Impaired intestinal barrier function in a mouse model of hyperuricemia

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Abstract. Previous studies have demonstrated the effects of hyperuricemia on the damage to target organs, including the kidneys, joints and the heart. However, it is unclear whether hyperuricemia results in damage to the intestines. The aim of the present study was to investigate intestinal barrier dysfunction in a mouse model of hyperuricemia constructed by knocking out the urate oxidase (Uox) gene. The morphology of the intestine was assessed via hematoxylin and eosin, and alcian blue staining. The serum and intestinal tissue levels of uric acid, tumor necrosis factor (TNF)-α and interleukin (IL)-6, in addition to the presence of uremic toxins, were assessed. The levels of diamine oxidase (DAO), D-lactate (D-LAC) and endotoxins in the serum, which are markers of the intestinal permeability, were measured using ELISA. The expression of the intestinal tight junction proteins zona occludens-1 (ZO-1) and occludin were detected by reverse transcription-quantitative polymerase chain reaction, western blotting and immunohistochemical analysis. The Uox-knockout mice spontaneously developed hyperuricemia. Histopathological analysis indicated intestinal defects including sparse villi, mucosal edema and a declining mucus layer in hyperuricemic mice. The expression levels of ZO-1 and occludin in the intestines were downregulated, and the serum levels of DAO, D-LAC and endotoxins were higher in the hyperuricemic mice, compared with control mice. The serum and intestinal tissue levels of IL-6 and TNF-α were significantly increased. Additionally, the expression levels of the serum uremic toxins, serum creatinine, blood urea nitrogen were significantly increased in hyperuricemic mice compared with the control mice, while only a marked increase in indoxyl sulfate (IS) and p-cresol sulfate was reported. Collectively, the results of the present study suggested that intestinal barrier dysfunction and subsequent enhanced intestinal permeability may occur as a result of hyperuricemia in mice. Furthermore, we proposed that the loss of intestinal epithelium barrier function may be associated with uric acid-induced inflammatory responses; however, further investigation is required.

Introduction

Hyperuricemia is a common metabolic disease caused by a disorder of purine metabolism in humans (1). In the majority of mammals, uric acid (UA) is converted to the more soluble form allantoin by urate oxidase (Uox), which is removed from the body. However, the Uox gene has undergone mutational silencing during the evolution of humans and hominoid apes; thus, humans are vulnerable to hyperuricemia (2). Hyperuricemia has been a global public health concern with an increasing incidence and prevalence in recent years (3,4). Hyperuricemia is a major risk factor for gout (5). In total, 10-20% patients with hyperuricemia may progress to gout (5). In addition, studies have suggested that hyperuricemia may also contribute to the development and pathogenesis of metabolic and systemic pathologic diseases, including diabetes, metabolic syndrome, hypertension, atherosclerosis and chronic kidney disease (6,7).

The intestine is the primary organ responsible for the clearance of UA and one third of UA in the blood is removed through the intestines (8,9). Certain patients with hyperuricemia present with gastrointestinal symptoms; however, the precise effects of hyperuricemia on the intestines remain unclear. A previous study showed that hyperuricemia is associated with an increased risk of colorectal carcinoma (10). Additionally, it was documented that supplementing exogenous UA via an intra-intestinal injection induces the influx of heterophils into the intestines of rabbits (11). Nevertheless, supplementing exogenous UA in experimental animals within a short time does not mimic human hyperuricemia adequately as the majority of cases of hyperuricemia in humans are sustained for at least a few years. In our previous study, a mouse model of hyperuricemia was established by knocking out the Uox
gene (12). The Uox-knockout (KO) mice spontaneously developed hyperuricemia (serum UA concentration >7 mg/dl) and survived ≤62 weeks (12). Therefore, we employed this model to investigate the effects of hyperuricemia effects on the intestines. In the present study, intestinal barrier function and intestinal permeability was assessed in the hyperuricemia mouse model, and the serum and intestinal tissue levels of UA, tumor necrosis factor (TNF)-α and interleukin (IL)-6, as well as serum uremic toxins were measured. We aimed to investigate intestinal barrier dysfunction in a mouse model of hyperuricemia, and further understand the underlying mechanisms.

Materials and methods

Animal model. In our previous study, a hyperuricemia mouse model was generated by knocking out the Uox gene using the transcription activator-like effector nuclease technique (12). The male Uox-KO mice and their wild-type (WT) counterparts (n=8 in each group) were used in the present study. The mice from 4 weeks (weight, 13-15 g) to 24 weeks (weight, 28-32 g) after birth were maintained individually in a temperature-controlled room at 22-24°C and 40-60% relative humidity under a 12-h light/dark cycle with ad libitum access to food and water. Blood and urine samples of all the animals were collected in the morning. The blood samples were centrifuged at 1,511 x g for 15 min at 4°C to obtain the serum. The mice were euthanized at 24 weeks after birth using CO₂ with 25% volume displacement per minute, and mortality was confirmed by observing respiration and by using the corneal reflection method. The intestines were harvested, and a section of the intestinal tissues were cut and washed with cold normal saline to remove blood; after drying on filter papers at room temperature, the intestinal samples were weighed, then immediately homogenized on ice. The homogenates were centrifuged at 5,000 x g for 10 min at 4°C to obtain the supernatants. All procedures were performed with the approval of The Animal Research Ethics Committee of The Affiliated Hospital of Qingdao University (approval no. AHQU20140612A).

Measurement of UA. Serum UA levels were analyzed in the mice from 4 to 24 weeks, and the levels of UA in the urine and the supernatant of the intestinal homogenate were measured at 24 weeks of age. The levels of UA in serum, urine and supernatant were measured using a UA determination kit R-i/ii (Beijing Leadman Biochemistry Co., Ltd.). The detection process was completed with an automatic biochemical analyzer (Toshiba Corporation, Minato, Japan) according to the manufacturer's protocols.

Histological examination. Intestinal samples were fixed by immersion in 10% neutral formalin at room temperature for 48 h, and subsequently embedded in paraffin. The paraffin block of tissue was sliced into 4-µm-thick sections, and stained with hematoxylin and eosin for 5 min at room temperature. Alcian blue staining was performed as described previously; the sections were stained with Alcian blue for 30 min at room temperature (13). Pathological changes in the intestine were observed under a light microscope (magnification, x400; Axiovert 200; Zeiss AG) and evaluated by an experienced pathologist.

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Total RNA was extracted from intestinal tissue using RNAiso Plus (Takara Bio, Inc.) and was reverse-transcribed into cDNA using a PrimeScript RT reagent kit with gDNA Eraser (Takara Bio, Inc.), according to the manufacturer's protocols. qPCR reaction was performed in a 25 µl reaction volume, using TB Green Premix Ex Taq reagent (Takara Bio, Inc.) using a fluorescent PCR device (CFX96, Bio-Rad Laboratories, Inc.). The thermocycling conditions were: i) 95°C For 30 sec; and ii) 40 cycles of 95°C for 5 sec and 60°C for 30 sec. The primer sequences were as follows: Zona occludens-1 (ZO-1) forward, 5'GCCACCTCTGGTCCAGCTCTTC-3', reverse, 5'ACCCGGAGTGAGGTTTTCT-3'; Ocludin forward, 5'CCTCAATTGGCAAGTGAAT-3', reverse, 5'GCTCCACCTGTCGTGATG-3', and β-actin forward, 5'CTCTGGAGAAAGCTATGA-3' and reverse, 5'GGCATAGGGTCTTACCGATG-3'. All samples were run in triplicate in a single 96-well reaction plate. The relative expression levels were analyzed using the 2^ΔΔCT method (14,15).

Immunohistochemistry detection of tight junction proteins. Immunohistochemistry for ZO-1 and occludin in intestinal tissue was performed as described previously (16). Briefly, the intestinal tissues from the Uox-KO mice and the WT mice were fixed and embedded in paraffin as aforementioned, and 4-µm sections were cut from the paraffin blocks. The sections were dewaxed in xylene (Beijing Zhongshan Golden Bridge Biotechnology Co., Ltd.; OriGene Technologies, Inc.) and antigen retrieval was performed using 10 mmol/l citrate buffer (pH 6.0; Beijing Zhongshan Golden Bridge Biotechnology Co., Ltd.; OriGene Technologies, Inc.) at 120°C for 3 min. Endogenous peroxidases activity were inhibited with 3% hydrogen peroxide for 10 min at room temperature. Then, the sections were incubated at 4°C overnight using rabbit anti-ZO-1 (1:400; cat. no. bs-1329R; BISOSS Biotech Co., Ltd) and rabbit anti-occludin (1:300; cat. no. A2601; ABlonatal Biotech Co., Ltd.). As a negative control, sections were incubated with PBS alone. Subsequently, the sections were incubated with a goat-anti-rabbit immune-globulin G secondary antibody (1:500; cat. no. PV9001; Beijing Zhongshan Golden Bridge Biotechnology Co., Ltd.; OriGene Technologies, Inc.) at room temperature for 1 h and stained with diaminobenzidine (OriGene Technologies, Inc.) at room temperature for 5 min. The sections were counterstained with hematoxylin at room temperature for 5 min and analysis was conducted with an optical microscope (magnification, x400). The immunoreactive positive signals appeared brown.

Western blotting analysis of tight junction proteins. Western blotting was performed as previously described (17). Briefly, total proteins were extracted from intestinal tissues using a protein lysis buffer (50 mM Tris-HCl, 150 mM NaCl, 1% Triton X-100, 0.5% sodium deoxycholate, 0.1% SDS; Applygene) and the protein concentration was measured using a bicinchoninic acid assay kit (Pierce; Thermo Fisher Scientific, Inc.). A total of 30 µg of protein from each sample were resolved using a 10% SDS-PAGE gel and transferred to a polyvinylidene difluoride
membrane (EMD Millipore), and blocked using 5% nonfat milk at room temperature for 1 h. Membranes were incubated with primary antibodies at 4°C overnight: Rabbit anti-ZO-1 (1:1,000; cat. no. 5406, Cell Signaling Technology, Inc.), rabbit anti-occludin (1:1,000; cat. no. A2601; ABclonal Biotech Co., Ltd.) and anti-GAPDH (1:1,000; cat. no. 8884; Cell Signaling Technology, Inc.). Subsequently, the membranes were incubated at 4°C with a horseradish peroxidase-conjugated secondary antibody (1:10,000; cat. no. 7074; Cell Signaling Technology, Inc.) for 1 h. An FR-980 Biological Image Analysis System (Furi) was used to detect immunoreactive bands and densitometry analysis was performed using ImageJ (version 1.8.0; National Institutes of Health).

Quantification of intestinal permeability. Diamine oxidase (DAO), D-lactate (D-LAC), and endotoxin levels in the serum are commonly regarded as indirect indicators of intestinal permeability (18). The concentrations of serum DAO, D-LAC and endotoxin were determined using ELISA kits (DAO ELISA Kit, cat. no. CSB-E10090m, Cusabio Technology LLC; D-lactate ELISA Kit, cat. no. JL48176, Shanghai Future Industrial Limited By Share Ltd.; endotoxin ELISA Kit, cat. no. CSB-E13066m, Cusabio Technology LLC), according to the manufacturer's protocols.

Analysis of serum uremic toxins. The levels of serum creatinine (SCR) and blood urea nitrogen (BUN) in serum samples (200 ul) were measured using an automatic biochemical analyzer (Toshiba Corporation, Minato, Japan) to assess kidney function. The detection of indoxyl sulfate (IS) and p-cresol sulfate (PCS) levels in the serum were performed using a mouse indoxyl sulfate ELISA Kit and p-cresol sulfate ELISA Kit (cat. no. JL44175 and cat. no. JL48074, Shanghai Future Industrial Limited By Share Ltd.) according to the manufacturer's protocols.

ELISA measurement of cytokines in the serum and intestinal tissues. The concentrations of TNF-α and IL-6 in the serum and intestinal tissues were quantified using the mouse TNF-α ELISA kit and mouse IL-6 ELISA kit, respectively (cat. nos. RK00027 and RK00008, ABclonal Biotech Co., Ltd.) according to the manufacturer's protocols.

Statistical analysis. All data were statistically analyzed using SPSS 21 (IBM Corp.) and GraphPad Prism 6 (GraphPad Software, Inc.). All values were presented as the mean ± standard error of the mean. Experiments were performed in triplicate. Quantitative data was assessed using independent-sample t-tests or one-way analysis of variance followed by a Tukey's post-hoc test. P<0.05 was considered to indicate a statistically significant difference.

Results

UA levels are elevated in the serum, urine and intestinal tissues of the Uox-KO mice. The Uox-KO mice had significantly elevated serum UA levels between 8.89-10.15 mg/dl; however, the levels appeared to be stable and 3-4 fold higher compared with WT mice (Fig. 1A and B). The urinary UA concentration in Uox-KO mice was ~80 mg/dl, ~10 fold higher compared with the WT littermate controls (Fig. 1C). The UA concentration in the intestinal tissues was ~1,310 µg/g in the hyperuricemic mice, which was significantly increased than that of the control group (Fig. 1D).

Histological observation of intestinal damage in the Uox-KO mice. Using light microscopy, regional epithelial shedding, fewer and fractured villi in the intestine of the Uox-KO mice were observed. Furthermore, mucosal and submucosal edema were observed in the Uox-KO mice. Conversely, the intestinal epithelial structure was intact, and no notable pathological changes were detected in the controls (Fig. 2A). Additionally, Uox-KO mice exhibited a notable reduction in mucus when compared with the WT mice as determined using alcin blue staining (Fig. 2B). These histological results suggested that the structure of the intestinal epithelium was disrupted in Uox-KO mice.

Expression of tight junction proteins is downregulated in Uox-KO mice. ZO-1 and occludin are considered to be the most important tight junction proteins, and contribute to the intestinal mucosal barrier function (19). Therefore, the mRNA and protein expression levels of ZO-1 and occluding were measured in the intestinal tissues. The results of RT-qPCR revealed that the mRNA expression levels of ZO-1 and occluding were significantly downregulated in the Uox-KO mice compared with the control (Fig. 3A). Consistent with the mRNA expression results, the protein expression levels of ZO-1 and occluding were significantly downregulated in the Uox-KO mice compared with the control as determined by western blotting (Fig. 3C). Immunohistochemistry demonstrated that the degree of positive staining for ZO-1 and occludin observed in the Uox-KO mice was markedly reduced compared with the control (Fig. 3B).

Intestinal permeability is increased in the Uox-KO mice. Impairments in the intestinal barrier were determined by intestinal permeability. Serum DAO, D-lactate and endotoxin levels were measured as markers of intestinal permeability. The results revealed that the serum levels of DAO, D-LAC and endotoxin were significantly increased in the Uox-KO mice compared with the WT mice (P<0.05; Fig. 4).

Levels of uremic toxin and cytokines are increased in the Uox-KO mice. To determine the potential underlying mechanism of intestinal barrier dysfunction in hyperuricemia, the serum uremic levels and cytokine levels were measured in the Uox-KO and WT mice. As presented in Fig. 5, the serum uremic toxins levels of SCR and BUN were significantly increased in Uox-KO mice compared with the control, whereas the levels of IS and PCS exhibited marked increases in the Uox-KO mice (Fig. 5). Thus, it is unlikely that the extent of damage observed in the hyperuricemic mice resulted from the slight increases in IS and PCS.

Inflammatory cytokines are known to contribute to intestinal damage (20). The levels of TNF-α and IL-6 in the serum and the intestinal homogenates were significantly higher in the Uox-KO mice compared with the WT mice (Fig. 6). The data indicated that the intestinal mucosal barrier damage in the hyperuricemic mice may be associated with the induction of the inflammatory response.
GUO et al: IMPAIRED INTESTINAL BARRIER FUNCTION IN HYPERURICEMIA

Figure 1. UA levels in serum, urine and intestinal tissues of the Uox-KO mice are increased. (A) Serum UA levels were higher in the Uox-KO mice between 4 and 24 weeks. The levels of UA in the (B) serum, (C) urine and (D) intestinal tissues of hyperuricemic mice were significantly increased compared with the WT littermate controls *P<0.05 vs. WT. UA, uric acid; UOX, urate oxidase; KO, knockout; WT, wild-type.

Figure 2. Histological analysis of intestinal tissue in the Uox-KO mice. (A) Hematoxylin and eosin staining showed that the intestines of hyperuricemic mice exhibited defects, including sparse intestinal villi, mucosal and submucosal edema. (B) Alcian blue staining revealed that the intestinal mucus layer was notably reduced in hyperuricemic mice compared with the control. Scale bar=50 µm. UOX, urate oxidase; KO, knockout; WT, wild-type.
Discussion

Numerous studies have confirmed that elevated serum UA levels contribute to the damage of multiple organs, including the kidneys and heart, and joints (5-7). Furthermore, a recent study indicated that UA has a role in impairment of the integrity of vascular endothelial cells (21). Notably, the structure and physiological characteristics between the vascular and
Figure 5. Uremic toxins levels are increased in the Uox-KO mice. Uremic toxins levels of SCR, BUN were significantly elevated, while those of IS and PCS were notably increased in Uox-KO mice compared with the control group. *P<0.05 vs. WT. UOX, urate oxidase; KO, knockout; SCR, serum creatinine; BUN, blood urea nitrogen; IS, indoxyl sulfate; PCS, p-cresol sulfate; WT, wild-type.

Figure 6. Cytokine levels are elevated in Uox-KO mice. The levels of TNF-α and IL-6 in the (A) serum (A) and (B) intestinal tissue were significantly increased in the Uox-KO mice compared with the control group. *P<0.05 vs. WT. UOX, urate oxidase; KO, knockout; TNF-α, tumor necrosis factor-α; IL, interleukin; WT, wild-type.
intestinal epithelium are homologous and similar; however, few studies have addressed the possible disruptive effects of hyperuricemia on the intestine. In the present study, the Uox-KO (hyperuricemic) mice exhibited impairments in the intestinal barrier, and pathological changes in the intestines were observed, including sparse intestinal villi, mucosal and submucosal edema, suggesting that hyperuricemia may have interfered with the epithelial integrity of the intestine, contributing to intestinal damage. The findings of the present study may improve current knowledge regarding the intestinal damage caused by hyperuricemia; however, damage to other organs under these conditions requires further investigation.

Hyperuricemic mice models are generally constructed by promoting purine synthesis, inhibiting UA degradation and supplementing exogenous UA, either alone or in combination (22,23). However, the UA levels in these models usually fluctuate by a large margin (23). Therefore, these models may not suited for simulating hyperuricemia in humans. In the present study, a mouse model of hyperuricemia was established by knocking out the Uox gene. Our previous study revealed that these mice exhibited a stably increased level of UA and had developed the symptoms of compromised renal function, hypertension, aberrant glycometabolism and lipometabolism, which are similar to those observed in patients with hyperuricemia (12). In the present study, impaired intestinal barrier function was also observed in these gene deficient mice. Collectively, these data suggest that the mouse model may be a more suitable model investigating hyperuricemia and its related complications.

The importance of the intestinal epithelium for excretion has been demonstrated (24). Approximately one-third of the circulating UA is cleared via urate transporters in the intestinal enterocytes (25). The intestinal epithelium and the apical junctional complex constitute the intestinal barrier, a dynamic interface between the internal and external environments, which serves a crucial role in the maintenance of intestinal homeostasis (26,27). In the present study, intestinal epithelial damage was observed in Uox-KO mice as detected by histological examination. Tight junctions between the intestinal epithelial cells are involved in the maintenance of the intestinal mucosal barrier function (28). ZO-1 and occludin are considered to be two of the primary tight junction proteins (19). In the present study, a significant decrease in the expression of ZO-1 and occluding in the intestine of Uox-KO mice was reported compared with the control. The mucus layer on the epithelial surface is also an essential component of the intestinal barrier (29). The thickness of the mucus layer was also decreased in hyperuricemic mice compared with the WT mice, as determined by alcian blue staining. These results suggested that intestinal mucosal barrier function may have been damaged in the hyperuricemic mice.

Impairments in the intestinal mucosal barrier manifest as increased intestinal permeability (30,31). DAO is an intracellular enzyme in the intestinal epithelium, whereas D-LAC and endotoxin are bacterial metabolites produced by the intestinal flora (32). When the permeability of the intestine is abnormally increased as a result of a sort of disruption, DAO, D-LAC and endotoxin in the lumen easily pass through the intestinal mucosa and into the peripheral blood (32,33). In the present study, increases in the serum levels of DAO, D-LAC and endotoxins were observed in the hyperuricemic mice compared with the controls, suggesting enhanced intestinal permeability in the hyperuricemic mice.

The present study suggested that the intestinal barrier was damaged and intestinal permeability was increased in the hyperuricemia mouse model; however, the exact mechanism underlying injury had not been determined. Previous studies demonstrated a high concentration of uremic toxins, including IS, PCS and advanced glycation end products could result in the destruction of the intestinal barrier structure in end-stage renal disease (34-36). Additionally, chronic renal failure induced by 5/6 nephrectomy results in increased levels of uremic toxins which damages the intestinal mucosal barrier in rats (37). Our previous study indicated that kidney impairment was similarly detected in the hyperuricemic Uox-KO mice, in which collapsed and necrotic nephrons and focal tubulointerstitial fibrosis were observed (12). In order to determine whether intestinal injury was caused by kidney damage in the hyperuricemic mice, the levels of uremic toxins, such as IS and PCS in the serum were measured. The results indicated that their serum levels were markedly increased in the hyperuricemic group compared with the controls. The effects of notably increased uremic toxin levels on intestinal damage may not fully account for the observed intestinal injury in these models.

Numerous studies have demonstrated that the increased production of inflammatory cytokines has a negative impact on intestinal mucosal barrier and epithelial function (20,38). For example, Xin et al (39) showed that impaired intestinal barrier in a rat model of chronic obstructive pulmonary disease was associated with the intestinal inflammatory response, in which elevated levels of the cytokines interferon-γ, TNF-α and IL-8 were observed. In vitro experiments have also revealed that TNF-α may result in structural changes to the tight junctions by decreasing occludin and ZO-1 expression (40). In humans, patients with hyperuricemia were found to have higher levels of inflammatory cytokine levels in the serum compared with the healthy controls (41). Similar results were observed in rabbits, in which the addition of exogenous UA to the intestinal lumen promoted an intestinal inflammatory reaction (11). In the present study, elevated production of the inflammatory cytokines TNF-α and IL-6 in the serum and intestinal homogenates were detected in the hyperuricemic mice. These results suggest there was a positive association between the increased levels of inflammatory cytokines and intestinal epithelial tight junction dysfunction. Therefore, hyperuricemia may affect intestinal barrier function through the initiation of inflammatory responses in the intestine. Further studies are required to determine the precise mechanism underlying intestinal mucosal barrier damage in hyperuricemic mice.

In conclusion, the present study demonstrated the presence of a damaged intestinal barrier and enhanced intestinal permeability in a mouse model of hyperuricemia. The impaired intestinal barrier function may be associated with the inflammatory process induced by UA. The findings of the present study may provide novel insight into organ damage as a consequence of hyperuricemia.
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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

YG and ZT designed the study. YG, HL and YY performed the experiments. HL, ZL, CL, YC and CJ contributed to data analysis. YG and HL drafted the manuscript. YG, YY, and ZT revised the manuscript. All authors have given approval to the final version of the manuscript.

Ethics approval and consent to participate

The present study was approved by the Animal Research Ethics Committee of the Affiliated Hospital of Qingdao University and the approval number was AHQU20140612A.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

References


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