal epithelial T84 cell monolayers incubated for 24 h in regular media and those incubated in media containing 42 or 72 mg/dl urea. *** $p < 0.001$.

Methods

Cell Culture and Incubation Studies

T84 cells were obtained from American Type Culture Collection (Manassas, Va., USA) and were grown in DMEM/F12 medium (Invitrogen Inc., Carlsbad, Calif., USA) containing sodium bicarbonate 1.2 g/l, L-glutamine 2.5 mM, HEPES 15 mM, and sodium pyruvate 0.5 mM (Invitrogen) supplemented with 10% fetal calf serum. To form polarized monolayers, cells were seeded (60,000 cells/cm²) in 12-well or 6-well plates with Transwell inserts (Fisher Scientific, Pittsburgh, Pa., USA) for measurement of TER and Western blot analysis. The T84 monolayers were maintained for 21 days at 37°C in complete medium in an incubator containing a mixture of air with 5% carbon dioxide. The medium was changed every other day and the TER was measured regularly using Millicell ERS-2 meter (EMD Millipore Inc., Billerica, Mass., USA). When a TER exceeding 1,000 mΩ·cm² was reached, the monolayers were incubated for 24 h at 37°C in DMEM/F12 medium alone or in medium containing 42 or 72 mg/dl (70, or 120 μM) urea (Sigma-Aldrich, St. Louis, Mo., USA) with or without urease (Sigma Inc.). The given urea concentrations were chosen to approximate the pre- and posthemodialysis values generally found in ESRD patients and as such were considered to be clinically relevant. Urease was added to the medium at 1 unit/μM urea. At the conclusion of the 24-hour incubation period, the TER was measured and cells were harvested and processed for Western blot analysis.

Western Blot Analyses

The cells were lysed with M-PER Mammalian Protein Extraction Reagent (Fisher Scientific). Protein concentration in the cell lysate was determined by BSA assay kit (Pierce, Rockford, Ill., USA) and 20 μg of total protein from each sample were fractionated on 4–12% Bis-Tris gradient gel (Invitrogen) at 120 V for 2 h and transferred to a nitrocellulose membrane. The membrane was then incubated with rabbit anti-claudin-1 or rabbit anti-occludin or mouse anti-ZO-1 (Invitrogen) antibodies at 1:250 dilu-

Fig. 2. Representative Western blots and group data depicting protein abundance of occludin, claudin-1 and ZO-1 in intestinal epithelial T84 cell monolayers incubated for 24 h in regular media and media containing 42 or 72 mg/dl urea. CTL = Control. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$ versus control.

tions, and anti-actin antibody (Sigma-Aldrich) at a 1:10,000 dilution overnight. The appropriate horseradish peroxidase-conjugated secondary antibodies (Sigma-Aldrich) were used at a 1:5,000 dilution. The membrane was visualized with SuperSignal West Pico (Pierce) and developed by autoluminography.

Data Analysis

Analysis of variance and Student's *t* test were used in the statistical evaluation of the data which are presented as mean $\pm$ SD. *p* values less than 0.05 were considered significant.

Results

Effect of Urea and Urea plus Urease on TER and pH

Incubation in media containing urea at 42 or 72 mg/dl resulted in a significant concentration-dependent fall in basal-apical electrical resistance pointing to the ability of urea to impair epithelial barrier function (fig. 1). Epithelial monolayers incubated in media containing both urea and urease exhibited detachment which precluded the ability to measure TER. Addition of urease alone to the culture medium had no significant effect on TER. The average pH of the culture medium rose from 7.1 to 7.3, 24 h after addition of urea plus urease.

Effect of Urea and Urea plus Urease on Epithelial TJ Proteins

The fall in the monolayer TER in cells incubated in urea-containing media was associated with significant concentration-dependent reductions in claudin-1, occludin, and ZO-1 abundance (fig. 2, 3). Addition of urease to the incubation media designed to simulate the presence of urease-possessing colonic bacteria led to a further and drastic fall in claudin-1, occludin and ZO-1 abundance (fig. 4).

Discussion

Systemic inflammation is a constant feature and a major mediator of cardiovascular disease, cachexia, anemia and numerous other morbidities in patients with advanced CKD [12–16]. Inflammation in these patients is commonly associated with endotoxemia in the absence of clinical infection [17–19]. The most likely source of endotoxemia in the infection-free patients is the gut microbial flora. However, entry of endotoxin from the intestinal lumen to the circulation can occur only when the mucosal barrier is impaired. As summarized in recent reviews [1, 2], there is compelling evidence supporting the impairment of the intestinal barrier function and its potential

Fig. 3. Representative Western blots and group data depicting protein abundance of occludin, claudin-1 and ZO-1 in intestinal epithelial T84 cell monolayers incubated for 24 h in media containing 42 mg/dl urea alone and those incubated in media containing 42 mg/dl urea plus urease. * $p < 0.05$; *** $p < 0.0001$ versus control.

Fig. 4. Representative Western blots and group data depicting protein abundance of occludin, claudin-1 and ZO-1 in intestinal epithelial T84 cell monolayers incubated for 24 h in media containing 72 mg/dl urea alone and those incubated in media containing 72 mg/dl urea plus urease. *** $p < 0.001$ versus control.

contribution to the prevailing inflammation in the uremic humans. This is based on the reported increase in intestinal permeability to high-molecular-weight polyethylene glycols [17, 20], presence of endotoxemia [17, 19], and histological evidence of chronic inflammation

throughout the gastrointestinal tract in this population [10, 11]. The TJ apparatus is essential for the epithelial polarity and restriction of paracellular permeability of intestinal epithelium. Consequently, disruption of the intestinal epithelial TJ can facilitate the influx of noxious

microbial and other luminal contents in the intestinal wall and their subsequent release throughout the body via the vascular and lymphatic channels. This can, in turn, lead to local and systemic inflammation and accumulation of uremic toxins generated by the gut microbiota.

The present study revealed that at clinically relevant concentrations, urea significantly lowered the TER in the TJ-forming polarized human colonic epithelial cell monolayer in vitro. The effect of urea was dramatically amplified by urease, which was used to simulate the effect of urease-possessing microbial species in the colon. These observations illustrate the potential contribution of influx of urea into the intestinal tract in the pathogenesis of the uremia-induced intestinal barrier dysfunction in the uremic patients. The observed fall in the TER was accompanied by and was largely due to the urea concentration-dependent depletion of occludin, claudin-1, and ZO-1 which are the key protein constituents of the epithelial TJ. The erosive effect of urea on the measured TJ proteins was greatly intensified in the presence of urease leading to near-complete depletion of ZO-1 and occludin and severe reduction in claudin-1 abundance. As noted above, accumulation of urea in the intra- and extracellular fluid compartments in patients and animals with advanced CKD results in its heavy influx into the gastrointestinal tract via passive diffusion and incorporation in the glandular secretions [7–9]. As noted earlier, within the intestinal lumen urea is hydrolyzed spontaneously and by microbial urease forming large quantities of ammonia [8, 9]. Ammonia is, in turn, converted to ammonium hydroxide which is a caustic base capable of causing cytotoxicity and tissue damage. Recent studies conducted

by our group have shown marked changes in the composition of microbial flora in humans and animals with advanced CKD [21]. Five of the 10 microbial families which were most abundant in the CKD patients possessed functional urease [2, 21]. Thus massive influx of urea into the gastrointestinal tract together with the dominance of urease-possessing bacteria in the uremic individuals work in concert to promote formation of ammonia and ammonium hydroxide in the intestinal tract [8, 9]. This can account for the dramatic impairment of the barrier function and destruction of TJ apparatus observed in the present in vitro experiments designed to simulate the effect of uremia in vivo.

In conclusion, the CKD-induced disruption of intestinal TJ and impairment of the gut barrier function is, in part, mediated by urea which heretofore was considered to be a nontoxic uremic retained metabolite. These findings illustrate the adverse effect of the high urea levels and reveal a novel mechanism for the salutary effect of urea-lowering strategies such as low-protein diet and longer and more frequent dialysis regimens in patients with advanced CKD.

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Disclosure Statement

The authors have no conflict of interest.

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