



Effects of repeated ochratoxin exposure on renal cells *in vitro*

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Abstract

In the present study an *in vitro* model of subchronic repeated exposure to OTA and OTB was employed to generate ochratoxin-derived subpopulations of human and porcine proximal tubular cells (HKC, IHKE, PKC, LLC-PK1). These cell subpopulations were subsequently used to investigate effects on cell proliferation rates, expression of marker proteins (cytokeratins, vimentin) and the acute cytotoxicity of OTA and OTB (MTT reduction, neutral red uptake, cell number). The hypothesis was tested whether repeated exposure at moderate concentrations of these toxins could provide for a reduced sensitivity of selected cell subpopulations to subsequent toxin exposure. Despite the observed increased cell population doubling times and the reduced sensitivity toward OTA and OTB exposure of some cell types, with the exception of the primary human epithelial cells, no overt changes in the expression of cytokeratin and vimentin could be determined. The presented data, however suggest that repeated exposure of renal epithelial cells to ochratoxins A or B will provide for a subpopulation of cells with reduced ochratoxin-sensitivity and alterations in growth characteristics.

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1. Introduction

Ochratoxins are secondary metabolites of certain ubiquitous *Aspergillus* and *Penicillium* species and are known contaminants of human food and animal feed often arising due to sub-optimal storage conditions (Speijers and van Egmond, 1993; Studer-Rohr, 1995). Chronic dietary exposure to ochratoxin A (OTA) has long been associated with the renal disease, known as Balkan Endemic Nephropathy (BEN) and with an increased incidence of urothelial tumors in humans. Presently, the hypothesis of BEN being an environmentally acquired disease with ochratoxins being one of the principal potential etiological factors (others include the aristolochic acid or the Pliocene lignite hypotheses), is widely accepted (Stefanovic et al., 2006; Pfohl-Leszkowicz et al., 2002; Tatu et al., 1998). The pathological characteristics of BEN include diffuse cortical fibrosis and degenerative changes of the renal epithelium. Not only humans can be

affected by exposure to ochratoxins. Indeed many similarities, including atrophic proximal tubules and fibrotic tissue, have been reported between the later stages of BEN and mycotoxin-induced porcine nephropathy (MPN) (Stoev, 1998). Rásony and co-workers reported a similar OTA-induced loss of proximal tubular cells, accompanied with increased regenerative processes, in rats following oral OTA exposure (Rásonyi et al., 1999).

The mechanism of OTA-mediated cell death remains a question of debate (for review see Dekant et al., 2005; O'Brien and Dietrich, 2005), but it seems that in all species investigated to date, including humans, the kidney attempts to balance cell loss with regenerative processes and the question of whether this loss of tissue can be replaced via proliferation of epithelial or fibroblastic neighboring cells remains unanswered. Regeneration is most likely driven by a combination of migration and/or proliferation and subsequent cell differentiation after sublethal injury. Consequently, this could lead to the appearance of an altered cell type, exhibiting neither functional nor morphological characteristics of the original cell type and as such, would be consistent with the genesis of fibrosis and of renal tumors.

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A complete outline of renal regeneration processes following acute injury is beyond the scope and intention of this research manuscript, however a good overview of these mechanisms is given by Nony and Schnellmann (2003).

Previous *in vitro* observations have demonstrated that even after acute exposure to highly toxic concentrations of ochratoxins ($\geq 10 \mu\text{M}$), approximately 10% of cells survive and can recommence proliferation (O'Brien et al., 2001). These cells have also been described to display a more fibroblast-like rather than epithelial appearance (Heussner et al., 2002). The latter findings, coupled with the observations that fibroblasts are less susceptible to ochratoxin-mediated toxicity (Grotegut et al., 2001; O'Brien et al., 2001), gave rise to the hypothesis that repeated exposure to ochratoxins alters normal renal epithelial cells, possibly converting them to a more fibroblast-like nature, thus, providing the rudiments for renal fibrosis (van Kooten et al., 2001). Support for this hypothesis was presented by the recent publication of Sauvants et al. (2005), who described an increase in collagen synthesis in OK cells and primary human renal proximal tubular epithelial cells (RPTEC) following exposure to OTA. All of these observations are mostly descriptive in nature and need to be further characterized. If indeed changes toward a more resistant cell type are involved in the regenerative processes following OTA insult, these should be reflected by alterations in parameters such as morphology, growth rate, protein expression and susceptibility to acutely cytotoxic concentrations of ochratoxins. It was therefore the aim of this study to determine whether OTA and/or OTB treatment of renal proximal tubular cells will result in the latter alteration of parameters.

For this, a model of repeated ochratoxin-induced cell loss with alternating regenerative periods was employed in this study in order to mimic the dietary situation of (sub)chronic interval exposure in humans and animals. Using this approach, various renal cell subpopulations of continuous renal cell lines and primary cells of porcine and human origin were generated and characterized with regard to morphology, growth rate, protein expression and susceptibility to acutely cytotoxic concentrations of ochratoxins.

2. Material and methods

2.1. Mycotoxins and reagents

Highly purified ochratoxin A and B (>98% purity, benzene free) were kindly provided by M.E. Stack, US FDA, Washington, DC. Sterile ochratoxin stock solutions were prepared in 0.1 M NaHCO_3 and freshly diluted (dilution factor = 40) in culture medium immediately before exposure. Media, FBS and other cell culture chemicals were purchased from PAA Laboratories GmbH, Cölbe, Germany. MEM/D-Val medium was purchased from Life Technologies, Karlsruhe, Germany. Unless otherwise sta-

ted, all other chemicals were purchased from Sigma-Aldrich GmbH, Seelze, Germany.

2.2. Biopsy material and animal tissue

Human renal biopsy material (male patients) was obtained from a local hospital (Klinikum Konstanz) in collaboration with the Department of Urology, Prof. Dr. R. Thiel, subsequent to informed patient consent. Tissue samples as well as patient data were handled in compliance with the stipulations put forth by the ethics committee of the University of Konstanz and in fulfillment of German law and the Declaration of Helsinki (1964) pertaining to personal data protection and handling of human biopsy material.

Whole kidneys from freshly-killed male improved German hybrid pigs were obtained from a local slaughterhouse.

2.3. Cell cultures

Primary male human kidney cells (HKCm) and primary male porcine kidney cells (PKCm) were prepared by standard collagenase digestion from renal cortices and cultured as previously described in detail (Dietrich et al., 2001). Briefly, minced renal cortices were incubated with buffered collagenase type 1 (1 mg/ml) at 37 °C for 30 min with gentle stirring and subsequently filtered through gauzes (200 and 40 μm). Cultures were initiated in Primaria tissue culture flasks (Becton Dickinson GmbH, Heidelberg, Germany) using MEM D-Val medium supplemented with 10% FBS and antibiotics (100,000 units/L penicillin, 100 mg/L streptomycin) to prevent fibroblast growth. Once small cell colonies were established, MEM D-Val was replaced by DMEM/F12 medium, supplemented with FBS and antibiotics.

LLC-PK1 cells were obtained from the European Collection of Cell Cultures, Salisbury, UK (ECACC No. 86121112) and IHKE cells were kindly provided by S. Mollerup, National Institute of Occupational Health (STAMI), Oslo, Norway. Both cell lines were cultured in DMEM, supplemented with 10% FBS and antibiotics as for primary cells.

All four cell types were cultured under standard conditions (37 °C, 5% CO_2).

2.4. Generation of ochratoxin cell subpopulations (study part A)

IHKE, LLC-PK1, HKCm and PKCm of passage number 4, 8, 1 and 1, respectively, were seeded at a density of 1×10^4 cells/cm² into 182 cm² culture flasks in their appropriate medium. Medium was replaced every 48 h. As indicated by preliminary experiments, after reaching 80–90% confluence, cells were exposed to nominal concentrations of 20 and 50 μM OTA or OTB until approximately 90% of the cells had died (depending on cell type, approximately

72–96 h, ochratoxins were replenished with each medium change). These concentrations were chosen on the basis of earlier cytotoxicity findings with 20 μM of either, OTA and OTB, representing a moderately toxic concentration ($\sim\text{EC}_{50}$) and 50 μM being highly toxic ($\geq\text{EC}_{80}$), depending on endpoint and exposure duration (Dietrich et al., 2001).

The ochratoxin-containing medium was then discarded, the cells were washed once with PBS (136.9 mM NaCl, 2.7 mM KCl, 8.1 mM Na_2HPO_4 , 1.5 mM KH_2PO_4 , pH 7.4, ≤ 300 mOsm) and further cultured with fresh medium until approximately 90% confluence was attained. This regimen of exposure and recovery was repeated a further two times to generate subpopulations of ochratoxin-exposed cells. In parallel, vehicle control cultures were exposed to a final concentration of 2.5 mM NaHCO_3 and passaged at approximately 90% confluence using trypsin/EDTA in order to provide comparable control cells. The resulting cell cultures were designated subpopulations after the third exposure interval or passage and used for further experimentation (see Table 1 for overview of subpopulations and Fig. 2 for repeated exposure time schedule).

2.5. Determination of growth characteristics (study part B)

For the determination of growth characteristics of all subpopulations, growth curves were established using a Coulter counter (Coulter Z1, Beckman Coulter GmbH, Krefeld, Germany) as previously described (Dietrich et al., 2001).

2.6. Immunodetection of cytokeratins and vimentin (study part B)

Confluent cell subcultures were harvested using trypsin/EDTA and intermediate filaments were isolated according to standard procedures (Eckert and Kartenbeck, 1997). Proteins were separated by 12% SDS-PAGE and electroblotted onto nitrocellulose membranes. Immunostaining was performed using mouse anti-cytokeratin (clone AE1/AE3, Biomedica Corp., Hayward, CA, USA), mouse anti-vimentin (clone V9, Biomedica Corp., Hayward, CA, USA) and HRP-conjugated rabbit anti-mouse IgG (Dako-Cytomation GmbH, Hamburg, Germany, cat. no. P0260) with subsequent chemiluminescent (ECL, Amersham Biosciences, Uppsala, Sweden) and colorimetric (DAB sub-

strate kit, Roche Diagnostics GmbH, Mannheim, Germany, cat. no. 1718096) detection according to the manufacturers' instructions.

2.7. Acute challenge experiments with ochratoxins (study part C)

Cell culture subpopulations were seeded at 1×10^4 cells/ cm^2 in serum replete medium, on normal tissue culture plastics for experiments as described previously (Dietrich et al., 2001). Exposure to OTA or OTB, each at nominal concentrations of 0.1, 1, 5, 10, 15, 20 and 50 μM or to vehicle (2.5 mM NaHCO_3) for 24, 48, 72 and 96 h began 24 h after seeding.

MTT reduction (MTT), neutral red uptake (NR) and cell number were assessed as previously described (Dietrich et al., 2001).

2.8. Calculations and statistical data analysis

Cytotoxicity data are presented as integrated best-fit IC_{50} (24–96 h) values (50% inhibitory concentrations) from non-linear regression $\pm 95\%$ confidence intervals based on asymptotic standard errors. These data were generated using GraphPad Prism version 4.0 for Windows and Excel software (Microsoft Excel 2000) from the mean percentage inhibition for each endpoint at each test substance concentration from at least three independent experiments carried out in duplicate. Values were calculated as the difference between the area under the control curve and the area under the curve for each substance concentration, integrated over 24, 48, 72 and 96 h, using the following equations:

$$A = \frac{N_1}{2} \cdot t_1 + \frac{N_1 + N_2}{2} \cdot (t_2 - t_1) + \frac{N_3 + N_4}{2} \cdot (t_4 - t_3) \quad (1)$$

$$I_A = \frac{A_c - A_t}{A_c} \cdot 100 \quad (2)$$

with A , area under the curve; N_n , endpoint value (% control MTT reduction, % control NR uptake or % control cell number) at timepoint n ; t_n , timepoint n (24, 48, 72 or 96 h); I_A , % inhibition; A_c , area under the control curve and A_t , area under the toxin curve.

Only intrapolated IC_{50} values are reported, extrapolated data, where calculated values were higher than the tested concentration range, were presented only as " \geq " or " $>$ " maximum tested concentration in order to avoid speculation with mathematically correct, but biologically irrelevant data.

Subpopulation growth characterization data were calculated as arithmetic means \pm SEM from three independent experiments carried out in duplicate.

Significance of differences compared to control cells or between subpopulations was tested using a one-way ANOVA followed by Tukey's multiple comparison test.

Table 1
Overview of cell culture subpopulations

Subpopulation no.	Repeated (3 \times) exposure to
Untreated	Vehicle
lowOTA	20 μM OTA
highOTA	50 μM OTA
lowOTB	20 μM OTB
highOTB	50 μM OTB

Vehicle = 2.5 mM NaHCO_3 .

3. Results

3.1. Generation of cell subpopulations (study part A)

The described study design (Fig. 1, part A) of repeated exposure of four different renal cells to ochratoxins resulted in five subpopulations for each cell type (Table 1). Routine microscopy of cell cultures (with approximation of level of confluence) showed that during the exposure intervals cells became less sensitive to ochratoxins as illustrated for IHKE cells in Fig. 2. This loss of sensitivity was time-dependent and more prominent for cell lines than for primary cells and was further characterized in subsequent experiments with the individual subpopulations as described in detail below.

Overt morphological transformations were not evident in the subpopulations generated by repeated exposure to ochratoxins.

3.2. Growth characteristics of cell subpopulations (study part B)

The results of growth characterization are summarized in Fig. 3. With the exception of the lowOTA-subpopulation (see Table 1), which was completely unaffected, HKCm subpopulations displayed an increased population doubling time (Fig. 3A), i.e. from 31.0 ± 0.5 (untreated) to 43.0 ± 3.4 h (highOTA) (see Table 1). Similarly, highOTA selection provided for overtly increased doubling time in LLC-PK1.

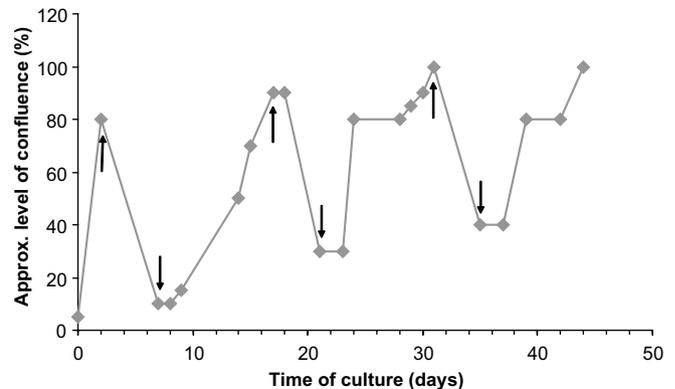


Fig. 2. Representative repeated exposure scheme. Exposure scheme of IHKE (lowOTA-subpopulation); ↑ start of exposure interval; ↓ end of exposure interval. The level of confluence was approximated during routine microscopy.

It was impossible to determine the doubling time for either HKCm OTB-subpopulation as it was higher than the time frame tested (96 h).

In contrast, PKCm subpopulations showed no changes in doubling time compared to untreated control cells. Interestingly, all IHKE subpopulations displayed a decreased population doubling time.

All PKCm and LLC-PK1 subpopulations showed an up to 50% reduction in cell number, when total cell number in the plateau phase was considered (Fig. 3B). This picture was not as consistent for HKCm cells, where the highOTA-subpopulation showed an increased total cell

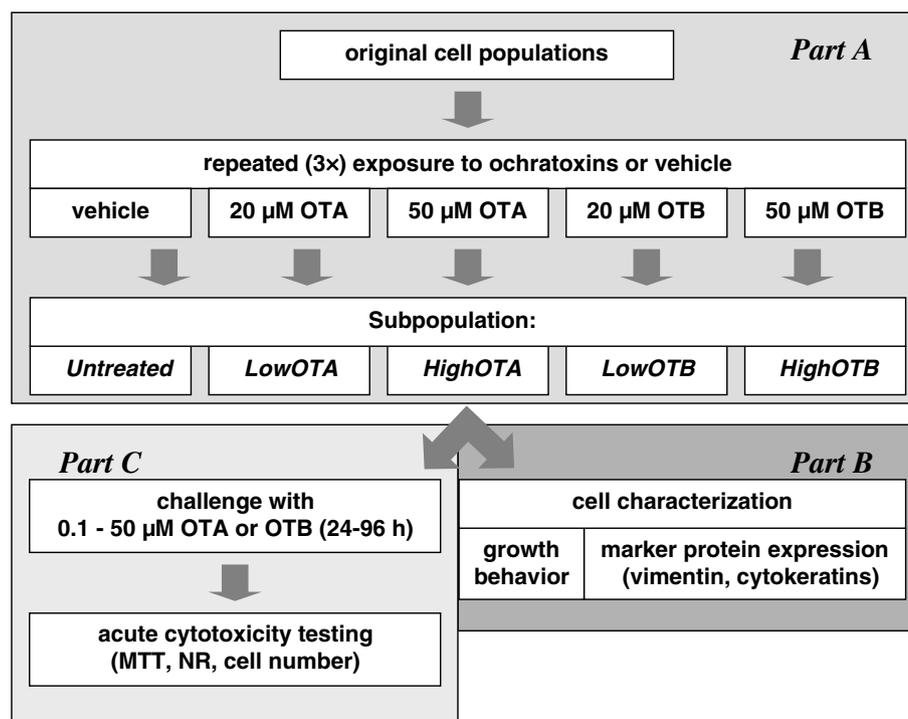


Fig. 1. Study design. Study part A: generation of cell subpopulations; study part B: characterization of cell subpopulations; study part C: determination of differential acute cytotoxic effects of ochratoxins in cell subpopulations.

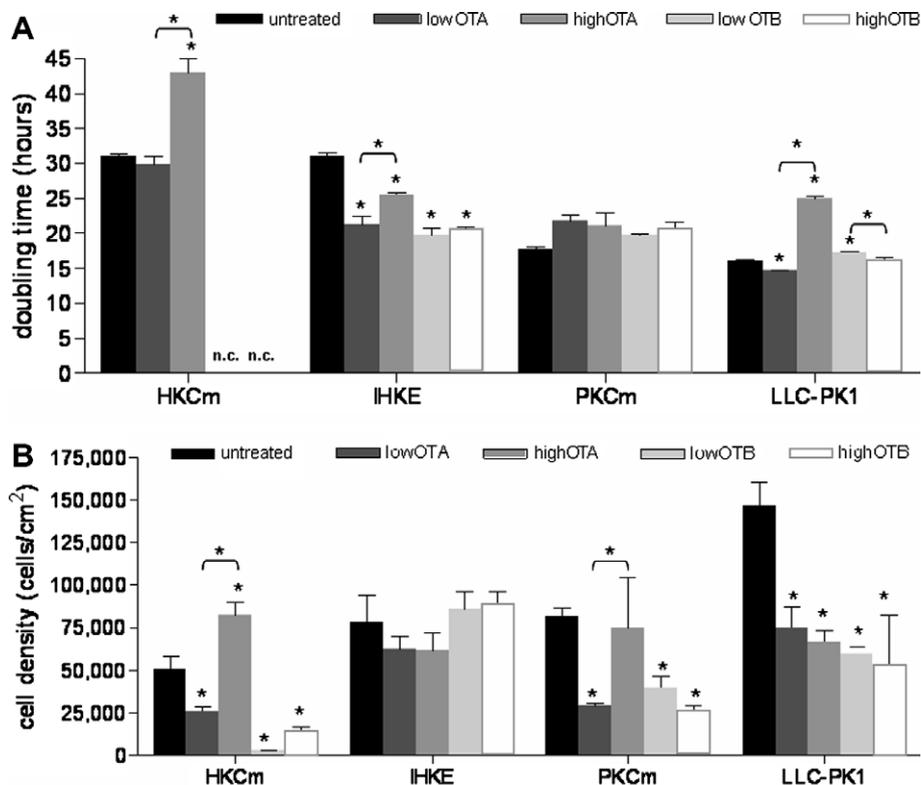


Fig. 3. Growth characteristics. A: Cell doubling times, B: cell densities in the plateau phase. Values represent the mean \pm SEM ($n = 3$). Significant differences compared to control or within groups were tested using a one-way ANOVA followed by Tukey's multiple comparison test and are indicated ($*p < 0.05$). Cell subpopulations are named as outlined in Table 1. n.c., not computable due to doubling times surpassing the 96 h window of experiment.

number. All other HKCm subpopulations displayed a decreased cell number. In contrast, none of the IHKE subpopulations displayed any significant difference in the total number of cells in the plateau phase.

In summary, repeated exposure to ochratoxins resulted in altered growth behavior in all four cell types.

3.3. Protein expression in cell subpopulations (study part B)

The expression of two common cytoskeletal marker proteins for cellular origin, namely cytokeratin and vimentin, were chosen as endpoints. Representative immunodetection results are shown in Fig. 4. The monoclonal mouse anti-human cytokeratin antibody used can potentially detect up to eleven different cytokeratins of the basic and the acidic subfamily in the range of 40–67 kD in a wide spectrum of human epithelial cells. This is dependent on the specific cell type and species of origin. In this study, a maximum of four specific bands could be visualized in human cells and only one in porcine cells (Fig. 4).

The HKCm subcultures highOTA, lowOTB and highOTB demonstrated a reduction in cytokeratin expression in comparison to their corresponding control. None of the other cell subpopulations displayed any difference in cytokeratin expression, which could be considered to be beyond normal protein loading and staining variance.

Vimentin expression was detected in the same manner in parallel. All control cells showed only very low expression

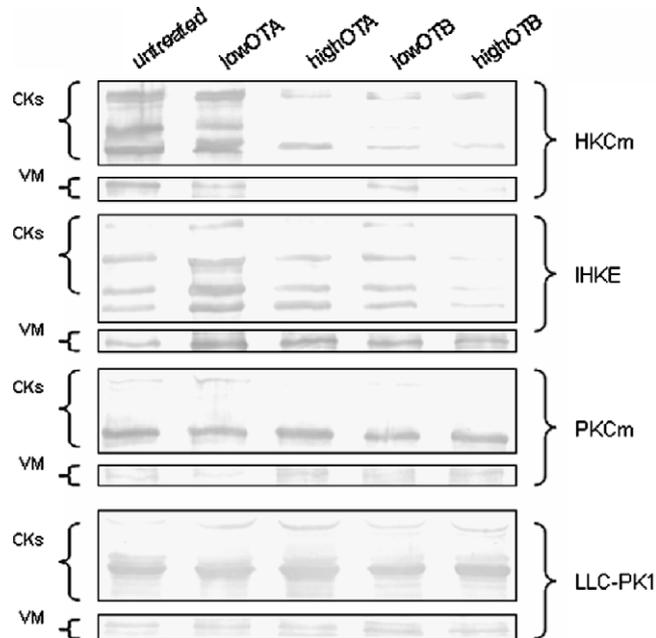


Fig. 4. Protein expression. Western blot analysis with DAB detection of cytokeratins (CKs) and vimentin (VM) expression in the cell subpopulations of HKCm (3 μ g total protein/lane), IHKE (4 μ g total protein/lane), PKCm (6 μ g total protein/lane) and LLC-PK1 (10 μ g total protein/lane). The monoclonal mouse anti-human cytokeratin antibody employed can potentially detect up to eleven different cytokeratins of the basic and the acidic subfamily in the range of 40–67 kD in a wide spectrum of human epithelial cells.

of vimentin. No increase could be detected in the ochratoxin subpopulations.

In summary, repeated exposure to ochratoxins resulted in altered cytokeratin expression in primary human cells only.

3.4. Cytotoxic effects of acute challenge exposure to ochratoxins (study part C)

Comparison of absolute IC₅₀ values (Tables 2 and 3) for both toxins of the four control cell culture subpopulations showed species- and cell type-related differences in relative endpoint sensitivity. MTT reduction was the most sensitive

endpoint for both human cell types (HKCm and IHKE), irrespective of the toxin used. In contrast, for porcine cells, cell number seemed to be a slightly more sensitive endpoint than either MTT reduction or NR uptake. Unless otherwise stated, in the following sections, the results of the control cells and the subpopulations are always described and compared using the most sensitive endpoint for the respective cell type.

OTA and OTB in control cells:

Primary porcine cells (PKCm) and the continuous porcine cell line (LLC-PK1) displayed similar sensitivities to OTA and OTB, however, PKCm seemed to be

Table 2
Results of cytotoxicity testing with OTA: comparison of integrated IC₅₀ values (24–96 h, μM)

Cell cultures	Endpoints	Subpopulation										
		Untreated			lowOTA		highOTA		lowOTB		highOTB	
		IC ₅₀	IC ₅₀	Mean fold change								
HKCm	NR	>50	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0
	MTT	13.1 ± 2.1	21.8 ± 10.8	1.7*	17.0 ± 6.9	1.3	≥50.0	≥3.8	26.7 ± 14.9	2.0*		
	CN	>50	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0
IHKE	NR	15.4 ± 11.8	>50	>3.3	>50	>3.3	>50	>3.3	>50	>3.3	>50	>3.3
	MTT	4.1 ± 1.4	7.4 ± 2.0	1.8	6.3 ± 1.4	1.5	6.1 ± 1.6	1.5	4.2 ± 1.8	1.0		
	CN	6.4 ± 4.6	7.1 ± 3.5	1.1	8.7 ± 3.5	1.4	8.2 ± 5.6	1.3	4.9 ± 2.5	0.8		
PKCm	NR	13.4 ± 5.0	49.2 ± 23.8	3.7*	27.5 ± 24.5	2.1*	>50	>3.7	40.1 ± 21.5	3.0*		
	MTT	10.2 ± 3.6	17.5 ± 6.4	1.7*	16.0 ± 3.9	1.6*	11.2 ± 3.2	1.1	10.6 ± 5.4	1.0		
	CN	8.1 ± 3.2	20.6 ± 7.0	2.5*	19.3 ± 4.2	2.4*	11.5 ± 3.9	1.4	17.7 ± 6.1	2.2*		
LLC-PK1	NR	24.1 ± 10.9	44.5 ± 24.5	1.8*	>50	>2.0	18.4 ± 5.6	0.8	>50	>2.0		
	MTT	23.0 ± 8.8	24.0 ± 14.6	1.0	19.8 ± 10.9	0.9	18.5 ± 6.8	0.8	22.7 ± 10.2	1.0		
	CN	13.9 ± 2.6	29.0 ± 8.7	2.1*	33.0 ± 10.2	2.4*	14.4 ± 5.0	1.0	14.8 ± 4.7	1.1		

Values represent integrated best-fit IC₅₀ (24–96 h, μM) of OTA from non-linear regression ±95% confidence intervals based on asymptotic standard errors. Significant differences compared to control are indicated (**p* < 0.05). ≥, values are at least as high as control, but higher than the tested concentration range (i.e. >50 μM); >, values are higher than control values, but higher than the tested concentration range.

Table 3
Results of cytotoxicity testing with OTB: comparison of integrated IC₅₀ values (24–96 h, μM)

Cell cultures	Endpoints	Subpopulation										
		Untreated			lowOTA		highOTA		lowOTB		highOTB	
		IC ₅₀	IC ₅₀	Mean fold change								
HKCm	NR	>50	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0
	MTT	15.2 ± 13.1	20.2 ± 5.1	1.3*	37.3 ± 21.0	2.5*	>50	>3.3	>50	>3.3	>50	>3.3
	CN	>50	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0	>50	≥1.0
IHKE	NR	19.5 ± 11.5	>50	>2.6	>50	>2.6	47.9 ± 10.4	2.5*	>50	>2.6	>50	>2.6
	MTT	4.6 ± 2.0	10.8 ± 5.2	2.3*	12.6 ± 7.1	2.7*	5.2 ± 2.2	1.1	7.8 ± 4.0	1.7		
	CN	8.5 ± 4.1	9.9 ± 3.9	1.2	10.1 ± 4.8	1.2	12.2 ± 5.6	1.4	9.6 ± 3.8	1.1		
PKCm	NR	21.5 ± 9.2	>50	>2.3	>50	>2.3	>50	>2.3	47.2 ± 24.0	2.2*		
	MTT	8.0 ± 2.9	>50	>6.3	25.4 ± 8.7	3.2*	18.9 ± 4.9	2.4*	27.5 ± 6.7	3.4*		
	CN	8.8 ± 2.9	33.5 ± 10.3	3.8*	>50	>5.7	10.4 ± 3.2	1.2	30.2 ± 6.9	3.4*		
LLC-PK1	NR	>50	>50	≥1.0	>50	≥1.0	32.0 ± 13.0	≤0.6	>50	≥1.0		
	MTT	>50	>50	≥1.0	>50	≥1.0	>50	≥1.0	29.2 ± 8.7	≤0.6		
	CN	16.8 ± 2.7	>50	>3.0	>50	>3.0	21.4 ± 9.8	1.3	30.1 ± 10.6	1.8*		

Values represent integrated best-fit IC₅₀ (24–96 h, μM) of OTA from non-linear regression ±95% confidence intervals based on asymptotic standard errors. Significant differences compared to control are indicated (**p* < 0.05). ≥, values are at least as high as control, but higher than the tested concentration range (i.e. >50 μM); >, values are higher than control values, but higher than the tested concentration range.

slightly more sensitive towards both ochratoxins than LLC-PK1 cells, when absolute IC₅₀ values for OTA and OTB were compared (Tables 2 and 3). In contrast, primary human cells (HKCm) displayed a lower sensitivity than the corresponding continual cell line IHKE, but also a similar cytotoxic response to both toxins. When the most sensitive endpoints are compared, no distinct species differences in cytotoxic effects of OTA and OTB were apparent.

Of the four cell types tested, the IHKE cell line seemed to be the most sensitive cell type to ochratoxin exposure.

OTA in cell subpopulations:

The effects of OTA exposure on the various cell subpopulations compared to their corresponding control cell cultures are shown in Table 2.

HKCm-subpopulations displayed a reduced sensitivity towards OTA-mediated cytotoxicity compared to their corresponding control. In contrast, none of the IHKE-subpopulations showed any significant alteration in sensitivity towards OTA cytotoxicity.

PKCm subpopulations (lowOTA, highOTA and high-OTB) also displayed a lower sensitivity to OTA exposure, whereas no altered effect could be observed in the PKCm lowOTB-subpopulation.

Both LLC-PK1 OTA-subpopulations also displayed a decreased sensitivity to OTA exposure, whereas the sensitivity of both OTB-subpopulations was unaltered.

OTB in cell subpopulations:

Table 3 shows the effects of OTB exposure on the various cell subpopulations in comparison with their corresponding control cell cultures.

All subpopulations of HKCm cells displayed a decreased sensitivity towards OTB cytotoxicity. Both IHKE OTA-subpopulations displayed a comparable decrease, whereas no significant effects were seen in either of the OTB-subpopulations.

All PKCm cell subpopulations showed an even more notable decrease in sensitivity (up to 6.3-fold) to OTB-challenge. LLC-PK1 cells generally showed similar trends, however, only few data were statistically significant.

Thus, although some contradictory observations were apparent, repeated exposure to OTA or OTB generally resulted in a decreased sensitivity towards subsequent OTA- and OTB-mediated cytotoxicity in human and porcine cells. This effect was most prominent in primary porcine cells.

4. Discussion

In order to generate ochratoxin-derived cell subpopulations as a model of repeated ochratoxin exposure, it was first necessary to induce renal cell regeneration processes due to cell loss. Long-term or extended *in vitro* studies in general are subject to the problem of limited life span of

primary cells. In order to overcome this problem, acutely toxic concentrations of OTA and OTB had to be used, although such high concentrations are unlikely to occur in the kidney *in vivo* under normal circumstances. Preliminary experiments indicated that the repeated exposure to either 20 μ M or 50 μ M OTA and OTB at approximately 90% confluency yields expandable cell subpopulations for subsequent analysis. This exposure regime resulted in five subpopulations for each of the four cell types, which could be used as a model in the presented study.

At first, these cell subpopulations were subjected to rudimentary characterization. The investigation of cell growth behavior demonstrated that the lowOTA-LLC-PK1 subpopulation and all IHKE cell subpopulations responded with higher cellular proliferation rates. All other cell subpopulations displayed slower or in some cases unaltered growth rates. The total cell number at confluence was decreased in most cell subpopulations, only IHKE cells did not show any alteration. These observations may be explained by the fact, that this cell line is originally a toxin-derived cell subpopulation (NiSO₄ clonogenicity) (Tveito et al., 1989), which probably represents a different control cell type compared to the other three cell types investigated. Therefore, caution is advised in the interpretation of results obtained using this cell type.

The investigated expression of marker proteins showed only effects in all four human primary cell subpopulations, where the cytokeratin expression seemed to be altered. All other cell subpopulations showed no changes. No increase in vimentin expression was detected in any cell type.

A comparison between the three different cytotoxicity endpoints used in this study demonstrated that not all endpoints provide similar results. The four investigated cell types showed a distinct species- and cell type-related endpoint-sensitivity. In both human cell types, the MTT reduction assay was the most sensitive endpoint, whereas total cell number seemed to be a slightly more sensitive endpoint in both porcine cell types. These results suggest that the endpoint for cytotoxicity testing in renal cells should be chosen according to their most sensitive behavior, which may differ between cell cultures of various sources.

Considering the data from the most sensitive endpoints, there is a clear decrease in cytotoxic response to OTA in all OTA-subpopulations and to OTB in all OTB-subpopulations, with the exception of IHKE cells, where sensitivity was unaltered.

No observable cytotoxicity difference between OTA and OTB was found in all control cell cultures and endpoints tested. At first, this appeared to stand in contrast to literature data including our own earlier findings (Dietrich et al., 2001; Heussner et al., 2006; O'Brien et al., 2001). The latter data showed a higher toxicity of OTA compared to that of OTB in various renal cell types, however these data always originated from single time-point short-term investigations. In the current study, an extended integrated time-interval of 96 h was investigated, obviously leading to a different outcome. More recent data from other investigators have

also described only minor cytotoxicity differences between OTA and OTB in LLC-PK1 cells when short-term effects are investigated, but no differences when higher concentrations or longer exposure times were employed (Mally et al., 2005).

Recently, Bruene and co-workers followed a similar repeated exposure approach of OTA (20 weeks) using HEK293 and NRK-52E cells. Here, also cell-type specific effects were detected. HEK293 cells displayed a transiently decreased sensitivity towards OTA combined with an increase in cell proliferation rates measured as an increased S-phase cell fraction. No such effects were detected in NRK-52E cells (B. Bruene, personal communication).

Thus it appears that repeated exposure to ochratoxins causes alterations in fundamental cellular characteristics and properties. If this is indeed associated with a real fibroblastic phenotype is unclear, however, the dedifferentiation of renal epithelial cells into fibroblast-like cells (and *vice versa*) has previously been described (Strutz et al., 1996). In rats, this transdifferentiation is accompanied by loss of cellular polarity, tight junctions, elongation, detachment and separation from neighboring cells (Ng et al., 1998). To date little is known about the mechanisms of transdifferentiation, however it seems often to be associated with changes in intermediary filaments, i.e. keratin expression is lost and vimentin appears, and collagen secretion is enhanced. Transdifferentiation has furthermore been discussed in relation to BEN (Markovic-Lipkovski, 2002). Indeed, Sauvant and colleagues hypothesized epithelial-to-mesenchymal transition because of OTA-induced enhancement of collagen secretion in OK cells and primary human proximal tubular epithelial cells (RPTECs) (Sauvant et al., 2005).

However, contrary to expectations based on the above publications, no overt morphological changes were seen in any of the cell subpopulations tested in this study. Moreover, no alterations in the expression of intermediary filaments (cytokeratins and vimentin) in PKCm, IHKE and LLC-PK1 cells, with the exception of lower cytokeratin expression in highOTA, lowOTB and highOTB HKCm subpopulations were observed. Whether or not the latter observations result from concentration effects or substance-specific mechanisms as previously suggested (O'Brien et al., 2001), cannot be derived from the current data.

In summary, the hypothesis of alteration of normal renal epithelial cells after repeated exposure to ochratoxins, detectable as alterations in parameters, e.g. morphology, growth rate and susceptibility to acutely cytotoxic concentrations of ochratoxins, was corroborated with the data presented in this study. These observations may be explained by the phenomenon of epithelial-to-mesenchymal transition (EMT), a selection for less ochratoxin-sensitive cells, or an adaptive behavior of the cells during repeated exposure.

Further investigations, including the introduction of model substances for cellular transformation (for example cyclosporine A, TGF- β 1, H₂O₂) (McMorrow et al., 2005;

Rhyu et al., 2005; Zhuang et al., 2005) and additional marker proteins (i.e. α -smooth muscle actin, cadherins, collagens, matrix metalloproteinases, etc.), recently shown to be involved in ochratoxin A-induced renal cortex fibrosis and EMT *in vivo* (Gagliano et al., 2005), are needed to elucidate the underlying mechanism.

The extrapolation of the *in vitro* data to a possible *in vivo* situation also suggests that ochratoxin-dependent increased cellular regeneration time may be an important factor contributing to the observed renal fibrosis in ochratoxin-exposed pigs and possibly also in humans. Indeed, slowed epithelial cell regeneration could explain earlier findings that ochratoxins can also induce invasion of the renal interstitia by fibroblasts as a measure of maintaining structural, but not functional, integrity of the kidney.

Whether the observed alterations are of stable and/or phenotypic or of temporary adaptive nature, should be investigated in the future to allow further insight into long-term ochratoxin effects.

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