Gingerol Fraction from *Zingiber officinale* Protects against Gentamicin-Induced Nephrotoxicity


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Nephrotoxicity is the main complication of gentamicin (GM) treatment. GM induces renal damage by overproduction of reactive oxygen species and inflammation in proximal tubular cells. Phenolic compounds from ginger, called gingerols, have been demonstrated to have antioxidant and anti-inflammatory effects. We investigated if oral treatment with an enriched solution of gingerols (GF) would promote a nephroprotective effect in an animal nephropathy model. The following six groups of male Wistar rats were studied: (i) control group (CT group); (ii) gingerol solution control group (GF group); (iii) gentamicin treatment group (GM group), receiving 100 mg/kg of body weight intraperitoneally (i.p.); and (iv to vi) gentamicin groups also receiving GF, at doses of 6.25, 12.5, and 25 mg/kg, respectively (GM + GF groups). Animals from the GM group had a significant decrease in creatinine clearance and higher levels of urinary protein excretion. This was associated with markers of oxidative stress and nitric oxide production. Also, there were increases in the mRNA levels for proinflammatory cytokines (tumor necrosis factor alpha [TNF-α], interleukin-1β [IL-1β], IL-2, and gamma interferon [IFN-γ]). Histopathological findings of tubular degeneration and inflammatory cell infiltration reinforced GM-induced nephrotoxicity. All these alterations were attenuated by previous oral treatment with GF. Animals from the GM + GF groups showed amelioration in renal function parameters and reduced lipid peroxidation and nitrosative stress, in addition to an increment in the levels of glutathione (GSH) and superoxide dismutase (SOD) activity. Gingerols also promoted significant reductions in mRNA transcription for TNF-α, IL-2, and IFN-γ. These effects were dose dependent. These results demonstrate that GF promotes a nephroprotective effect on GM-mediated nephropathy by oxidative stress, inflammatory processes, and renal dysfunction.

Gentamicin (GM) is a typical aminoglycoside antibiotic agent that is widely used in clinical practice for the treatment of Gram-negative infections (1, 2). However, its use is complicated by a great risk of nephrotoxicity, which affects 10 to 30% of patients, especially after long-term use (3–5).

In its elimination route, GM accumulates in the renal proximal tubular cells through the megalin/cubilin complex receptor, which is responsible for GM transportation inside the cell. Aminoglycoside-induced nephrotoxicity is characterized by tubular cell apoptosis and/or necrosis, predominantly in the proximal tubules (3, 6). Clinically, it results in acute kidney injury (AKI) with elevations of serum creatinine (SCr) and urea, in addition to proteinuria (5, 7). Pathological findings can be seen as proximal tubular edema, tubular necrosis, and desquamation, as well as inflammation and diffuse interstitial edema.

Several studies have shown the involvement of reactive oxygen species (ROS) in gentamicin-induced AKI. Gentamicin has been shown to enhance the generation of superoxide anion (O$_2^-$), peroxynitrite anion (ONOO$^-$), hydrogen peroxide (H$_2$O$_2$), and hydroxyl radical (OH$^-$) production from renal cortical mitochondria, which is associated with an increase in lipid peroxidation and decrease in antioxidant enzymes (8, 9). These ROS act directly on cells, affecting their structure and functionality (2, 5).

Several phytotherapeutic agents have been used to prevent or ameliorate GM-induced AKI (5, 10–12), including ginger extract. Ginger (*Zingiber officinale*) is one of the world’s best-known spices and is cultivated in several countries. It has been used since antiquity for its health benefits (13). Extracts obtained from its roots usually contain polyphenol compounds, such as [6]-gingerol, [8]-gingerol, and [10]-gingerol, which have been cited as the main components responsible for its pharmacological effects (14, 15). Among other potential mechanisms, ginger has antioxidant (15–17) and anti-inflammatory properties (15, 18, 19).

Ginger has been shown to protect against renal ischemia-reperfusion injury and cisplatin-related nephrotoxicity (20, 21). In the former case, its use has been associated with an antiapoptotic effect. However, ginger’s anti-inflammatory effects on acute nephrotoxicity have scarcely been studied. In the present study, we aimed to evaluate the effects of a gingerol-enriched fraction of ginger extract in a model of gentamicin-induced nephrotoxicity, focusing mainly on gentamicin’s oxidative and inflammatory effects (15, 18).

**MATERIALS AND METHODS**

**Animals and experimental drug.** Experiments were performed with male Wistar rats weighing 240 to 280 g. The animals were housed under standard laboratory conditions and maintained on a 12-h light-dark cycle, and they had free access to food (Biotec, Agua Fria, Santa Catarina, Brazil) and water. The experimental protocols and all procedures were conducted according to the norms of the National Council for Control of Animal...
Experimentation (CONCEA) and were approved by the Ethics Committee of Animal Research of the Federal University of Ceará, Fortaleza, Brazil (protocol 68/12).

GM was purchased from Sigma and was administered intraperitoneally (i.p.) for 7 days. The GM solution was prepared so that the maximum injected volume was 0.5 to 0.6 ml/rat.

GF. Nine grams of ginger extract obtained from fresh ginger rhizomes was fractionated into 14 fractions (F1 to F14) by column chromatography (18.0 cm by 4.2-cm internal diameter (i.d.)) on silica gel (70-230 mesh) with gradients of n-hexane and ethyl acetate. After high-pressure liquid chromatography (HPLC) analysis, the gingerolides were identified in F3 (gingerol fraction [GF]). The standard gingerolides used in the analysis had already been isolated and identified in previous studies (22). The gingerolides present in the GF were quantified on a semipreparative HPLC system through a sample injection (50 mg of GF), collection of the fractions that contained each gingerol, and further weighing (22). The GF contained 41.7% [6]-gingerol (the main ginger compound), 4.8% [8]-gingerol, 2.4% [10]-gingerol, and 51.1% other minor compounds. GF was diluted in a 2% Tween 80 solution and orally administered in a maximum volume of 2.5 to 3.5 ml per rat.

Experimental design. Animals were assigned to 6 different groups containing 7 or 8 animals each. The sham group (CT group) received 0.4-mI.p. injections of saline solution (0.9%) for 7 consecutive days and oral treatment with 2% Tween 80 solution in the last 5 days. The second group (GF group) also received 0.4-mI.p. injections of saline for 7 consecutive days, plus 5 days of gingerol-enriched solution (25 mg/kg of body weight) through the oral route. Another group (GM group) received i.p. injections of GM (100 mg/kg) for 7 consecutive days (23) and oral treatment with 2% Tween 80 solution for 5 days. Finally, three other groups (GM + GF groups) received i.p. injections of GM (100 mg/kg) for 7 consecutive days and three different doses of GF (6.25, 12.5, and 25 mg/kg) for 5 days. The oral administration of 2% Tween 80 or GF was always given on the fifth day after the first GM or saline injection.

Evaluation of renal functions. The rats were kept individually in metabolic cages, and urine was collected for a 24-h period after the last oral administration. The animals were anesthetized, and a blood sample was obtained from the abdominal aorta. Blood plasma was separated for biochemical measurements. The left kidney was removed and immediately stored at −80°C. The right kidney was stored in 10% formalin for the histological studies. Plasma and urine samples were used for the analysis of urea, uric acid, and urinary protein by use of standard diagnostic kits (Labtest, Fortaleza, Brazil). Creatinine was also measured spectrophotometrically. The left kidney was removed and immediately homogenized in an ice-cold 0.02 M EDTA solution. Aliquots (400 µl) of tissue homogenate were mixed with 320 µl of distilled water and 80 µl of 50% (wt/vol) trichloroacetic acid in glass tubes and centrifuged at 3,000 × g for 15 min. Supernatants (400 µl) were mixed with 800 µl Tris buffer (0.4 M; pH 8.9), and 20 µl of 5,5-dithiobis(2-nitrobenzoic acid) (DTNB; 0.01 M) was added. After shaking of the reaction mixture, the absorbance was measured at 412 nm within 5 min of DTNB addition, against a blank with no homogenate. The absorbance values were extrapolated from a glutathione standard curve, and the level of GSH was expressed in grams of GSH per milligram of protein.

SOD activity. Superoxide dismutase (SOD) activity was measured according to the method of Sun et al. (27). The activity of this enzyme was evaluated by measuring its capacity to inhibit the photochemical reduction of nitroblue tetrazolium (NBT). In this assay, the photochemical reduction of riboflavin generates O2, which reduces NBT to produce formazan salt, which has maximal absorbance at 560 nm. In the presence of SOD, the reduction of NBT is inhibited, as the enzyme converts the superoxide radical to peroxide. The results are expressed as amounts of SOD needed to inhibit the rate of NBT reduction by 50%, in units of enzyme per gram of protein. Homogenates (10% tissue in phosphate buffer) were centrifuged (1 min, 2,608 × g, 4°C), and the supernatant was removed and centrifuged a second time (20 min, 15,294 × g, 4°C). The supernatant was then assayed. In a dark chamber, 1 ml of the reactant (50 mM phosphate buffer, 100 mM EDTA, and 13 mM L-methionine, pH 7.8) was mixed with 30 µl of the sample, 150 µl of NBT (75 µM), and 300 µl of riboflavin (2 µM). The tubes containing the resulting solution were exposed to fluorescent light bulbs (15 W) for 15 min and then read using a spectrophotometer at 560 nm.

TABLE 1 Oligonucleotide sequences of primers used for qPCR

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<th>Gene</th>
<th>Primer direction</th>
<th>Primer sequence (5′-3′)</th>
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<tr>
<td>TNF-α</td>
<td>Sense Antisense</td>
<td>GTACCACCTCTGAGGAAAAAC AGTTGGTGTGATCTGACAGTG</td>
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<tr>
<td>IL-1β</td>
<td>Sense Antisense</td>
<td>GACCTGTTCTTGGAGGCTGACA CTGATCGGAGCCGCAAGTC</td>
</tr>
<tr>
<td>IL-2</td>
<td>Sense Antisense</td>
<td>CAAAGGAGGCACAGATTGCA CGGATGGATCTTCCAGGCA</td>
</tr>
<tr>
<td>IFN-γ</td>
<td>Sense Antisense</td>
<td>ACAGTAAGGAAAAAGGATTGCA GCTGGATCTGTGGTGTTC</td>
</tr>
<tr>
<td>YWHAZ</td>
<td>Sense Antisense</td>
<td>GCTACTTGGCTAGGTTTGC TGTCGTAGCCTGGTTCACAAAT</td>
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Reduced glutathione (GSH) content in renal tissue was estimated according to the method described by Sedlak and Lindsay (26), with a few modifications. For the GSH assay, each kidney was homogenized in an ice-cold 0.02 M EDTA solution. Aliquots (400 µl) of tissue homogenate were mixed with 320 µl of distilled water and 80 µl of 50% (wt/vol) trichloroacetic acid in glass tubes and centrifuged at 3,000 × g for 15 min. Supernatants (400 µl) were mixed with 800 µl Tris buffer (0.4 M; pH 8.9), and 20 µl of 5,5-dithiobis(2-nitrobenzoic acid) (DTNB; 0.01 M) was added. After shaking of the reaction mixture, the absorbance was measured at 412 nm within 5 min of DTNB addition, against a blank with no homogenate. The absorbance values were extrapolated from a glutathione standard curve, and the level of GSH was expressed in grams of GSH per milligram of protein.

Inflammatory status. Gene expression of tumor necrosis factor alpha (TNF-α), interleukin-1β (IL-1β), IL-2, and gamma interferon (IFN-γ) was assayed using a CFX96 Touch detection system (Bio-Rad). Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein zeta polypeptide (YWHAZ) was used as the reference (28). DNA primers for all genes (Table 1) were designed on the basis of mRNA sequences obtained from the National Center for Biotechnology Information (http://www.ncbi.nlm.nih.gov; accessed 4 February 2011).

Real-time PCR assays were performed in a final volume of 25 µl containing 12.5 µl iQ SYBR green supermix (Bio-Rad), 200 nM (each) primers, and 1 µl cDNA from the sample. Negative samples were also tested, with the cDNA being replaced with autoclaved Milli-Q water. The PCR conditions were as follows: an initial denaturation period of 3 min at 95°C followed by 40 cycles of gene amplification. Each cycle consisted of an initial denaturation step of 20 s at 95°C, followed by an annealing step of 20 s at 60°C and an extension step of 45 s at 72°C. The samples were then subjected to an extension step of 3 min at 72°C.

To measure the specificity of the applied amplifications (i.e., to deter-
mine whether the formed products were specific for the tested genes, we performed a melting curve analysis in which the reaction temperature was increased 0.5°C every 15 s, beginning at the annealing temperature of the tested set of primers and ending at 95°C. Throughout the curve construction process, the changes in fluorescence were measured, and the data obtained, using CFX Manager software (version 3.0; Bio-Rad), were based on the values for the threshold cycle, i.e., the cycle where the observed fluorescence was 10-fold higher than the basal fluorescence for each quantitative PCR (qPCR) assay. Gene expression was obtained by applying the mathematical 2^(-ΔΔCt) method (29).

Histological and morphological analyses. The right kidneys were fixed with paraformaldehyde. Next, they were dehydrated with 70% ethanol and processed in paraffin. The resulting blocks were sliced into 5-μm-thick sections, stained with hematoxylin and eosin (H&E), and observed under a light microscope (×400).

Statistical analysis. Values were expressed as means ± standard errors of the means (SEM) for statistical analysis. One-way analysis of variance (ANOVA) followed by the Student-Newman-Keul post hoc test was used for parametric data. Gene expression data were evaluated by the Mann-Whitney test. P values of <0.05 were considered significant. Analysis was performed using GraphPad Prism 5.0.

RESULTS
Effect of GF on gentamicin-induced AKI. The GM group had a significant reduction in CLCR in comparison with both the CT and GF groups. This CLCR reduction was accompanied by increases in uric acid levels and the urine protein excretion rate (Table 2).

There was a progressive amelioration of CLCR and a serum creatinine (Scr) reduction in the GM+GF groups, with statistical significance at the 12.5- and 25-mg/kg doses. The 25-mg/kg dose appeared to confer better protection, so it was used in some experiments to explore GF’s protective effects.

Gentamicin-induced lipid peroxidation is attenuated by gingerosols. Gentamicin treatment increased MDA levels in renal tissue compared to those in the control group. In the GM + GF groups, there was a progressive reduction in the MDA level as the GF dose was increased. At the dose of 25 mg/kg, kidney MDA levels in the GM+GF group were similar to those in the control group (Fig. 1).

Effect of GF on gentamicin-induced nitrosative stress. The serum nitrite levels in kidney tissue were significantly elevated by GM administration. Animals in the GM + GF group receiving the GF dose of 25 mg/kg (GM+GF25 group) had serum nitrite levels that returned to control (CT) values (Fig. 2).

The protective effect of gingoressols is associated with increased antioxidant enzyme activity. The seven consecutive days of gentamicin treatment significantly reduced the levels of reduced GSH (Fig. 3) and SOD activity (Fig. 4). This reduction was significantly attenuated by oral treatment with GF at a dose of 25 mg/kg. GF alone at a dose of 25 mg/kg did not alter these parameters.

TABLE 2 Effects of GF on parameters of overall renal function in GM-induced nephrotoxicity

<table>
<thead>
<tr>
<th>Parameter</th>
<th>CT</th>
<th>GM</th>
<th>GM+GF (25 mg/kg)</th>
<th>GM+GF (6.25 mg/kg)</th>
<th>GM+GF (12.5 mg/kg)</th>
<th>GM+GF (25 mg/kg)</th>
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<tr>
<td>Serum creatinine (mg/dl)</td>
<td>0.62 ± 0.03</td>
<td>1.05 ± 0.08</td>
<td>0.51 ± 0.01#</td>
<td>0.87 ± 0.06</td>
<td>0.78 ± 0.07#</td>
<td>0.63 ± 0.03#</td>
</tr>
<tr>
<td>Creatinine clearance (ml/min)</td>
<td>1.02 ± 0.18</td>
<td>0.41 ± 0.08</td>
<td>0.97 ± 0.1#</td>
<td>0.55 ± 0.08</td>
<td>0.68 ± 0.09</td>
<td>0.88 ± 0.14#</td>
</tr>
<tr>
<td>Urea (mg/dl)</td>
<td>47.3 ± 2.61</td>
<td>80.6 ± 7.77</td>
<td>37.9 ± 2.8#</td>
<td>62.5 ± 6.7</td>
<td>53.8 ± 6.9#</td>
<td>41.7 ± 1.8#</td>
</tr>
<tr>
<td>Urinary protein (mg/dl)</td>
<td>40.5 ± 4.48</td>
<td>93.5 ± 9.28</td>
<td>44.5 ± 4.82#</td>
<td>78.3 ± 9.93</td>
<td>55.6 ± 7.74#</td>
<td>51.3 ± 3.24#</td>
</tr>
<tr>
<td>Uric acid (mg/dl)</td>
<td>1.33 ± 0.21</td>
<td>3.74 ± 0.52</td>
<td>1.57 ± 0.29#</td>
<td>2.5 ± 0.5</td>
<td>2.33 ± 0.71</td>
<td>1.66 ± 0.33#</td>
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*-, statistically significant (P < 0.05) compared to control (CT); #, statistically significant (P < 0.05) compared to GM.
Gingerol’s protective effects are associated with reduced inflammatory gene expression. To further explore GF’s protection-related mechanisms, we studied the expression of four mRNAs belonging to the innate and adaptive immune responses in kidney tissue samples (mRNAs for TNF-α, IL-1β, IL-2, and IFN-γ). As stated previously, only one group taking GM and GF was evaluated (GM+GF25 group). We observed that TNF-α, IL-1β, IFN-γ, and IL-1β expression increased after GM administration (Fig. 5) compared to that in controls (CT) \((P < 0.05)\). Animals in the GM+GF25 group had significant reductions in the expression of these genes only for TNF-α and IFN-γ. There was a tendency for mRNA expression of IL-1β and IL-2 to be blocked after GF treatment, but the difference was not statistically significant. Again, GF did not modify gene expression in animals not treated with GM.

Morphological analyses. In accordance with clearance experiments, morphological analyses demonstrated that the GM+GF groups had less severe hydropic degeneration of the proximal tubular epithelium than the GM group. Also, inflammatory infiltrates were found more often in GM-treated animals than in the GM+GF groups (Fig. 6).

**DISCUSSION**

In the present study, we demonstrated the protective effects of gingerols in a gentamicin-induced nephrotoxicity model. Attenuation of the GM-induced CLCR reduction by gingerols was associated with reduced oxidative stress, reduced nitric oxide metabolites, and reduced gene expression of important inflammatory cytokines. Previously, ginger extract has been evaluated only fairly in the context of GM-induced nephrotoxicity, with incomplete results (i.e., no data about the glomerular filtration rate—CLCR—or the mechanism associated with this protection) \((30)\).

In a recent report, about 30% of patients treated with GM for more than 7 days showed some sign of nephrotoxicity \((4, 31)\). The severe complications resulting from GM-induced nephropathy are limiting factors for its clinical use \((32, 33)\). Gentamicin-induced kidney injury is characterized by renal tubular epithelial cell necrosis, apoptosis, inflammatory responses, oxidative stress, and vascular contraction \((34, 35)\).

Gingerols are the most abundant pungent compounds in fresh ginger roots. These phenolic compounds have been cited in several studies as the main compounds responsible for the pharmacological effects of ginger, with the most abundant being [6]-gingerol \((36)\). These compounds have, among others, anti-inflammatory and antioxidant properties. These effects have been demonstrated in various organs obtained from diverse injury models. In the context of renal tissue, these compounds have been studied in relation to ischemia-reperfusion injury, cisplatin nephrotoxicity, renal damage caused by carbon tetrachloride, and diabetic nephropathy \((37–40)\).

In our model of GM-induced nephrotoxicity, the beneficial effects of gingerols were remarkable, with almost complete restoration of CLCR by use of a GF dose of 25 mg/kg. This prevention of the CLCR decrease was accompanied by less cell damage, as seen in the histological analysis, and by a reduction in the protein excretion rate. In proximal tubular damage, the renal tubule capacity to reabsorb normally filtered low-weight proteins is impaired. The near normalization of urine protein excretion seen in animals treated with GF is a marker of functional preservation of this tubular segment.

In the present study, we aimed to study several mechanisms associated with gingerol-induced nephroprotection. According to previous reports on the antioxidant effects of ginger extract, rats receiving GM and gingerols show a reduction in GM-induced oxidative stress. Lipid peroxidation is an initial event in the GM-induced nephrotoxicity injury cascade. This view was supported by an increase in MDA level (an index of lipid peroxidation), depletion of the kidney GSH content, and a decrease in SOD activity. In a recent study \((41)\), aminoglycoside and other bactericidal antibiotics induced oxidative damage to DNA, proteins, and membrane lipids. The antioxidant N-acetyl-L-cysteine ameliorates the oxidative stress-related effects. Oxidative substances have been implicated in GM-associated mitochondrial lesions in renal
proximal tubular cells as the possible mechanism through which gingerol and other antioxidant substances ameliorate GM-induced nephrotoxicity. Other substances with antioxidant properties have been tested in aminoglycoside-induced nephrotoxicity (42–44). In the present study, we used a model that approximates clinical practice: an in vivo study, protective substance administration initiated after the use of the toxic agent, and evaluation of functional parameters with clinical relevance (e.g., CLCR), in addition to histopathology, inflammatory gene expression, and oxidative status.

Vasodilator effects of NO appear to play a role in GM-induced nephrotoxicity (45). Moreover, NO seems to increase renal injury through its reaction with superoxide radical (O\(_2^-\)), generating the very cytotoxic peroxynitrite species (46). However, the detrimental effect of NO in GM-induced nephrotoxicity appears to be mediated only by inducible nitric oxide synthase (iNOS), not by its constitutive form (eNOS). While treatment with a specific inhibitor of iNOS, aminoguanidine, reduces GM-induced oxidative damage, eNOS inhibition can indeed aggravate it (47, 48). Unfortunately, we were not able to determine which NOS form was attenuated by gingerol administration. We suggest that its inhibitory effect is similar to that attributed to [6]-gingerol, as it inhibited NO production in lipopolysaccharide (LPS)-activated J774.1 macrophages and reduced iNOS protein levels in these cells (49).

**FIG 5** Effects of GF on relative expression of the TNF-α, IL-1β, IL-2, and IFN-γ genes in gentamicin-induced nephrotoxicity. Statistical analysis was performed by the Mann-Whitney test (n = 6). *, significant difference compared to saline (CT); #, significant difference compared to gentamicin (GM) (P < 0.05); **, significant difference compared to saline (P < 0.01); ##, significant difference compared to gentamicin (P < 0.01).

**FIG 6** Photomicrographs depicting H&E-stained sections of kidneys from rats receiving saline (control) (A), GM (100 mg/kg) (B and C), saline plus oral treatment with GF at a dose of 25 mg/kg (D), or gentamicin plus GF at 25 mg/kg (E and F). There was vacuolar degeneration in proximal tubule cells (B) and an inflammatory infiltrate (C) associated with GM administration. These features were attenuated by concomitant GF treatment (E and F). Magnification, ×400.
Reduction of NO synthesis can ameliorate GM-induced nephrotoxicity by itself, but it can also act through the reduction of oxidative stress caused by gingerols.

An increased or unbalanced ROS production and oxidative stress mediate the inflammatory response unleashed by GM. Superoxide anion and hydrogen peroxide activate NF-κB (6, 50), a key mediator of several inflammatory pathways, which induces the expression of proinflammatory cytokines and iNOS (35, 51, 52).

Gentamicin was associated with an increase in inflammatory gene expression of TNF-α, IL-2, IL-1β, and IFN-γ. This upregulation was attenuated by gingerols, except for that of IL-1β and IL-2, for which there was no statistical significance. It has been demonstrated that gingerols can inhibit TNF-α expression in the liver, nervous system, and other tissues (16, 53). Also, it is known that gentamicin-induced nephrotoxicity: do we have a promising therapeutic approach to blunt it? Pharmaco. Res. 62:179–186. http://dx.doi.org/10.1016/j.phrs.2010.04.004.

following the pathogenesis of tubulointerstitial impairment through the promotion of leukocyte attraction and adhesion to inflamed renal tubular cells. In our study, the reduction in inflammatory gene expression is in accordance with the reduced inflammatory infiltrate seen in GM+ GF25 animals.

Our data demonstrate a dose-dependent protective effect for all evaluated parameters. Although we were unable to determine the low or high threshold of the dose-effect curve, this relationship is relevant to future clinical studies and to understanding the effects of these substances on oxidative stress. We used a gingerol extract fraction with a predominance of [6]-gingerol, just as reported by others. Although it comprises more than 85% of all gingerols administered, we cannot attribute all beneficial effects to [6]-gingerol. Recently, it was demonstrated that the less abundant compound [10]-gingerol, not [6]-gingerol, was responsible for the anti-inflammatory effects on neuronal cells (53).

In conclusion, a gingerol-enriched fraction reduced GM-induced nephrotoxicity, and this protection was associated with reductions in oxidative stress and NO production and with inhibition of inflammatory gene expression.

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