Gentamicin Induced Apoptosis of Renal Tubular Epithelial (LLC-PK1) Cells

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Nephrotoxicity is a major limiting factor in the use of aminoglycoside antibiotics, the mechanisms for which are still speculative. To clarify the mechanisms of renal tubular cell death induced by aminoglycosides, we examined the renal proximal tubule-like cell line, LLC-PK₁, after inducing apoptosis through a chronic treatment with gentamicin (GM). Changes in the expression of the Fas were also investigated. On flow cytometric analysis, $5.7 \pm 3.3\%$ of the control cells appeared in a region of decreased forward light scatter and increased side light scatter, where both indices represent the characteristics of apoptotic cell death. Compared to the control, treatment with 10 mM of GM for 15 days significantly increased the proportion of cells in the apoptotic region to $23.9 \pm 8.5\%$. This finding was supported by electrophoretic analysis of the DNA extracted from the GM-treated cells, where a series of bands corresponding to integer multiples of 180 to 200 base pairs was visualized. However, the 15-day GM treatment did not cause a significant elevation in the expression of the 45 kD Fas protein, the cell surface molecule that stimulates apoptosis, by Western blot analysis.

In conclusion, long-term exposure to GM induces apoptosis of the renal tubular epithelial cells, and this process may contribute to some of the aminogly coside nephrotoxicities. Further studies are needed on the mechanism(s) of apoptosis induced by GM.

Key Words : Gentamicins; LLC-PK1 cells; Apoptosis

INTRODUCT IO N

Aminoglycoside antibiotics, including gentamicin (GM), are widely used in the treatment of gram-negative infections. A major complication of the use of these drugs is nephrotoxicity, accounting for 10% to 15% of all cases of acute renal failure^{1,2)}. Based on comparative clinical nephrotoxicity, GM is the most nephrotoxic drug²⁾. The specificity of GM for renal toxicity is apparently related to its preferential accumulation in the renal proximal convoluted tubules. Although derangement of many cellular alterations has been proposed, the specific mechanism whereby GM induces proximal tubular cell injury remains to be determined.

Apoptosis has recently been recognized as an important mechanism of cell death mediating chemical

and hypoxic injury to epithelial cells^{3,4)}. Apoptosis is an active form of cell death whose morphological features consist of both cytosolic and nuclear shrinkage, quite distinct from necrosis. Apoptotic cells and bodies are rapidly phagocytosed and destroyed by macrophages and neighboring cells, so apoptosis is usually inconspicuous in tissue sections even when **i** is responsible for extensive and rapid cell loss. Recently, apoptosis has been demonstrated to occur in the kidney following ischemic injury or partial ureteral obstruction³). Therefore, the role played by apoptosis in amino-glycoside nephrotoxicity has become an important issue. Whether or not apoptosis of the renal proximal tubular epithelial cells is induced by GM has not been previously studied.

In order to further clarify the mechanisms of renal tubular cell death induced by aminoglycosides, we examined the renal proximal tubule-like cell line, LLC-PK₁, after chronic treatment with GM to produce ap-

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optosis. The change in the expression of the Fas was also investigated.

MATERIALS AND METHODS

Reagents: Gentamicin sulfate, Dulbecco's modified Eagle's medium and fetal calf serum were obtained from Sigma Chemical Co. (St. Louis, MO, USA). The concentration of GM was determined based on the weight of GM sulfate salt in 100 mM Tris/HCl (pH 7.5).

1. Cell culture

The porcine proximal tubule-like cell line LLC-PK (ATCC CRL-1392) was obtained from the American Type Culture Collection (Rockville, MD, USA). The cells were cultured on 100 mm culture-grade plastic dishes in nutrient -modified Eagle's medium supplemented with 3% fetal calf serum (FCS) in an atmosphere of 5% CO₂ at 37 . The monolayers were plated at $3x 10^5$ cells per well on six-well plates and incubated four to six days until confluent. A fresh medium was provided on days three and five after plating. The experiments were performed on confluent cells between the 20th and 50th passages.

GM treatment: The confluent monolayers were washed three times in HBSS and provided with either a fresh medium (control) or a medium containing GM in various concentrations (0.1, 1, and 10 mM) and incubated for 15 days. Fresh study media were provided every three days until the cells were harvested for the apoptosis analysis.

2. The assessment of apoptosis

Fluorescence-activated cell sorter (FACS) analysis: A total of 950 μ L of cell suspension at a concentration of 10° cells/mL was incubated with 50 μ L propidium iodide (PI, 100 μ g/mL, Calbiochem-Novabiochem Co., La Jolla, CA. USA) in the dark at 4 . The cells were analyzed by FACScan flow cytometry (FACSort; Becton Dickson, USA). The selected gate corresponded to the cells according to the dot-blot histogram (10,000 to 50,000 cells were counted). The inter- and intra-assay variation coefficient was less than 6%.

DNA fragmentation on agarose gek: The DNA fragmentation assay was performed according to the

method published by Van de Water et al.61 Adherent cells were scraped into phosphate-buffered saline and pelleted by centrifugation at 200 g for 5 min. The spontaneously detached cells were processed separately and combined with the adherent cells before extracting a single fraction of DNA for electrophoresis. The cell pellets were lysed by overnight incubation in a 300 µL lysis buffer containing 10 mM Tris-HCl, 10 mM EDTA, 50 mM NaCl, 0.2% SDS, and 200 µg/mL proteinase K, pH 7.6 at 42 . The lysate was added to the pellet of the pooled washings and placed on ice for 15 min. The lysates were centrifuged at 13,000 g for 20 minutes and the fragmented DNA present in the supernatant was extracted with phenol-chloroformisoamyl(25:24:1) and ethanol precipitation. The pellet was resuspended in 20-30 µL of Tris-EDTA buffer containing RNAse A, incubated for 30 minutes at 37 and then separated by electrophoresis on a 1% agarose gel containing ethidium bromide. The DNA was visualized by ultraviolet examination. The size of the DNA fragments was compared with a 123-bp standard DNA ladder.

3. Fas expression

Each well was washed with ice-cold phosphatebuffered saline and treated with a lysis buffer (150 mM NaCl, 1% NP-40, 0.5% DOC, 50 mM Tris, and 0.1% SDS, pH 8.0) including protease inhibitors for an hour on ice. The lysed cells were centrifuged at 12,000 g for 10 minutes at 4 . Twenty micrograms of the cell extracts were resolved on 10% SDS-PAGE. The separated proteins were then transferred to a nitrocellulose membrane. After blocking with 5% nonfat dry milk, the membrane was incubated for one hour with 1:1000 polyclonal (rabbit) anti-Fas antibody (Santa Cruz Biotechnology, Santa Cruz, CA, USA) diluted in 5% milk non-fat proteins/Tris-buffered salinetween buffer. The blot was washed again and incubated with a goat anti-rabbit horseradish peroxidase conjugated secondary antibody (Amersham, Pharmacia Biotech UK Ltd., Buckinghamshire, England). The blot was developed using enhanced chemiluminescence.

4. Statistical analysis

The data were expressed as $mean \pm S.D$. The statistical significance was assessed using a one-way analysis of variance combined with Scheffe's test and a probability level of p < 0.05 was regarded as significant.

RESULTS

1. Effects of GM on apoptosis

On the flow cytometric assessment of apoptosis using light scatter measurements, $5.7 \pm 3.3\%$ of the control cells appeared in a region of decreased forward light scatter (FSC) and increased side light scatter (SSC), where both indices represent the characteristics of apoptotic cell death. Compared to the control, treatment with GM for more than 12 days significantly increased the proportion of cells in the apoptotic region in a dose dependent manner (Table 1, Figure 1). The maximum apoptosis occurred at 15 days post-treatment (5.7 ± 3.3% vs $23.9 \pm 8.5\%$ at 10 mM GM, p < 0.05). No significant apoptosis was observed in the cells treated with GM for less than 12 days.

These findings were supported by electrophoretic analysis of the DNA extracted from the GM-treated cells. The DNA extracted from the GM-treated cells was visualized as a series of bands corresponding to integer multiples of 200 base pairs that are characteristic of apoptosis (Figure 2).

2. The effect of GM on Fas expression

To assess the changes in the Fas expression following GM exposure, Western blots were performed on the cells exposed to 10 mM GM for 7 to 15 days every 24 hours. Although there was a significant increase in the number of apoptotic cells, there as no significant difference in the expression of the 45 kD Fas protein between the GM-treated and control cells (Figure 3).



% Apoptosis
5.7 ± 3.3
4.8 ± 4.1
9.2 ± 5.8
$23.9 \pm 8.5*$

*: p < 0.05, compared to control

Cells were treated with gentamicin for 15 days.



Figure 1. Flow cytometric assessment of the effects of gentamicin (GM) on the size and granularity of the LLC-PK cells. Approximately 5% of the control cells appeared in a region of decreased forward light scatter and increased side scatter, where both indices represent the characteristics of apoptotic cell death (A). Treatment with GM at the indicated concentrations for 15 days increased the proportion of cells in the apoptotic region in a dose- dependent manner (B: GM 1 mM, C: 10 mM).

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Figure 2. The effect of gentamicin on DNA fragmentation in the LLC- $PK_{\rm L}$ cells.

Lane 1: 123-base pair standard ladder, lane 2: control, lane 3: GM 10 mM $\,$

Gel electrophores is of the DNA extracted from the GM-treated cells showed a DNA ladder pattern characteristic of apoptos is.

DISCUSSION

Aminoglycosides are polycationic compounds that are not metabolized and are minimally protein bound¹⁾. After glomerular filtration, aminoglycosides bind to the apical membrane of renal proximal tubular cells and are internalized by adsorptive endocytosis⁷⁾. After uptake into the cells, a number of intracellular processes are disrupted by the presence of aminoglycosides. Although the mechanism of aminoglycosideelicited renal proximal tubular cell injury has been



Figure 3. The effect of gentamicin on Fas expression in the LLC-PK cells. (A) Western blot analysis of the Fas expression. Lane 1: control, Lane 2: GM 10 mM for 15 days. (B) The results of a densitometric analysis of four independent experiments were shown. There was no significant difference between the control and GM (10 mM) treatment (p> 0.05).

investigated in many studies, the exact biochemical determinant of tubular cell injury remains unknown.

Apoptosis is a physiologic, programmed cell death that occurs under both physiological and pathological conditions. In particular, apoptosis has been reported to be an important mechanism that mediates ischemic, chemical hypoxic and oxidant injury of renal tubules⁸). Duncan-Achanzar et al.⁹⁾ reported that apoptosis in response to 35 μ M HgCl was confirmed by observing the morphological features characteristic of apoptotic cells. Lieberthal et al.¹⁰⁾ also suggested apoptosis as the mechanism of death induced by cis-platin in proximal tubular epithelial cells. In the present study, we demonstrated that apoptosis occurred in LLC-PK cells after treatment with GM for 15 days. Nakagawa et al.¹¹⁾ reported the GM-induced apoptosis of vestibular hair cells of guinea pigs. To our knowledge, our finding is the first evidence of GM-induced apoptosis in renal tubular epithelial cells.

Chronic exposure of renal tubular epithelial cells to high levels of GM leads to the accumulation of GM within the lysosomes accompanied by cell injury. Ford et $al^{(12)}$ reported that LLC-PK₁ cells exposed to 2 mM GM for six days displayed a significant decrease in the specific activities of lysosomal and apical membrane enzymes with an increased release into the medium. Hori et al.13) and Schwertz et al.14) also evaluated the alterations in enzyme activities and cellular lipid levels, as well as performing a morphological examination of renal tubular epithelial cell injuries, after exposure to GM for three and seven days. In our study, treatment with GM at a concentration of 10 mM for 15 days induced apoptosis in the LLC-PK cells. Based on these findings, we postulate that apoptosis occurs in renal tubular epithelial cells after long-term exposure to GM, and this process may be crucial for cell injury, subsequent, and eventual regeneration. It remains to be determined, however, whether the 10 mM of GM can be achieved in the renal proximal tubular cells in experimental animals and in humans treated with GM.

The process leading to apoptosis is complex and various primary biochemical events that may initiate the apoptosis process are believed to exist including deprivation of growth factors, mitochondrial injury, oxidative stress, F-actin cytoskeletal injury and perturbation of intracellular Ca2+ homeostasis4.8). The mechanism(s) responsible for apoptosis in GM-induced injury to LLC-PK cells is not known. The ability of GM to alter mitochondrial respiration has been well documented in reports of both in vitro and in vivo studies^{15,16)}. Walker and Shah¹⁵⁾ observed that the generation of hydrogen peroxide by the mitochondria was enhanced from less than 0.5 nmol in the absence of GM to more than 6 nmol/mg/min in the presence of GM. Recently, GM has also been shown to enhance the generation of superoxide anion and hydroxyl radical in renal cortical mitochondria. Reactive oxygen species play an important role in the activation of the endonuclease and consequent DNA damage4,10). Based on these findings, we postulated that apoptosis may occur through the enhanced generation of reactive oxygen metabolites following exposure to GM.

We have determined experimentally that the DNA

ladder pattern characteristic of apoptosis occurs in the LLC-PK1 cell monolayers 14 days after 10 mM GM exposure. This result demonstrates that the endonuclease-mediated cleavage of the DNA into nucleosome-sized fragments occurs after long-term exposure to GM. Reactive oxygen species have been reported to play an important role in the activation of endonuclease and consequent DNA damage. In addition, Kozek et al.¹⁷⁾ reported that the treatment with GM upsets the process of intracellular Ca2+ homeostasis, which may induce apoptosis due to its activation by calcium of the Ca^{2+}/Mg^{2+} -dependent endonucleases that are involved in the cleavage of DNA. Therefore, it can be postulated that the activation of endonuclease following the perturbation of cytosolic calcium caused by GM is another factor for producing apoptosis.

The Fas and its ligand are important mediators of apoptosis and show several structural and functional similarities to the tumor necrosis factor receptor system^{8,18)}. The ability of the Fas to induce death is dependent on the balance between the expression of death factors, such as the interleukin-1-converting enzyme and the expression of protective factors, such as Bcl-2 and Fas-associated phosphatase-18). In the LLC-PK cells, a cyclosporine A treatment caused an increase in the expression of the 45 kD Fas protein. Healy et al.¹⁸⁾ suggested that the Fas may be an important mediator of cyclosporine A-induced apoptosis in renal proximal tubular cells. In this study, however, we were unable to observe that the GM exposure was associated with an increase in the expression of the Fas protein.

The use of whok-animal models to study the primary aminoglycoside-induced proximal tubule cell injury is complicated by the structural and functional heterogeneity of the kidney. Furthermore, tubular cell necrosis resulting from aminoglycoside exposure is focal in nature, and cellular regeneration coexists temporally¹⁹⁾. Many of these complicating factors can be eliminated by the use of a cultured cell model. The LLC-PK₁ cell line has renal proximal tubule cell-like properties, including high levels of proximal tubular marker enzymes⁹⁾. Hori et al.¹²⁾ concluded from their investigation that the LLC-PK₁ cell line could be a useful model for studying aminoglycoside nephrotoxicity in vitro. Thus, we have used the LLC-PK₁ cells as a model system for GM nephrotoxicity.

In conclusion, long-term exposure to GM induces

apoptosis of the renal tubular epithelial cells and this process may contribute to certain aminoglycoside nephrotoxicities. Further studies are needed on the mechanism(s) of apoptosis induced by GM.

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