# Early exposure to germs modifies kidney damage and inflammation after experimental ischemia-reperfusion injury

Hye Ryoun Jang,<sup>1</sup> Maria Teresa Gandolfo,<sup>2</sup> Gang Jee Ko,<sup>1</sup> Shailesh Satpute,<sup>1</sup> Lorraine Racusen,<sup>2</sup> and Hamid Rabb<sup>1</sup>

<sup>1</sup>Nephrology Division, Department of Medicine, and <sup>2</sup>Department of Pathology, Johns Hopkins University School of Medicine, Baltimore, Maryland

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Jang HR, Gandolfo MT, Ko GJ, Satpute S, Racusen L, Rabb H. Early exposure to germs modifies kidney damage and inflammation after experimental ischemia-reperfusion injury. Am J Physiol Renal Physiol 297: F1457-F1465, 2009. First published August 12, 2009; doi:10.1152/ajprenal.90769.2008.-Kidney ischemia-reperfusion injury (IRI) is, in part, mediated by immune and inflammatory factors. Since microbial stimuli are known to alter immune and inflammatory responses, we hypothesized that differences in perinatal microbial status would modify renal injury following IRI. We performed bilateral renal IRI on 6-wk-old germ-free and control mice and studied the effects on kidney lymphocyte trafficking, cytokines, function, and structure. Compared with control mice, normal kidneys of germ-free mice exhibited more NKT cells and lower IL-4 levels. Postischemia, more CD8 T cells trafficked into postischemic kidneys of germ-free mice compared with control mice. Renal structural injury and functional decline following IRI were more severe in germ-free mice compared with control mice. When germ-free mice were conventionalized with the addition of bacteria to their diet, the extent of renal injury after IRI became equivalent to age-matched control mice, with similar numbers and phenotypes of T cells and NKT cells, as well as cytokine expression in both normal kidneys and postischemic kidneys of conventionalized germ-free mice and age-matched control mice. Thus microbial stimuli influence the phenotype of renal lymphocytes and the expression of cytokines of normal kidneys and also modulate the outcome of IRI.

germ-free status; immune modulation

ISCHEMIA-REPERFUSION INJURY (IRI) is a leading cause of acute kidney injury (AKI) in both native and transplanted kidneys, worsening both early and late allograft outcomes (39). IRIinduced AKI is associated with a robust inflammatory response caused by both innate and adaptive immune components in tubulointerstitial tissues (12, 23). The concept of renal IRI as an inflammatory disease mediated by components of the immune system is supported by a large body of work demonstrating the renoprotective effects of therapy targeting complements, chemokines, and adhesion molecules (13, 34, 38), and also lymphocytes (26, 48). While most work to date has focused on the different components of the immune system, there has been minimal work on elucidating the basic drivers of the immune response and whether there is a role for environmental factors such as microbial stimuli on the tissue responses in kidney IRI.

Several epidemiological studies have revealed a steep increase in the prevalence of allergies and autoimmune diseases in Western societies over the past few decades (1, 29, 37, 42). The "hygiene hypothesis" was suggested to explain the increasing prevalence of allergic disease and proposed that reduced exposure to infections in early childhood due to several factors such as improved personal hygiene, standards of living, and diminished family size may have resulted in increased risk of allergic diseases (37). Two major mechanisms, missing immune deviation and reduced immune regulation, were proposed to explain the hygiene hypothesis (21, 28, 45, 46). The increased prevalence of autoimmune diseases was also linked to an altered immune system influenced by the change in the microbial burden (4). Lymphocytes, especially T cells, came to the forefront of studies investigating the relationship between the microbial burden and the changes in the immune system involved in the increased prevalence of allergy and autoimmune disease.

The pathophysiological role of T cells has been identified in renal IRI (7, 26, 48). Direct or indirect blockade of T cells attenuated renal injury both functionally and structurally in a murine renal IRI model. CD4 and CD8 double-knockout mice were significantly protected from early renal injury in vivo, while T cells exhibited increased adherence to renal tubular epithelial cells after in vitro hypoxia reoxygenation (26). Athymic *nu/nu* mice, another T cell knockout mouse strain, were also protected from initial injury, and the adoptive transfer of T cells into these mice reconstituted renal injury following IRI (5). T cell-targeting medications such as FK506 and mycophenolate mofetil were found to significantly attenuate renal injury after IRI, supporting the importance of T cells in the pathogenesis of renal IRI. T cells have also been demonstrated to participate in IRI of the liver, lung, brain, and intestine (49).

Based on the data that T cells are a mediator of renal IRI and that the immune response during renal IRI could have similar features to allergy and autoimmune disease, we hypothesized that "normal nonpathogenic" germ exposure could modify renal injury following IRI. We first examined the phenotype of lymphocytes and cytokine expression in normal kidneys of control and germ-free mice. We then compared kidney lymphocyte trafficking, structure, and function in germ-free vs.

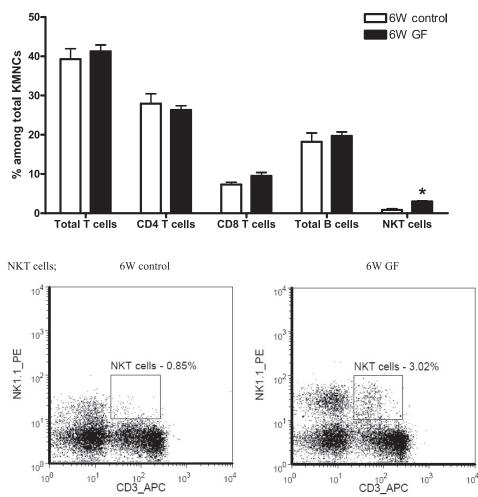
Table 1. Number of kidney-infiltrating mononuclear cells

	6W Control	6W GF	11W Control	11W CV-GF
Day 0	2.36±0.29	$2.44 \pm 0.11$	$2.73 \pm 0.39$	2.39±0.39
Day 1	$1.95 \pm 0.25$	$1.86 \pm 0.25$	$1.80 \pm 0.15$	$1.88 \pm 0.22$

Values are means  $\pm$  SE expressed as  $\times 10^{6}$ . 6W control, 6-wk-old control mice; 6W GF, 6-wk-old germ-free mice; 11W control, 11-wk-old control mice; 11W CV-GF, 11-wk-old conventionalized germ-free mice kept in a conventional specific pathogen-free environment and fed with fecal materials from control mice for 5 wk before ischemia.

Address for reprint requests and other correspondence: H. Rabb, Div. of Nephrology, Johns Hopkins University School of Medicine, Ross Bldg., Rm. 965, 720 Rutland Ave., Baltimore, MD 21205 (e-mail: hrabb1@jhmi.edu).

Fig. 1. Baseline phenotype of kidney lymphocytes in control and germ-free mice. Kidney mononuclear cells (KMNCs) were isolated from naive control and germ-free mice (n =7/group) and analyzed with flow cytometry. There was no difference in the percentages of total T cells, CD4 T cells, CD8 T cells, and total B cells among KMNCs in the kidneys of control and germ-free mice before surgery. However, there were more NKT cells in the kidneys of germ-free mice. 6W control, 6-wkold control mice; 6W GF, 6-wk-old germ-free mice. \*P < 0.05 compared with 6W control.

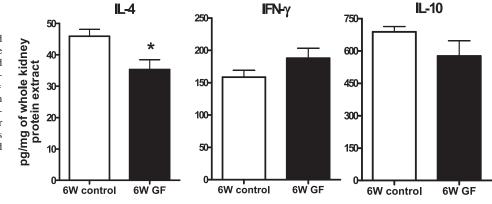


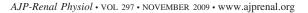
control mice after renal IRI. Finally, we "conventionalized" germ-free mice with microbes and examined how this changed the kidney IRI responses in these mice. These studies open a new line of investigation into kidney IRI that will focus on germ-immune interactions on the outcome of AKI.

#### MATERIALS AND METHODS

*Mice and germ conventionalization technique.* Germ-free mice on a Swiss-Webster background were born by Caesarean section and raised in flexible plastic isolators using strict gnotobiotic techniques. Control Swiss-Webster mice were delivered by normal vaginal delivery and maintained in a conventional specific pathogen-free (SPF) environment. Male germ-free mice and male Swiss-Webster mice were purchased from Taconic Farms (Hudson, NY) and studied simultaneously. All experiment protocols were approved by the Institutional Animal Care and Use Committee. The first group of 6-wk-old germ-free (6W GF) and 6-wk-old control (6W control) mice was either killed immediately upon arrival or received bilateral IRI surgery after a 24-h acclimation period in a sterile biosafety hood. Mice that were killed were used to compare basal intrarenal-residing lymphocytes and cytokine levels (n = 7/group). Mice that received IRI surgery were maintained with autoclaved chow and water under the biosafety hood for 24 h, followed by death (n = 10/group). Functional

Fig. 2. Baseline levels of IL-4, IFN- $\gamma$ , and IL-10 in kidneys of control and germ-free mice. IL-4, IFN- $\gamma$ , and IL-10 were measured in protein samples extracted from naive kidneys of control and germ-free mice (n = 7/group). IL-4 was significantly decreased in germ-free mice. IL-10 trended lower in germ-free mice, whereas there was a tendency for increased IFN- $\gamma$  expression in renal tissues from germ-free mice. \*P < 0.05 compared with 6W control.





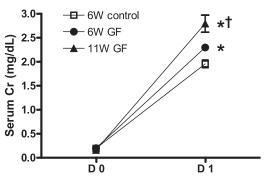


Fig. 3. Germ-free mice show higher serum creatinine than control mice at 24 h after renal ischemia-reperfusion injury (IRI). The serum creatinine (Cr) level was measured to assess the renal function at 24 h after bilateral IRI (n = 10/group). Germ-free mice had higher serum creatinine compared with control mice. 11W GF, 11-wk-old germ free mice; D0, before bilateral IRI; D1, 24 h after bilateral IRI. \*P < 0.01 compared with 6W control. †P < 0.001 compared with 6W GF.

and structural changes of 11-wk-old germ-free (11W GF) mice were also compared with the 6W GF mice at 24 h after bilateral IRI surgery. The second group of germ-free and control mice, which were the same age as the first group and also delivered at the same time, was kept in a conventional SPF environment for 5 wk. Fecal material from control mice was added to the germ-free mice with usual chow to conventionalize the germ-free mice. After 5 wk, conventionalized germ-free (11W CV-GF) mice and age-matched control (11W control) mice were either killed to analyze basal lymphocytes and cytokine changes

(n = 5/group) or received bilateral IRI surgery with the same protocol as the first group (n = 10/group).

*Quantification of microbial burden*. Stool from control, 6W GF, and 11W CV-GF mice was collected with rectal swabs using sterile cotton tips, weighed, and diluted with sterile saline. Stool (8 mg) was inoculated in fluid thioglycollate medium (BD Diagnostics, Sparks, MD), and 0.16 mg of each sample was placed on trypticase soy agar with 5% sheep blood (TSA II)/MacConkey II agar plate medium (BD Diagnostics). Test tubes containing the inoculated thioglycollate medium were kept in 37°C, and absorbance at 490 nm was measured at 48 h after inoculation. Plate mediums were kept at room temperature, and colony-forming units (cfu) were counted at 72 h after inoculation.

*Renal ischemia-reperfusion model.* An established murine renal IRI model was used (27). Briefly, mice were anesthetized with an intraperitoneal injection of pentobarbital sodium (75 mg/kg). Following an abdominal midline incision, both renal pedicles were bluntly dissected and clamped with a microvascular clamp (Roboz Surgical Instrument, Gaithersburg, MD) for 35 min. During the procedure, 2 ml of sterile saline at 40°C (1 ml during ischemia and 1 ml during reperfusion) was instilled into the peritoneal cavity. After the clamps were removed, the wounds were sutured and the mice were allowed to recover with free access to chow and water. Throughout the procedure, mice were placed on a heating pad (40°C).

Assessment of renal function. Blood samples were obtained from the tail vein before and at 24 h after renal IRI. Serum creatinine levels (mg/dl) were measured with a Cobas Mira plus autoanalyzer (Roche Diagnostics, Indianapolis, IN).

*Tissue histological analysis.* Mice were killed at 24 h after IRI, and kidneys were harvested after exsanguination. Tissue samples were

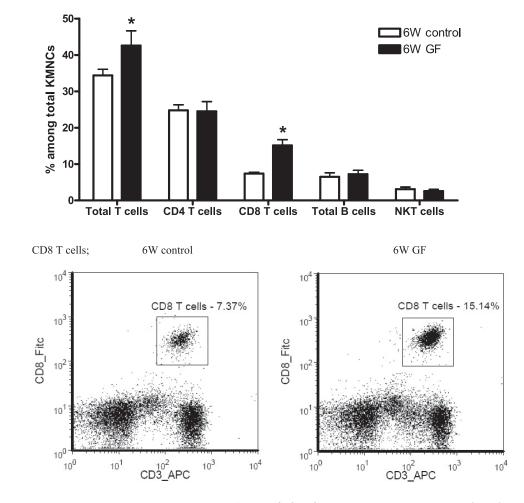
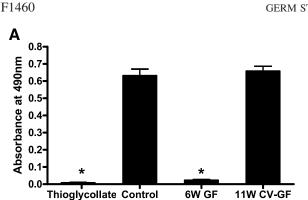


Fig. 4. More CD8 T cells are trafficked into the postischemic kidneys of germ-free mice. Increased trafficking of CD8 T cells into the postischemic kidneys of germ-free mice caused the increase in total T cell trafficking. There was no difference in the percentages of total CD4 T cells, total B cells, and NKT cells among KMNCs isolated from the 2 groups (n = 10/group). \*P < 0.05 compared with 6W control.

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Fig. 5. Effect of conventionalization on intestinal flora of germ-free mice. A: to conventionalize, germ-free mice were kept in a conventional specific pathogen-free (SPF) environment and were fed with fecal materials from control mice for 5 wk. The absorbance of thioglycollate medium with no inoculation and that of thioglycollate medium inoculated with germ-free mice stool were similar. The absorbance of thioglycollate medium inoculated with control mice tool and 11W CV-GF mice stool was significantly higher compared with thioglycollate medium with no inoculation. B: the stool culture of 11W CV-GF mice showed a similar degree of microbial burden as control mice. Thioglycollate, thioglycollate medium with no inoculation; control, stool culture from 6W control mice; 6W GF, stool culture from 11-wk-old conventionalized germ-free mice.\*P < 0.001 compared with both control and 11W CV-GF.

fixed with 10% buffered formalin followed by paraffin embedding, and afterward renal sections were stained with hematoxylin and eosin (H&E). Renal tubular damage was scored in a blinded fashion by a renal pathologist.

Isolation of kidney-infiltrating mononuclear cells. After anesthesia, mice were perfused with  $\sim 30$  ml of warm sterile saline, and kidneys were harvested. Kidney mononuclear cells (KMNCs) were isolated according to the previously described method (3). Briefly, decapsulated kidneys were immersed in RPMI buffer (Mediatech, Manassas, VA) containing 5% fetal bovine serum and disrupted mechanically using a Stomacher 80 Biomaster (Seward, Worthing, West Sussex, UK). Samples were strained, washed, and resuspended in 36% Percoll (Amersham Pharmacia Biotech, Piscataway, NJ) followed by gentle overlaying onto 72% Percoll. After centrifugation at 1,000 g for 30 min at room temperature, KMNCs were collected from the Percoll interface, washed twice, and counted on a hemocytometer using trypan blue exclusion.

*Flow cytometric analysis of KMNCs.* Isolated KMNCs were preincubated with anti-CD16/CD32 Fc receptor-blocking antibody for 10 min to minimize nonspecific antibody binding. Cells were then incubated with various combinations of anti-mouse antibodies (all from BD Biosciences except for anti-mouse PE-conjugated anti-CD25 antibody from eBioscience) for 25 min at 4°C, washed with FACS buffer, and fixed using a 1% paraformaldehyde solution. Four-color immunofluorescence staining was acquired and analyzed using a FACSCalibur instrument (BD Biosciences, San Jose, CA) and FCS Express 3 (De Novo Software, Los Angeles, CA), respectively. Each assay included at least 10,000 gated events.

Bioplex protein array system. A panel of cytokines was measured in whole kidney protein extracts obtained from naive control and germ-free mice with the Bioplex Protein Array system (Bio-Rad, Hercules, CA). This is a multiplexed, particle-based, flow cytometric assay that utilizes anti-cytokine monoclonal antibodies linked to microspheres incorporating distinct properties of two fluorescent dyes. Our assay was designed to detect and quantify IL-4, IL-10, and IFN- $\gamma$ . Each cytokine value was normalized by dividing the raw cytokine concentration (pg/ml) with kidney protein concentration (mg/ml) measured by a Bradford assay.

Statistical analyses. All data are expressed as means  $\pm$  SE. Group means were compared with a Mann-Whitney test using SPSS 12.0K and ANOVA followed by a Newman-Keuls post hoc analysis using GraphPad Prism version 4. Statistical significance was determined when the *P* value was <0.05.

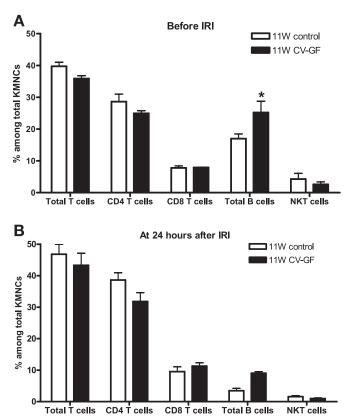


Fig. 6. Conventionalized germ-free mice show similar phenotypes of kidneyresident lymphocytes as age-matched control mice. A: KMNCs isolated from 11W CV-GF mice were analyzed and compared with KMNCs from agematched 11W control mice (n = 5/group). There was no difference in the percentages of total T cells, CD4 T cells, CD8 T cells, and NKT cells. Total B cells were increased in the naive kidneys of 11W CV-GF mice. B: KMNCs were analyzed again at 24 h after bilateral renal IRI (n = 10/group). The percentages of total T cells, CD4 T cells, CD8 T cells, total B cells, and NKT cells were similar between groups. 11W control; 11-wk-old control mice; 11W CV-GF; 11-wk-old conventionalized germ-free mice kept in a conventional SPF environment and fed fecal materials from control mice for 5 wk before ischemia. \*P < 0.05 compared with 11W control.

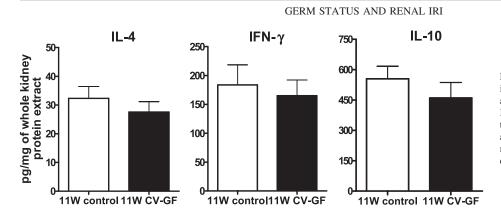


Fig. 7. Baseline levels of IL-4, IFN- $\gamma$ , and IL-10 in kidneys of conventionalized germ-free mice and age-matched control mice. IL-4, IFN- $\gamma$ , and IL-10 were measured in protein samples extracted from naive kidneys of 11W control mice and 11W CV-GF mice (n = 5/group). There was no difference in the expression level of the three cytokines. Groups are defined as in Fig. 6.

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## RESULTS

Baseline characteristics of mononuclear cells in kidneys from control and germ-free mice. There was no difference in the total number of KMNCs among groups on both day 0 (before ischemia) and day 1 after IRI (Table 1). More NKT cells were found in the kidneys of 6W GF mice (6W control vs. 6W GF,  $0.85 \pm 0.34$  vs.  $3.02 \pm 0.13\%$  of KMNCs, P < 0.05), whereas the percentages of total T cells, CD4 T cells, CD8 T cells, and total B cells were similar between the two groups (Fig. 1). KMNCs isolated from 6W GF mice had reduced effector memory CD8 T cells (6W control vs. 6W GF, 26.94  $\pm$ 5.43 vs. 7.58  $\pm$  1.08% of total CD8 T cells, P < 0.05), but similar percentages of effector memory CD4 T cells (control vs. germ free,  $23.69 \pm 2.70$  vs.  $20.22 \pm 1.36\%$  of total CD4 T cells, P > 0.05) compared with 6W control mice. There was no difference in the percentage of regulatory T cells expressing CD4 and CD25 among total KMNCs between the two groups (6W control vs. 6W GF, 0.84  $\pm$  0.10 vs. 1.01  $\pm$  0.17%).

Baseline expression of IL-4, IFN- $\gamma$ , and IL-10 in the renal tissues of control and germ-free mice. Normal kidneys of germ-free mice expressed less IL-4 compared with control mice (6W control vs. 6W GF, 45.95 ± 2.18 vs. 35.29 ± 3.15 pg/mg of total protein extracted from whole kidney, P < 0.05). IL-10 expression trended to be lower, while IFN- $\gamma$  trended to be higher in normal germ-free mice kidneys (Fig. 2).

Germ-free mice had worse kidney function and more CD8 T cells in their postischemic kidneys at 24 h after IRI. Germ-free mice had greater renal functional impairment than control mice at 24 h after bilateral renal IRI (Fig. 3). More total T cells trafficked into the postischemic kidneys of germ-free mice, and this increment of total T cells was accompanied by significantly increased trafficking of CD8 T cells (Fig. 4).

Conventionalization of germ-free mice reconstituted intestinal flora. Stool culture using fluid thioglycollate medium and TSA II/MacConkey II agar plate medium showed similar amounts of microbial burden in control mice and 11W CV-GF mice (Fig. 5). There was no difference in colony counts on both TSA II and MacConkey II agar plates between control and 11W CV-GF groups (TSA II:MacConkey II, control 430  $\pm$ 98.3:427  $\pm$  10.4, 6W GF 0:0, 11W CV-GF 404  $\pm$  47.5:394  $\pm$ 29.5 cfu/mg stool).

*Exposure to germs abolished the difference in basal NKT cell trafficking and cytokine expression in normal kidneys.* KMNCs isolated from 11W control mice and 11W CV-GF mice were similar in the total number and the percentages of T cells, CD4 T cells, CD8 T cells, and NKT cells. However, there

were more B cells in the normal kidneys of 11W CV-GF mice (Fig. 6). There was no difference in IL-4, IFN- $\gamma$ , and IL-10 levels measured in normal kidneys between the two groups (Fig. 7). Both 11W CV-GF mice and 11W control mice showed similar renal functional impairment (Fig. 8) and lymphocyte trafficking into the postischemic kidneys (Fig. 6).

Effects of conventionalization of bacteria on tubular injury following IRI. When GF mice are conventionalized to standard bacterial flora, age increases during the conventionalization process. Prior reports have shown that aging increases susceptibility to kidney IRI (25, 40). We found that 11W control mice showed significantly increased tubular injury in the outer medulla and inner medulla after the same duration of kidney ischemia compared with 6W control mice (Fig. 9, A and C). 11W GF mice also showed significantly increased tubular injury in the outer medulla and inner medulla compared with 6W GF mice (Fig. 9B). After conventionalization, 11W CV-GF mice had less tubular injury in all three areas of the postischemic kidney compared with 11W GF mice. Unexpectedly, the 11W CV-GF mice also had less tubular injury compared with 6W GF mice in the outer medulla and inner medulla (Fig. 9, B and D).

## DISCUSSION

These data demonstrate that germ-free status alters NKT cells and IL-4 content of normal mouse kidneys and worsens kidney histological and functional responses to IRI, with enhanced CD8 T cell trafficking. Furthermore, conventionalizing germ-free mice by feeding intestinal microbiota normalized the

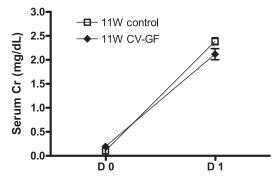
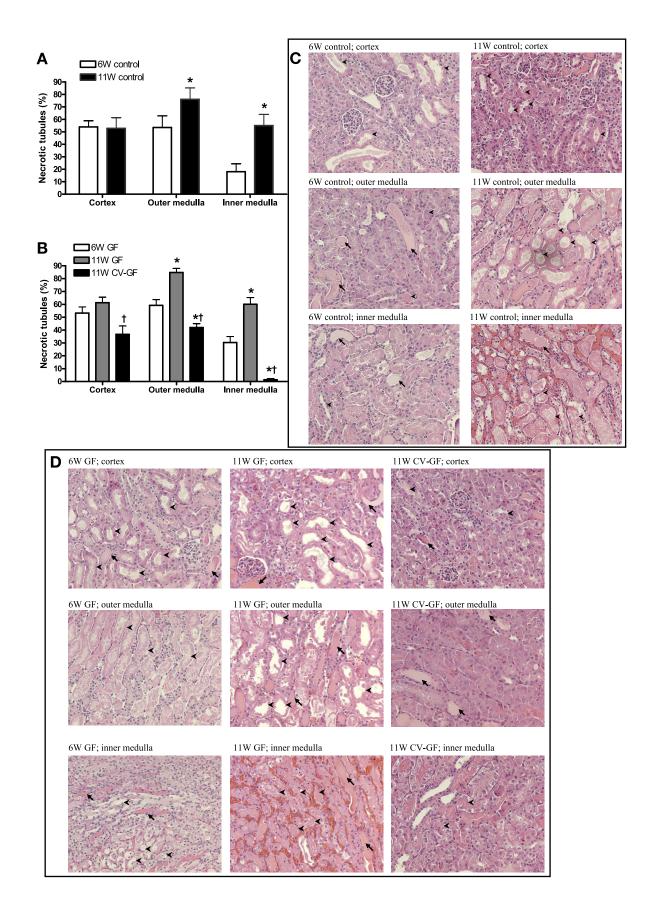


Fig. 8. Introduction of microorganisms in germ-free mice leads to the development of equivalent kidney dysfunction after IRI to age-matched control mice. There was no difference in serum creatinine (Cr) levels measured at 24 h after bilateral renal IRI between the 2 groups. Groups are defined as in Fig. 6.



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kidney IRI functional response, making germ-free mice similar to control mice.

Previous reports demonstrated that germ-free mice were protected from intestinal IRI due to increased production of IL-10 (35, 36). In contrast, we found that the germ-free mice had an enhanced kidney dysfunction and worse structural injury after IRI. At baseline, kidneys of germ-free mice expressed less IL-4 but showed a tendency toward increased IFN- $\gamma$  compared with control mice. These cytokine profiles suggest immune deviation in the germ-free mice, similar to Th1-skewed autoimmune disease. According to the hypothesis of missing immune deviation, the reduced microbial burden during childhood caused by a Westernized lifestyle induces an imbalance of Th1 and Th2 responses, resulting in an exaggerated Th2 response in allergic disease and an exaggerated Th1 response in autoimmune disease (4, 8). Thus our results are more similar to that seen in autoimmune disease rather than allergic disease. Our results are consistent with previous reports on liver and kidney IRI showing that CD4 T cells of the Th1 phenotype are pathogenic and those of the Th2 phenotype are protective (33, 47). A number of studies reported that preceding or concomitant activation of the immune system with whole bacteria or bacterial-derived products led to protection from experimental allergic disease (10, 14, 19, 41) and autoimmune disease (6, 9, 17, 24, 30, 31).

Regulatory T cells and IL-10 have been implemented in the pathogenesis of autoimmune disease and allergic disease as major mediators in the hypothesis of reduced immune regulation. This hypothesis proposes that the increased prevalence of allergic disease is a consequence of reduced stimulation of regulatory T cells caused by a reduced microbial burden during childhood. Despite ongoing debates for the underlying pathogenic mechanisms, several studies have reported the protective role of regulatory T cells in autoimmune disease (15, 22, 32) and allergic disease (11, 20, 50). One recent study reported that there is a small percentage of natural regulatory T cells expressing CD4 and CD25 in naive mice kidneys (2). In the current study, we also found a small number of natural regulatory T cells in normal kidneys of control and germ-free mice, but there was no difference between the two groups, suggesting that the difference in microbial burden does not affect the population of intrarenal regulatory T cells. There was also no significant difference in the level of renal IL-10 expression, implying that reduced immune suppression may not occur in the kidneys of germ-free mice.

We found that NKT cells were increased in the normal kidneys of germ-free mice compared with control mice. NKT cells have been found in normal mice kidneys (2) and post-ischemic kidneys (3). Recently, two studies reported that NKT cells play an important role in early renal injury following IRI (16, 18). NKT cells were suggested to contribute to renal injury by mediating neutrophil infiltration and production of IFN- $\gamma$  (18). Our findings of an increased number of NKT cells and an increased tendency of IFN- $\gamma$  expression in normal kidneys of

germ-free mice are consistent with an enhanced injury response.

Regarding effector memory T cells, the percentage of effector memory CD8 T cells in naive germ-free mice kidney was less than that of naive control mice, whereas there was no difference in the percentage of effector memory CD4 T cells between the two groups. Memory T cells are antigen-generated, generally long-lived, and quiescent cells that respond very rapidly and effectively to a subsequent challenge with the same antigen. Normal mice kidneys are known to contain more effector memory CD4 and CD8 T cells which express high CD44 and low CD62L than the spleen or blood (2). Our results imply that the number of kidney-infiltrating effector memory CD8 T cells is influenced by the previous microbial burden but that those cells are not involved in the pathogenesis of ischemic AKI because the kidneys of germ-free mice, with less effector memory CD8 T cells, had more severe injury after renal IRI.

At 24 h after renal IRI, more CD8 T cells trafficked into the postischemic kidneys of germ-free mice, suggesting the important role of CD8 T cells in the initial injury phase of renal IRI. CD4 T cells have been identified as a crucial mediator of renal IRI. CD4-deficient mice were significantly protected from renal IRI (5), and less CD4 T cell infiltration mediated by IL-16 deficiency also showed a protective effect (43). However, the role of CD8 T cells in renal IRI is not known. Our results suggest that naive CD8 T cells, having less chance of microbial stimuli, could infiltrate postischemic kidneys and play more roles in establishing renal injury following IRI. CD8 T cells isolated from postischemic kidneys were recently reported to produce more IFN- $\gamma$  compared with normal and sham-operated kidneys (2).

Normal kidneys of conventionalized germ-free mice revealed similar expression of IL-4, IL-10, and IFN-y compared with age-matched control mice. Thus expression of these cytokines in normal kidneys is influenced by microbial stimuli. There was also no difference in the number of NKT cells between conventionalized germ-free mice and age-matched control mice. Conventionalized germ-free mice showed equivalent renal functional injury to age-matched (11-wk-old) control, whereas germ-free mice showed significantly more renal functional injury than control mice. The degree of tubular injury with the percentage of necrotic tubules in the sets of 6W control vs. 11W control and 6W GF vs. 11W GF vs. 11W CV-GF mice was compared. Older control (11W control) mice showed more tubular injury compared with 6W control mice. There was more tubular injury in 11W GF mice compared with 6W GF mice. Conventionalized germ-free (11W CV-GF) mice showed less tubular injury in all three areas of the postischemic kidney compared with 11W GF mice and in the medulla of postischemic kidney compared with 6W GF mice, suggesting that exposure to "nonpathogenic" microbes has some protective effect on initial renal injury following renal IRI.

Commensal microbes are also reported to have a protective role in the pathogenesis of type I diabetes, a debilitating

Fig. 9. Introduction of microorganisms in germ-free mice attenuates tubular injury. A: tubular injury was scored with the percentage of necrotic tubules among total tubules. Old control (11-wk-old) mice showed significantly increased tubular injury in both the outer medulla and inner medulla compared with control (6-wk-old) mice. \*P < 0.05 compared with 6W control. B: 11W CV-GF mice showed decreased tubular injury compared with germ-free mice. \*P < 0.05 compared with 11W GF. C: both the outer medulla and inner medulla of old control (11-wk-old) mice exhibited higher proportions of necrotic tubules than control (6-wk-old) mice. D: postischemic kidneys of 11W CV-GF mice showed mitigated tubular injury compared with both 6W GF and 11W GF mice. Arrowheads indicate necrotic tubules, and arrows indicate tubular casts. Groups are defined as in Figs. 1, 3, and 6.

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#### GERM STATUS AND RENAL IRI

autoimmune disease caused by T cell-mediated destruction of the pancreas (44). In this study, germ-free, MyD88 (an adaptor for multiple innate immune receptors that recognize microbial stimuli)-negative NOD mice developed robust type I diabetes, while specific pathogen-free NOD mice lacking MyD88 did not. Conventionalization of germ-free, MyD88-negative NOD mice attenuated the development of diabetes.

Our study demonstrates, for the first time, the important role of microbial stimuli in kidney lymphocytes, cytokine expression, and the response to kidney IRI. Future studies can be geared to identify the details of the microbiota that modulate these effects in the kidney and alter microbiota therapeutically to improve outcomes from kidney IRI.

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