Sensitive biomarkers are needed to detect kidney injury at the earliest stages. The objective of this study was to determine whether the appearance of kidney injury molecule-1 (Kim-1) protein ectodomain in urine and kidney injury molecule-1/ hepatitis A viral cellular receptor-1 (Kim-1/Havcr1) gene expression in kidney tissue may be more predictive of renal injury after exposure to nephrotoxants when compared to traditionally used biomarkers. Male Sprague-Dawley rats were injected with a range of doses of gentamicin, mercury (Hg; HgCl₂), or chromium (Cr; K₂Cr₂O₇). The results showed that increases in urinary Kim-1 and kidney Kim-1/Havcr1 gene expression paralleled the degree of severity of renal histopathology and were detected at lower doses of nephrotoxants when compared to blood urea nitrogen (BUN), serum creatinine, and urinary N-acetyl-β-D-glucosaminidase (NAG). In a time course study, urinary Kim-1 was elevated within 24 h after exposure to gentamicin (100 mg/kg), Hg (0.25 mg/kg), or Cr (5 mg/kg) and remained elevated through 72 h. NAG responses were nephrotoxicant dependent with elevations occurring early (gentamicin), late (Cr), or no change (Hg). At 72 h, after treatment with any of the three nephrotoxants, there was increased Kim-1 immunoreactivity and necrosis involving ~50% of the proximal tubules; however, only urinary Kim-1 was significantly increased, while BUN, serum creatinine, and NAG were not different from controls. In rats treated with the hepatotoxicant galactosamine (1.1 mg/kg), serum alanine aminotransferase was increased, but no increase in urinary Kim-1 was observed. Urinary Kim-1 and kidney Kim-1/Havcr1 expression appear to be sensitive and tissue-specific biomarkers that will improve detection of early acute kidney injury following exposure to nephrotoxic chemicals and drugs.

Key Words: acute kidney injury; nephrotoxicity biomarkers; kidney injury molecule-1; mercury; chromium; gentamicin.
we will use Kim-1/Havcr1 to designate this gene.) Furthermore, the Kim-1 protein ectodomain is cleaved and released into urine as demonstrated in a number of nephrotoxic models in rats and in humans with AKI following exposure to chemical agents, ischemia, and protein overload (Han et al., 2002; Ichimura et al., 2004). The objectives of the present study were to: (1) determine if elevated levels of Kim-1 protein in urine and increases in Kim-1/Havcr1 gene expression in kidney represent more sensitive biomarkers of AKI compared to traditional, routinely used biomarkers of nephrotoxicity; (2) compare the temporal relationship between Kim-1 and NAG excretion in urine; (3) determine the correlation of the appearance of Kim-1 in urine and Kim-1/Havcr1 expression in kidney with changes in nephropathology and Kim-1 immunoreactivity of renal tissue; and (4) evaluate the specificity of the appearance of Kim-1 in urine for renal injury. The study was conducted using three well-established animal models of AKI: inorganic mercury (Hg)-, hexavalent chromium (Cr)-, and gentamicin-induced nephrotoxicity.

MATERIALS AND METHODS

Animal care and treatments. All procedures requiring the use of animals were conducted in accordance with the U.S. Public Health Service Guide for the Care and Use of Laboratory Animals in an Association for Assessment and Accreditation of Laboratory Animal Care International-accredited facility. Male Sprague-Dawley rats (Harlan, Indianapolis, IN), weighing 270-300 g (10-12 weeks old) were housed in standard plastic cages and maintained on a 12-h light/dark cycle at 22-24°C with access to food and water ad libitum. In the dose-response study of gentamycin, rats received a daily sc injection for 3 days of gentamycin sulfate (Sigma, St Louis, MO) (25, 50, 100, 150, 200, and 400 mg gentamicin/kg) in distilled water at an injection volume of 2 ml/kg. In the dose-response study of Hg and Cr, rats were anesthetized with isoflurane and injected with HgCl₂ (Sigma) (0.1, 0.25, 0.5, or 1 mg Hg/kg, iv) or K₂Cr₂O₇ (Sigma) (1, 2.5, 5, 10, and 20 mg Cr/kg, sc) in 0.9% saline at an injection volume of 2 ml/kg. Immediately following the single injection of Hg or Cr and 24 h after the third injection of gentamycin, rats were placed individually in plastic metabolism cages with free access to food and water collected for 24 h. For the gentamycin time course study, rats received a sc injection of 100 mg/kg gentamicin daily for 3 days. A 24-h pre-study urine sample was collected prior to the injections of gentamycin, and urine was collected from the same rats at intervals of 0-4, 4-8, 8-24, 24-48, and 48-72 h commencing immediately after the third sc injection. For the Hg and Cr time course studies, rats were given a single injection of 0.25 mg/kg Hg, iv, or 5 mg/kg Cr, sc. A 24-h pre-study urine sample was collected, and urine was collected from the same rats at intervals of 0-4, 4-8, 8-24, 24-48, and 48-72 h commencing immediately after the single injection. All control rats received either an iv or sc injection of 0.9% saline (Hg and Cr vehicle) or a daily sc injection of distilled water (gentamicin vehicle) for 3 days. Urines were centrifuged at 3000 × g for 10 min at 4°C, and supernatants were collected and stored in multiple aliquots at −80°C. At the end of each experiment, rats were euthanized by terminal exsanguination under isoflurane anesthesia. Blood was collected from the inferior vena cava in heparinized syringes for biochemical analyses. Kidneys were removed immediately and sections preserved for histopathologic and immunohistochemical studies and in RNA Later (Ambion, Austin, TX) to preserve RNA for determination of Kim-1/Havcr1 expression.

To determine that the increase in urinary Kim-1 is specific for renal injury, a separate group of rats received a single ip injection of the hepatotocicant galactosamine (Sigma) (1.1 g/kg) or saline vehicle. At 12 and 24 h after treatment, blood and urine were collected for measurement of alanine aminotransferase (ALT) (Vetscan serum chemistry analyzer; Abaxis, Union City, CA) and Kim-1 protein, respectively.

Biochemical analyses for renal injury. Heparinized blood was analyzed for BUN and creatinine using a Reflotron Clinical Chemistry Analyzer (Boehringer Mannheim-Roche Diagnostics, Indianapolis, IN). Urine N-acetyl-β-D-glucosaminidase (NAG) was measured by NAG assay kit (Bio-quant, San Diego, CA). Urine creatinine was determined using QuantiChrom creatinine assay kit (BioAssay Systems, Hayward, CA). Urine Kim-1 protein was measured using Microsphere-based Lumxin xMAPTM technology with monoclonal antibodies raised against rat Kim-1 in the V.S.V.J.V.B laboratory. This technique is an adaptation of the recently developed and validated sandwich enzyme-linked immunosorbent assay (ELISA) described in Vaidya et al. (2005), J. Am. Soc. Nephrol. 16, 192A, [abstract]. This assay has been extensively evaluated with respect to its assay range (4-40,000 pg/ml), sensitivity (lowest limit of detection: 8.8 pg/ml), specificity, recovery analysis (90-100% recovery), precision profile (inter- and intra-assay variability <10%), and dilution linearity. The validation studies showed a significant correlation (92%) between the ELISA and microbead-based assay. This assay has been used to determine urinary levels of Kim-1 in several recent studies using rat models (de Borst et al., 2007; Perez-Rojas et al., 2007; van Timmeren et al., 2006).

RNA isolation and Kim-1/Havcr1 mRNA quantitation. Cross-sections of kidney were placed in RNA Later and stored at 4°C for 24 h to allow RNA Later to penetrate tissues. RNA Later was then removed, and the samples were stored at −80°C. RNA was isolated using RNeasy kits (Qiagen, Valencia, CA). Frozen kidney sections were homogenized using a VirTis Tempest rotor-stator homogenizer (VirTis, Gardiner, NY) in Qiagen RNA lysis buffer and processed according to RNeasy kit protocols. Qualitative analyses were performed on each sample using the RNA 6000 Nano Analyser on an Agilent 2100 bioanalyzer (Agilent Technologies, Palo Alto, CA). All samples had an RNA integrity number of 8 or greater. RNA samples were quantitated on a NanoDrop ND-1000 spectrophotometer (NanoDrop Technologies, Wilmington, DE), aliquoted, and frozen at −80°C.

Real-time quantitative PCR was used for relative quantitation of Kim-1/Havcr1 mRNA levels. cDNA was generated from total RNA using random hexamer primers with TaqMan Reverse Transcription Reagents kits (Applied Biosystems, Foster City, CA). PCR was performed using SYBR Green PCR Master Mix reagents (Applied Biosystems) with the ABI Prism 7900HT sequence detection system as described in User Bulletin #2 (updated 10/2001) using four replicate wells per sample. Relative amounts of target were calculated from standard curves for Kim-1/Havcr1 and the endogenous reference (ribosomal protein L7) prepared from five fourfold serial dilutions of a positive control sample (cDNA prepared from kidney RNA pooled from three Sprague-Dawley rats treated with 400 mg/kg gentamicin for 3 days). Gene expression data were normalized by dividing the amount of target mRNA by the endogenous reference. Relative fold changes in gene expression were calculated for individual animals by dividing the amount of normalized target mRNA by the mean of the three control animals for any given dose.

Primers sequences used in reverse transcriptase (RT)-PCR—Kim-1/Havcr1 forward primer: 5’-GTGAGTGGAGCCAGCAACAC-3’ and Kim-1/Havcr1 reverse primer: 5’-AATCCCTTGATCCATTGTTTTCTT-3’.

Histopathology and immunohistochemistry. Kidneys were fixed in 10% zinc-formalin (Anatech Ltd, Battle Creek, MI) for 48 h and then stored in 70% ethanol at room temperature prior to processing. Kidney sections were mounted on glass slides and stained with hematoxylin and eosin for histopathologic evaluation. Nephropathy scores were assigned semiquantitatively by light microscopy on a scale of 0-5: 0 = normal histology; 1 = degeneration only without necrosis; and 2-5 = < 25, 50, < 75, and > 75%, respectively, of epithelial cells of the proximal convoluted tubules showing necrosis, degeneration, regeneration, tubular dilatation, protein casts, and interstitial lymphocytic infiltration. For detection of immunoreactivity of renal tissues, formalin-fixed, paraffin-embedded renal tissue sections were mounted to glass

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slides coated with poly-L-Lysine (American HistoLabs, Inc., Gaithersburg, MD). Preparation of these slides included microwave irradiation of the sections in a pressure cooker with antigen retrieval Citra solution (BioGenex, San Ramon, CA). For blocking endogenous peroxidase activity, sections were incubated with 0.3% hydrogen peroxidase in methanol for 30 min and then with 5% normal horse serum for 30 min. Sections were incubated overnight at 4°C with mouse monoclonal antibody against Kim-1 (Brigham and Women’s Hospital and Harvard Medical School, Boston, MA) at a dilution of 1:50, followed by incubation with a biotinylated secondary antibody (Vector, Burlingame, CA) for 1 h, and then with avidin-biotinylated horseradish peroxidase complex (Vector) for 30 min. The peroxidase reaction was carried out with 0.05% 3,3′-diaminobenzidine in 0.1M Tris-HCl buffer and 0.01% hydrogen peroxide for 10 min. The sections were counterstained with hematoxylin for 5 min. For negative staining, the primary antibody was omitted from the incubation step.

Statistical analyses. Data for biomarker assays are expressed as fold changes to make appropriate comparisons between biomarkers. For dose-response studies, fold-change data are expressed as box-and-whisker plots and show median, 25th and 75th percentiles, and the smallest and largest values in the range of data, to help illustrate and define the data distribution. For the time course studies using same rats to collect urine at sequential intervals, fold-change data are expressed as mean ± SD. A one-way ANOVA followed by a Bonferroni post hoc test was employed to compare differences between the treatment groups and control group. A Kruskal-Wallis rank test was used to assess the differences in the nephropathology scores. All analyses were conducted using GraphPad Prism version 4.03 for Windows (GraphPad Software, San Diego, CA). A two-tailed p value < 0.05 was considered statistically significant.

RESULTS
Renal Pathology and Immunohistochemistry

Representative photomicrographs of lesions in rat kidney 72 h after administration of nephrotoxic agents are presented in Figure 1. No lesions were found in the kidney from control rats (Fig. 1A). Gentamicin-, Hg-, or Cr-induced renal lesions were basically similar and consisted mainly of epithelial cell necrosis (Figs. 1C, E, and G), degeneration, and regeneration of the proximal convoluted tubules. Dilatation of the tubules, protein casts in the tubular lumina, glomerular vacuolization, and interstitial lymphocytic infiltration were also observed. In the time course study of Cr, two of six rats had severe tubular necrosis and regeneration in addition to renal cell carcinoma (Fig. 1I) and adenofibroma. These tumors were unlikely to be age related or related to treatment due to the short exposure time. Immunoperoxidase staining for Kim-1 was not evident in control kidneys (Fig. 1B). In contrast, positive immunoperoxidase staining for Kim-1 was observed in tubular epithelial cells of the kidneys from gentamicin-, Hg-, or Cr-treated rats (Fig. 1D, F, and H), including in the renal tumor-positive rats (Fig. 1J).

Dose-Response Studies

In the gentamicin dose-response study, BUN, serum creatinine, and urinary NAG levels in rats were significantly elevated above control values only at the highest dose of 400 mg/kg, a dose sufficient to produce ARF (Fig. 2). The mean relative fold changes were 5, 6, and 43 for BUN, blood creatinine, and urinary NAG, respectively, which were statistically significant. In contrast, both urinary Kim-1 protein and kidney Kim-1/Havcr1 mRNA expression levels increased in a dose-related manner after injection of gentamicin at doses
The mean increases of urinary Kim-1 were 11-, 30-, 82-, and 260-fold higher than control urinary Kim-1 (165.9 ± 141 pg/mg creatinine) and the mean increases of kidney Kim-1/Havcr1 mRNA levels were 41-, 112-, 240-, and 514-fold higher than control for gentamicin doses of 100, 150, 200, and 400 mg/kg, respectively. Moreover, the increase of urine Kim-1 and kidney Kim-1/Havcr1 expression paralleled dose-dependent changes in nephropathy score (Fig. 2).

BUN, blood creatinine, and urinary NAG levels significantly increased during 24 h posttreatment only at doses of Hg ≥ 0.5 mg/kg (Fig. 3). At Hg doses of 0.5 and 1 mg/kg, BUN, creatinine, and urinary NAG increased 3- and 6-fold, 4- and 7-fold, 8- and 17-fold, respectively, compared to control. In contrast, urinary Kim-1 protein values were 14- and 25-fold higher than control values (165.9 ± 141 pg/mg creatinine) at Hg doses of 0.25 and 0.5 mg/kg, respectively; these increases paralleled changes in
nephropathy scores. Unexpectedly, urinary Kim-1 only increased fivefold after injection of 1 mg/kg Hg, which is not statistically a significant, compared to control, likely due to severe renal damage. From our previous experience, this dose of Hg leads to significant mortality rate after 48–72 h, which may account for this result. Kidney Kim-1/Havcr1 mRNA expression increased 71-, 187-, and 167-fold at Hg doses of 0.25, 0.5, and 1 mg/kg, respectively, compared to control. Elevations in Kim-1/Havcr1 gene expression paralleled dose-dependent changes in nephropathy scores.

In the Cr dose-response study, BUN and blood creatinine levels both significantly increased by 1.5-fold after injection of 10 mg/kg (Fig. 4). At 20 mg/kg Cr, BUN and creatinine increased six- and threefold above control, respectively. A statistically significant elevation in urine NAG was observed at Cr doses as low as 5 mg/kg and higher; NAG levels were three- , five- and eightfold higher than control at 5, 10, and 20 mg Cr/ kg, respectively. Urinary Kim-1 values were twofold higher than control (165.9 ± 141 pg/mg creatinine) at Cr doses of 5 mg/kg; three- and fivefold increases were observed at doses of

FIG. 3. Comparison of urinary Kim-1 and kidney Kim-1/Havcr1 expression with routinely used biomarkers BUN, creatinine, and NAG and nephropathy score after treatment with various doses of Hg (HgCl2). Immediately following the single injection of Hg as described in the Materials and Methods, rats were placed individually in plastic metabolism cages and urine collected for 24 h, after which rats were euthanized and blood and kidneys were collected. Urinary NAG and Kim-1 are normalized by urinary creatinine. Relative fold changes of BUN, serum creatinine, urinary NAG, urinary Kim-1, and kidney Kim-1/Havcr1 expression for each animal were calculated by dividing the value for each individual animal level by the level averaged from all corresponding vehicle (saline or distilled water)-treated animals and are displayed in box and whisker plots. BUN (control) = 16.6 ± 3.3 mg/dL; creatinine (control) = 0.5 ± 0.0 mg/dL; NAG (control) = 20.3 ± 10.6 (U/g creatinine); urinary Kim-1 (control) = 165.9 ± 141 (pg/mg creatinine); kidney Kim-1/Havcr1 = 1.2 ± 0.7. Asterisks indicate statistically significant difference (p < 0.05) compared to control group (n = 6 for each group).
10 and 20 mg/kg, respectively, which were statistically significant. Kidney Kim-1/Havcr1 gene expression increased 5-, 14-, and 37-fold at doses of 5, 10, and 20 mg/kg Cr, respectively. The increases in urinary NAG, Kim-1, and kidney Kim-1 are normalized by urinary creatinine. Relative fold changes of BUN, creatinine, NAG, urinary Kim-1, and kidney Kim-1 expression for each animal were calculated by dividing the value for each individual animal level by the level averaged from all corresponding vehicle (saline or distilled water)-treated animals and are displayed in box and whisker plots. BUN (control) = 16.6 ± 3.3 mg/dL; serum creatinine (control) = 0.5 ± 0.0 mg/dL; NAG (control) = 20.3 ± 10.6 (U/g creatinine); urinary Kim-1 (control) = 165.9 ± 141 (pg/mg creatinine); kidney Kim-1/Haverl = 1.2 ± 0.7. Asterisks indicate statistically significant difference ($p < 0.05$) compared to control group ($n = 6$ for each group).

**Time Course Studies**

A statistically significant, threefold increase in urinary NAG was observed during 0–4 and 4–8 h after the third sc injection of 100 mg/kg gentamicin (Fig. 5). In subsequent collection intervals encompassing 8–72 h, NAG values were lower and not significantly different from controls. In contrast, urine Kim-1 increased 7-, 12-, 26-, 78-, and 73-fold for the 0- to 4, 4- to 8, 8- to 24, 24- to 48, and 48- to 72 h intervals, respectively, above controls (59.1 ± 17.6 pg/mg creatinine), and the increases were statistically significant compared to controls for intervals between 24–72 h (Fig. 5).

After treatment with 0.25 mg Hg/kg, no significant increases in urinary NAG were observed relative to controls for all intervals from 0 to 72 h (Fig. 6). In contrast, urinary Kim-1
increased 30-, 140-, 114-fold at the 8- to 24, 24- to 48, and 48- to 72 h intervals, respectively, above controls (59.1 ± 17.6 pg/mg creatinine) and the increases were statistically significant compared to saline controls between 24 and 72 h.

After treatment with 5 mg Cr/kg, urinary NAG increased twofold higher than controls for the 8- to 24, 24- to 48, and 48- to 72 h intervals, but the increases were statistically significant only at the earlier two intervals (Fig. 7). Urinary Kim-1 concentrations increased 6-, 58- and 212-fold at the 8- to 24, 24- to 48, and 48- to 72 h intervals, respectively, above controls (59.1 ± 17.6 pg/mg creatinine) and were statistically different from controls from 24 to 72 h.

At 72 h after treatment with any of the three nephrotoxins, rats were euthanized and blood collected for BUN and blood creatinine analyses and kidneys preserved for histopathological and immunohistochemical evaluation. These data were compared with urinary biomarkers for the last collection interval (48–72 h; Fig. 8). Renal histopathology scores indicated a significant degree of renal injury compared to controls after treatment with gentamicin (100 mg/kg), Hg (0.25 mg/kg), or Cr (5 mg/kg). Urinary Kim-1 levels significantly increased after exposure to each nephrotoxicant and were statistically significant, in parallel with the nephropathy scores. Other
biomarkers—BUN, creatinine, and urinary NAG—were not different from controls 72 h after treatments.

**Galactosamine Hepatotoxicity and Urinary Kim-1**

At 12 and 24 h after treatment with galactosamine, serum ALT activity was markedly elevated, suggestive of frank liver injury (Table 1). In contrast, no increase in urinary Kim-1 was detected with galactosamine-induced hepatotoxicity compared to saline controls, indicating that Kim-1 excretion in urine is a specific marker of renal injury and not hepatic injury.

**DISCUSSION**

The development of sensitive, specific, and early biomarkers of AKI will enhance the detection of the nephrotoxic potential of drug candidates and compounds released unintentionally from medical devices during preclinical studies. Better nephrotoxicity biomarkers will permit earlier diagnosis of AKI and allow medical personnel to intervene sooner to preserve residual renal function in the clinical or hospital setting and improve health outcomes. In the present study, our goal was to determine if Kim-1, a new nephrotoxicity biomarker, is a specific marker of renal injury and not hepatic injury.

**Table 1**

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Serum ALT (U/l)</th>
<th>Urinary Kim-1 (pg/ml)</th>
</tr>
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<tbody>
<tr>
<td>Control</td>
<td>45 ± 6</td>
<td>102.8 ± 49.2</td>
</tr>
<tr>
<td>Galactosamine, 12 h</td>
<td>557 ± 355*</td>
<td>73.6 ± 13.2</td>
</tr>
<tr>
<td>Galactosamine, 24 h</td>
<td>2377 ± 1851*</td>
<td>90.5 ± 34.9</td>
</tr>
</tbody>
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*p < 0.05 compared to control group (n = 5 for each group).
Kim-1, a type 1 transmembrane protein that contains immunoglobulin (Ig) and highly O- and N-glycosylated mucin domains in its ectodomain and a relatively short cytoplasmic tail (Ichimura et al., 1998). The upregulation of Kim-1/Havcr1 mRNA expression with renal pathology was investigated. In our experience, a high rate of kidney injury after exposure to elemental or inorganic forms of Hg; convoluted portions (S1 and S2 segments) of the proximal tubule can also be involved when the toxic effect is more severe (Diamond and Zalups, 1998). Hg has a strong affinity for thiol groups on molecules such as glutathione, cysteine, metallothionein, and other proteins in the body (Johnson, 1982; Zalups, 2000). Binding to key enzymes and structural or functional proteins has been proposed to cause the Hg toxicity (Boadi et al., 1991). The effects of Hg on intracellular thiol metabolism, oxidative stress, lipid peroxidation, mitochondrial function, and intracellular distribution of calcium ions may also play a role in nephropathy induced by Hg (Mahboob et al., 2001; Zalups, 2000).

Therefore, Kim-1 has been suggested to serve as a useful general biomarker for renal proximal tubule injury in preclinical and clinical studies of drug safety evaluation, chemical-related renal injury, and the monitoring of renal disease states. In the current study, three well-established AKI models—gentamicin, a clinically relevant aminoglycoside antibiotic, and Hg and Cr, two nephrotoxic metals—were used to compare the sensitivity of Kim-1 to BUN and serum creatinine, two routinely used biomarkers in preclinical and clinical regimens, and urinary NAG, a lysosomal enzyme found predominantly in proximal tubules. Furthermore, the relationship between urinary Kim-1 excretion and kidney Kim-1/Havcr1 gene expression with renal pathology was investigated.

In the dose-response study, urinary Kim-1 and kidney Kim-1/Havcr1 mRNA expression are very sensitive to the renal injury produced by gentamicin, a potent broad-spectrum aminoglycoside that is commonly used in the treatment of severe Gram-negative bacterial infections (Cunha and Schor, 2002). Gentamicin accumulates in the epithelial cells of renal proximal tubules, leading to structural changes and functional impairments of the plasma membrane, mitochondria, and lysosome (Ali, 1995; Olbricht et al., 1991). Nephrotoxicity is a dose-limiting side effect of gentamicin. In the present study, histopathology observations showed that a single dose of 150 mg/kg gentamicin for 3 days produced a modest degree of renal tubule injury (necrosis in ≤ 25% of the tubules). At this low degree of injury, urinary Kim-1 protein increased 30-fold and kidney Kim-1/Havcr1 expression increased 112-fold, while BUN, serum creatinine, and urinary NAG did not change after treatment with this dose. In fact, these routinely used nephrotoxicity biomarkers were not significantly elevated except at the highest dose of 400 mg/kg, a dose used to produce AKI and tubular necrosis in laboratory animal models.

The results of the current study indicated that urinary Kim-1 and kidney Kim-1/Havcr1 expression were more sensitive biomarkers for nephrotoxicity induced by Hg. Kidney proximal tubules, in particular the pars recta or S1 segments, are the primary targets where mercuric ions accumulate and express toxicity after exposure to elemental or inorganic forms of Hg; convoluted portions (S1 and S2 segments) of the proximal tubule can also be involved when the toxic effect is more severe (Diamond and Zalups, 1998). Hg has a strong affinity for thiol groups on molecules such as glutathione, cysteine, metallothionein, and other proteins in the body (Johnson, 1982; Zalups, 2000). Binding to key enzymes and structural or functional proteins has been proposed to cause the Hg toxicity (Boadi et al., 1991). The effects of Hg on intracellular thiol metabolism, oxidative stress, lipid peroxidation, mitochondrial function, and intracellular distribution of calcium ions may also play a role in nephropathy induced by Hg (Mahboob et al., 2001; Zalups, 2000). In the current study, a dose of 0.25 mg/kg Hg induced renal injury involving 25–50% of the proximal tubules while a dose of 0.5 mg/kg Hg induced histopathologic lesions involving 50–75% of tubules. Increases of urinary Kim-1 (14-fold) and kidney Kim-1/Havcr1 mRNA (71-fold) were detected following a single injection of 0.25 mg/kg Hg. Increases in BUN, creatinine, and urinary NAG were evident only at doses of 0.5 mg Hg/kg or higher, and the fold-increase was much smaller than that of Kim-1. An unusual response was observed at the highest dose of Hg, 1 mg/kg, where mean urinary Kim-1 elevated fivefold, but this level was not significantly different from control levels while kidney Kim-1/Havcr1 expression increased by 167-fold. A possible explanation for the markedly reduced appearance of Kim-1 in urine may be related to the severe renal injury produced at the highest dose of 1 mg/kg which resulted in a marked decrease in Kim-1 protein synthesis. In our experience, a high rate of mortality is observed in rats treated with 1 mg/kg after 3–4 days of exposure. Other possible mechanisms may explain this unexpected response. First, the urinary excretion of Hg increases due to the administration of high-dose Hg in part due to the release of Hg from necrotic proximal tubular epithelial cells (Madsen and Hansen, 1980). The Ig domain of Kim-1 contains six cysteine residues. The high bonding affinity between Hg and sulfhydryl groups in cysteine might change the structure of Kim-1 Ig domain. The sandwich Kim-1 ELISA assay employed in this study uses monoclonal antibodies made against the purified ectodomain of Kim-1. Therefore, the high concentration of urinary Hg may interfere with the Kim-1 assay while not affecting kidney Kim-1/Havcr1 mRNA expression measured by RT-PCR via a similar mechanism. Alternatively, the cleavage of Kim-1 ectodomain into urine has been suggested to be mediated by a matrix metalloproteinase (Bailly et al., 2002). A high dose of Hg has been reported to markedly inhibit the activities of several enzymes, e.g., glutathione disulfide reductase and glutathione peroxidase in proximal tubule epithelial cells (Addya et al., 1984; Chung et al., 1982). The metalloproteinase might be inhibited after a high dose of Hg but not at lower doses of Hg. Another possibility is that the structure of the proteolytic site of Kim-1 might be altered by Hg which blocked the cleavage and shedding of the ectodomain, resulting in a reduced Kim-1 excretion into urine.
Acute tubular necrosis induced by Cr salts has been used as a model to study AKI (Biber et al., 1968; Wedeen and Qian, 1991). Convoluted proximal tubules (S1 and S2 segments) constitute the site where Cr selectively accumulates and produces necrosis (Wedeen and Qian, 1991). Oxidative and nitrosative stress-related injury has been associated with Cr-induced nephrotoxicity (Barrera et al., 2003; Pedraza-Chaverri et al., 2005). In the current study, results from the Cr dose-response suggested that urinary NAG is a good biomarker for Cr nephrotoxicity. An increase of NAG by threefold at 5 mg/kg Cr was observed which was statistically significant from control. At the same dose, a trend for increases in urinary Kim-1 (twofold) and kidney Kim-1/Havcr1 expression (fivefold) were observed; however, statistically significant increases in these two markers were not evident except at higher doses ≥ 10 mg/kg Cr, an outcome related perhaps to the high variability in the urinary Kim-1 data in the control group for the Cr experiments.

A time-dependent increase in nephrotoxicity was observed for each of the three nephrotoxic agents for intervals from 0 to 72 h posttreatment. With gentamicin, a progressive increase in urinary Kim-1 levels was observed as early as 0–4 h and levels continuously increased to 72-fold compared to controls at 24–48 h and 48–72 h. An unexpected early and transient response for urinary NAG was observed after gentamicin treatment. Following treatment with either Hg or Cr, the appearance of Kim-1 in urine was evident as early as 8–24 h after treatment with Hg (30-fold increase) and Cr (fivefold increase). Marked increases in urinary Kim-1 of magnitudes ranging from 50- to 200-fold were observed from 24–48 h and 48–72 h after treatment with Hg or Cr. While the histopathology indicated ~50–75% injured tubules at 72 h after nephrotoxicant treatment (Fig. 8), only Kim-1 increased dramatically while some nonresponses (e.g., Hg) or transient, mild responses after some treatments (gentamicin, Cr) were observed with NAG. Thus, the marked increases of Kim-1 at these same time points were no different from controls. This experiment was critical because a splice variant of Kim-1/Havcr1, a gene isolated from monkey kidney cells that codes for a receptor for the hepatitis A virus, is expressed in liver (Kaplan et al., 1996) and several human organs and tissues (Feigelstock et al., 1998). In studies interrogating the liver genome after treatment with hepatotoxic agents, no increase in HAVcr-1 gene expression has been reported (Jessen et al., 2003). Results from the current study and other genomic studies suggest that the appearance of Kim-1 in urine is tightly linked to the renal injury and not liver injury.

The value of a biomarker also relies on whether the method of assay is simple, cheap, and fast. A sensitive high-throughput method has been developed to quantify urinary Kim-1 by a sandwich ELISA (Vaidya et al., 2006) and now a Luminex-based bead assay (Liangos et al., 2007). Another methodological advantage of Kim-1 is that it is stable in urine and no stabilizing buffer needed for the collected urine samples prior to freezing and storage for later analysis.

In conclusion, a consistent pattern of marked elevations in urinary Kim-1 concentrations and kidney Kim-1/Havcr1 expression was observed using three different agents with known nephrotoxic potential. Under the experimental conditions of the current studies, these two biomarkers were elevated earlier during renal tubular pathogenesis than the currently accepted, routinely used nephrotoxicity biomarkers and, hence, appear to be more sensitive nephrotoxicity biomarkers. The increases in urinary Kim-1 protein and kidney Kim-1/Havcr1 mRNA levels in response to the three nephrotoxics were also more closely anchored to the increases in severity in renal histopathology than BUN and serum creatinine. As a noninvasive indicator of renal proximal tube injury, urinary Kim-1 was elevated at early stages of the
injury, and the elevations were sustained throughout the progression of cell injury to proliferation and regeneration. Positive immunoreactivity of Kim-1 in kidneys from rats treated with nephrotoxins confirmed that this protein is synthesized in response to renal injury. Additional studies are needed to establish the kinetics and potential recovery/reversibility of these responses under chronic exposure conditions to wider spectrum of nephrotoxics and control compounds. Therefore, the data presented in the report suggest that urinary Kim-1 and kidney Kim-1/Havcr1 gene expression may serve as potentially valuable nephrotoxicity biomarkers. These biomarkers could improve the identification of nephrotoxic compounds in preclinical evaluations of compounds, for monitoring the progression of renal pathogenesis in clinical evaluations of drugs and exposures to residues and contaminants associated with medical devices, and for exposures to toxicants in occupational and environmental exposures.

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