

Biological similarities between murine chemical-induced and natural human bladder carcinogenesis

CARLOS PALMEIRA^{1,6}, PAULA A. OLIVEIRA⁸, CATARINA LAMEIRAS²,
TERESINA AMARO³, VICTOR M. SILVA⁴, CARLOS LOPES⁷ and LÚCIO SANTOS^{5,6}

Departments of ¹Immunology, ²Microbiology, ³Pathology, ⁴Urology, and ⁵Surgical Oncology, Portuguese Institute of Oncology, Porto; ⁶Health School, University Fernando Pessoa, Porto; ⁷Department of Pathology and Molecular Immunology, Instituto de Ciências Biomédicas de Abel Salazar, University of Porto, Porto; ⁸Department of Veterinary Sciences, CEVAC, University of Trás-os-Montes and Alto Douro, Vila Real, Portugal

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Abstract. The present study investigated the similarities between rodent and human urothelial carcinogenesis models using DNA content, p53 and Ki-67 immunoreexpression as surrogate markers of bladder carcinogenesis. Following N-butyl-N-(4-hydroxybutyl)-nitrosamine exposure, 49 human cystectomy specimens of bladder cancer and 53 rat bladder specimens were studied. All of the tumours and adjacent mucosa present in each specimen were evaluated. High similarities were observed between the rodent urothelium carcinogenesis process and the corresponding process in humans, in regards to the histopathological features and biological alteration profile: DNA aneuploidy, p53 overexpression and high proliferative index measured by Ki-67 immunoreexpression. Despite these similarities, a higher frequency of alterations was observed in earlier stages in the rat chemical-induced carcinogenesis, namely in 5c aneuploid cells, p53 overexpression and higher Ki-67 labelling index. These results confirm that this experimental animal model is a suitable and reproducible model of bladder carcinogenesis, particularly in regards to high-risk non-invasive and invasive urothelial carcinomas. These features mandate its use in the identification of new molecular targets and evaluation of tumour response to new cytotoxic drugs or drug combinations in bladder cancer therapeutic intervention.

Introduction

The importance of chemical carcinogens in the development of bladder cancer has been well established and generally

accepted. This carcinogenesis process occurs as a multistep process. Preneoplastic lesions, differently progressed, but with clonally related genomes, exist prior to *in situ* and invasive carcinoma, and may locally contribute to the independent formation of tumours (1).

Animal models of cancer have been fundamental for the demonstration of this multistep nature (2). Several rodent models have been established to study these different stages of urinary bladder chemical carcinogenesis, namely chemical initiation and promotion, and the stages of progression (3,4). The administration of N-butyl-N-(4-hydroxybutyl)-nitrosamine (BBN) to mice (5-7) or rats (7-11) is one of the most widely used chemical carcinogens.

The resultant induced tumours in these experimental models have been histologically well studied. However, little is known regarding the biological features and genetic similarities between rodent and human bladder cancer models.

This study aimed to identify biological similarities between human bladder carcinogenesis and rat chemical-induced bladder carcinogenesis by assessing DNA content alterations and p53 and Ki-67 immunoreexpression, in order to use this preclinical animal model in future therapeutic experimental studies.

Materials and methods

In this study, we compared biological data collected from our previous studies, in which human (12) and rat (10) bladder samples with premalignant and malignant lesions and adjacent mucosa were studied with regards to histopathology, DNA content, p53 immunoreexpression and the Ki-67 labelling index.

Human bladder samples. Consecutive radical cystectomy specimens from 49 patients with previous bladder cancer (non-invasive and invasive urothelial cell carcinomas), consecutively admitted and treated at the Instituto Português de Oncologia do Porto, Portugal, between 1989 and 1996, with available archival material for the image cytometric study and patient consent, were included in the present study. Only high-

Correspondence to: Dr Carlos Palmeira, Department of Immunology, Portuguese Institute of Oncology, and Health School of University Fernando Pessoa, Rua Dr António Bernardino de Almeida, 4200-072 Porto, Portugal
E-mail: palmeira.ca@gmail.com

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risk non-invasive urothelial cell carcinomas received prior intravesical Bacillus Calmette-Guérin treatment. Clinical data were collected from the patient records and the paraffin blocks from the Department of Pathology. All of the lesions present in each cystectomy specimen were studied, including the tumour area and the adjacent mucosa (12).

Rat bladder samples. Fifty-three female Fisher 344 rats were obtained at the age of 5 weeks from Harlan (Amsterdam, The Netherlands). Details concerning the animals, diet, carcinogens and experimental protocol were approved by the Portuguese Ethics Committee for Animal Experimentation (Direcção Geral de Veterinária, approval no. 520/000/000/2003) and are fully described in our previously published study (10). Briefly, animals were randomly separated in five groups. BBN was administered in drinking water at a concentration of 0.05% during a period of 20 weeks to all animal groups, with the exception of group 1, which served as the control and was not given any chemical supplement. After this BBN exposure, animals were maintained on normal tap water until the end of the experiment. Groups 2 and 3 were exposed only to BBN, and animals were sacrificed after an intraperitoneal administration of sodium pentobarbital 1 week (group 2) and 7 weeks (group 3) after BBN exposure. One week after BBN exposure, groups 4 and 5 began intravesical instillations with physiological saline solution (PSS) and mitomycin C (MMC), respectively, once a week for 6 weeks. After this treatment period, the animals were sacrificed. The tumours and adjacent mucosa were studied for each bladder specimen.

For intravesical immunotherapy, the dosing schedules were based on those commonly used in clinical work; 300 μ l of MMC solution (1 mg/1 ml) were used (10). The preparation procedures of the bladder for macroscopic examination and tissue processing were previously described (10).

Histopathological analysis. The diagnostic pathology slides were re-evaluated by experienced uropathologists (T.A. for the human series, and C.L. and P.O. for the rat specimens). Pathological staging was performed according to the American Joint Committee on Cancer Staging (13). The histological classification and grading were performed using the criteria from the 2004 World Health Organization (WHO) guidelines (14,15).

DNA content analysis. The DNA content quantification was performed using image cytometric analysis and a CAS 200 Image Analysis System (Cell Analysis Systems, Inc., Elmhurst, IL, USA). All methodological and DNA histogram analysis procedures were previously described in detail (6). Briefly, after the calibration of the image system using a control slide with rat hepatocytes with a known quantity of DNA, 20-30 lymphocytes and a minimum of 100 intact non-overlapping urothelial nuclei were measured and analysed for each case. The G0/G1 peak was visually identified in each DNA histogram, and the mean, standard deviation (SD) and coefficient of variation (CV) values were calculated. The DNA index (DI) describes the relative DNA content of the study population and was defined as the ratio of mean DNA content of the urothelial G0/G1 peak divided by the mean DNA content of the resting diploid lymphocyte G0/G1 peak. The 5cER was also evaluated and defined as the percentage of

cells with values $>5n$. Lesions were considered aneuploid only if a separate G0/G1 peak was distinguishable on the histogram and it differed from the reference lymphocyte population by >2 SD. A DNA diploid lesion showed a single distinct G0/G1 peak with a DI within 2 SD of the control lymphocytes and usually with $<1\%$ 5cER.

Immunohistochemical analysis

Human specimens. For the immunohistochemical analysis a standard avidin-biotin peroxidase method was used, as described in a previous study (16). The immunoexpression of p53 and Ki-67 was evaluated with the primary antibodies D07 (1:50 dilution; Dako®) and MIB1 (1:50 dilution; Novocastra®), respectively.

Paraffin-embedded tissues, known to express nuclear p53 (colon carcinoma) and have a high proliferation (lymphoma), were used for titration and positive controls. Negative controls were performed replacing the primary antibody with 2.5% bovine serum albumin in PBS.

Rat specimens. A three-step streptavidin-biotin immunoperoxidase method was used, as described in a previous study (10). The primary antibodies AB-1 (1:50 dilution; Neomarkers-Labvision) for p53 expression and Ki-67 (1:20 dilution; Dako) for proliferative activity, were used. Paraffin sections from colon and breast cancer with known immunoreactivity to p53 and Ki-67 antigens, respectively, were used as positive controls. Negative controls were carried out by replacing the primary antibodies with PBS.

Slides were re-evaluated by two independent observers (T.A. and L.S. for the human cases and C.L. and P.O. for the rat cases) in a blinded fashion, without knowledge of the clinical data. The evaluation method was also described previously (16). The entire lesion was screened to find the region with the maximum fraction of positive- and contiguous-stained nuclei for p53 and the region with the maximum fraction of positivity for Ki-67. The percentage of positive-stained nuclei was scored in this region using a $\times 40$ objective. Whenever possible, at least 100 cells were scored in each histological lesion. In the human series, the cut-off values used to distinguish positivity for p53 and Ki-67 were >18 and $>32\%$, respectively, based on studies reported previously (16). In the rat samples, the p53 and Ki-67 immunoexpression was calculated as the percentage of positive nuclei divided by the total number of cells examined (10).

Statistical analysis. The statistical analysis was carried out using the SPSS 15.0 statistical package for Windows (SPSS Inc.). A descriptive analysis was performed for the variables studied. The incidences of differences in molecular alterations between the histological groups were evaluated by the Pearson's Chi-square and the Fisher's exact (when $n < 5$) tests. The mean percentage of positivity of p53 and Ki-67 of the histological types was compared by analysis of variance (ANOVA). $P < 0.05$ was defined as statistically significant.

Results

Evaluation of human carcinogenesis. Of the 49 patients included in the present study, 19 cases (38.8%) were classified as high-grade papillary urothelial carcinoma (HGP) and 30

Table I. Frequency of molecular alterations for each histological type observed in the human series.

Variables	Normal n=27 n (%)	HYP n=9 n (%)	DYS n=2 n (%)	CIS n=11 n (%)	HGP n=19 n (%)	INV n=30 n (%)	p-value
DNA ploidy							<0.001
Diploid	26 (96.3)	7 (77.8)	1 (50)	1 (9.1)	7 (36.8)	2 (6.7)	
Aneuploid	1 (3.7)	2 (22.2)	1 (50)	10 (90.9)	12 (63.2)	28 (93.3)	
5cER							<0.001
Without	26 (96.3)	7 (77.8)	1 (50)	1 (9.1)	9 (47.4)	2 (6.7)	
With	1 (3.7)	2 (22.2)	1 (50)	10 (90.9)	10 (52.6)	28 (93.3)	
p53							0.001
Negative	26 (96.3)	8 (88.9)	1 (50)	7 (63.6)	14 (73.7)	15 (50.0)	
Positive	1 (3.7)	1 (11.1)	1 (50)	4 (36.4)	5 (26.3)	15 (50.0)	
Ki-67							<0.001
Negative	26 (96.3)	9 (100)	2 (100)	4 (36.4)	7 (36.8)	8 (26.7)	
Positive	1 (3.7)	0 (0)	0 (0)	7 (63.6)	12 (63.2)	22 (73.3)	

Normal, morphological normal urothelium; HYP, hyperplasia; DYS, dysplasia; CIS, carcinoma *in situ*; HGP, high-grade papillary urothelial carcinoma; INV, invasive urothelial carcinoma.

Table II. Incidence of urothelial histopathological lesions in rats exposed to N-butyl-N-(4-hydroxybutyl)-nitrosamine and treated with Mitomycin C and physiological saline solution.

Group (n)	Normal n=10 n (%)	HYP n=6 n (%)	DYS n=41 n (%)	CIS n=7 n (%)	LGP n=40 n (%)	HGP n=30 n (%)	INV n=10 n (%)
1 (10)	10 (100)	0 (0.0)	0 (0.0)	0 (0)	0 (0.0)	0 (0.0)	0 (0.0)
2 (12)	0 (0)	2 (16.6)	10 (87.3)	0 (0)	10 (83.3)	2 (16.6)	2 (16.6)
3 (10)	0 (0)	2 (20.0)	10 (100.0)	0 (0)	10 (100.0)	9 (90.0)	1 (10.0)
4 (11)	0 (0)	1 (9.0)	11 (100.0)	0 (0)	11 (100.0)	11 (100.0)	5 (45.4)
5 (10)	0 (0)	1 (10.0)	10 (100.0)	7 (70)	9 (90.0)	8 (80.0)	2 (20.0)

Normal, morphological normal urothelium; HYP, hyperplasia; DYS, dysplasia; CIS, carcinoma *in situ*; LGP, low-grade papillary urothelial carcinoma; HGP, high-grade papillary urothelial carcinoma; INV, invasive urothelial carcinoma.

(61.2%) as invasive urothelial carcinoma (INV). For all of the tumours the adjacent mucosa was evaluated and showed differential histological patterns dependent on the adjacent tumour. Morphological normal urothelium (n=13, 68.5%), hyperplasia (n=3, 15.8%) and carcinoma *in situ* (CIS) (n=3, 15.7%) lesions were found in the HGP adjacent mucosa, while the INV adjacent mucosa showed morphological normal urothelium (n=14, 46.7%), hyperplasia (n=6, 19.9%), dysplasia (n=2, 6.7%) and CIS (n=8, 26.7%) lesions.

Each one of these histological patterns found in the adjacent mucosa showed the same molecular profile, in regards to the biological variables studied, independently of the adjacent tumour.

Considering each histological type, we compared the distribution of the molecular alterations for the biological variables studied. Statistical significant differences were

observed according to the potential of aggressiveness of the lesion (Table I).

Evaluation of rat carcinogenesis. The histopathological lesions observed in each experimental group of rats are described in Table II. No histopathological alterations in urothelial cells were observed in the control group (group 1). The frequency of lesions increased with a longer observation period after BBN exposure. A higher number of HGP and INV cases was observed in the animal group exposed to BBN and submitted to the intravesical instillation of physiological saline solution (group 4). CIS lesions were detected only in animals in group 5, exposed to BBN and treated with MMC.

The molecular alterations found in each histological type are presented in Table III. Despite the period of observa-

Table III. Frequency of molecular alterations for each histological type observed in the rat series.

Variables	Normal n (%)	HYP n (%)	DYS n (%)	CIS n (%)	LGP n (%)	HGP n (%)	INV n (%)	p-value
DNA ploidy								<0.001
Diploid	10 (100.0)	4 (80.0)	9 (56.3)	0 (0.0)	17 (70.8)	0 (0.0)	0 (0.0)	
Aneuploid	0 (0.0)	1 (20.0)	7 (43.8)	7 (100.0)	7 (29.2)	23 (100.0)	3 (100.0)	
5cER								<0.001
Without	8 (80.0)	4 (80.0)	9 (56.3)	0 (0.0)	19 (79.2)	2 (8.7)	0 (0.0)	
With	2 (20.0)	1 (20.0)	7 (43.8)	7 (100.0)	5 (20.8)	21 (91.3)	3 (100.0)	
p53 (positivity) mean value	10 (0.0)	6 (20.7)	17 (35.5)	7 (37.4)	34 (36.0)	26 (44.0)	4 (38.9)	<0.001
Ki-67 (positivity) mean value	10 (0.0)	6 (16.8)	17 (23.5)	7 (29.3)	34 (23.3)	26 (34.0)	4 (23.9)	<0.001

Normal, morphological normal urothelium; HYP, hyperplasia; DYS, dysplasia; CIS, carcinoma *in situ*; LGP, low-grade papillary urothelial carcinoma; HGP, high-grade papillary urothelial carcinoma; INV, invasive urothelial carcinoma.

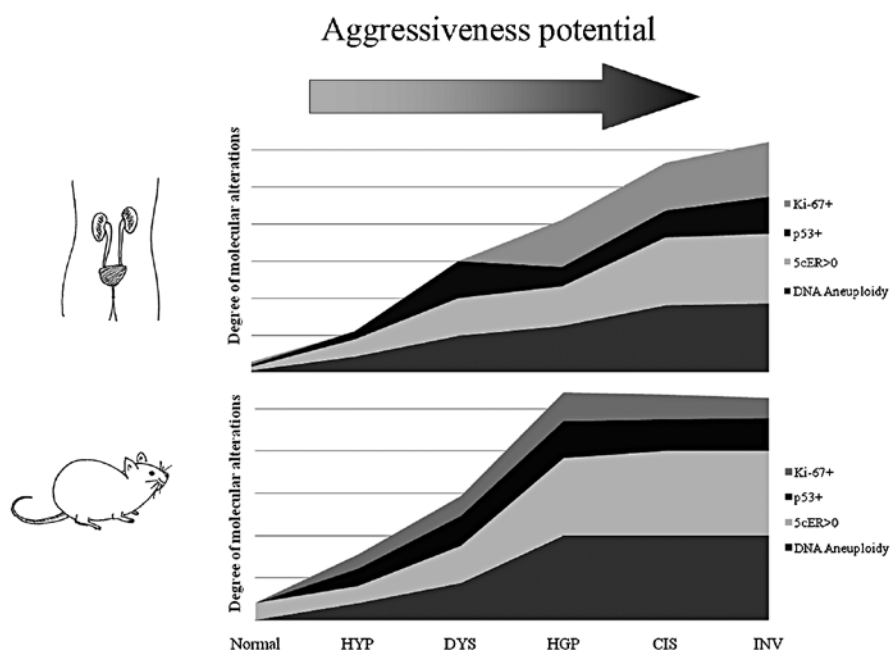


Figure 1. Frequency of molecular alterations for each histological type observed in the human and rodent models. Normal, morphological normal urothelium; HYP, hyperplasia; DYS, dysplasia; HGP, high-grade papillary urothelial carcinoma; CIS, carcinoma *in situ*; INV, invasive urothelial carcinoma.

tion and type of intravesical instillation, the frequency of DNA aneuploidy, 5c aneuploid cells, p53 overexpression and abnormal Ki-67 immunoexpression increased with the degree of aggressiveness of the urothelial lesion.

The comparison between the two cancer models, in regards to the molecular alterations found in each histological pattern, is illustrated in Fig. 1. High similarities were observed between the rodent urothelium carcinogenesis process and the corresponding process in humans, in regards to histopathological features and the profile of biological alterations: DNA aneuploidy, p53 overexpression and high proliferative index measured by Ki-67 immunoexpression. Despite these similarities, a higher frequency of alterations was observed in

earlier stages in rat chemical-induced carcinogenesis, namely in 5c aneuploid cells, p53 overexpression and higher Ki-67 labelling index.

Discussion

Cancer involves abnormal cellular growth and appears to have a common molecular basis. Comparing the genetic signatures of human and murine bladder carcinogenesis, William and colleagues (7) found that cell cycle-related genes are involved in the two processes, supporting our options to select DNA content, p53 alterations and Ki-67 labelling index as surrogate markers of bladder carcinogenesis.

The present study found compelling similarities between the rodent urothelium carcinogenesis process and the corresponding process in humans, in regards to histopathological features and the profile of biological alterations: DNA aneuploidy, p53 overexpression and high proliferative index measured by Ki-67 immunoeexpression.

The histopathological alterations observed by our group in rat chemical-induced bladder carcinogenesis, namely hyperplasia, dysplasia, low- and high-grade papillary urothelial cell carcinoma, CIS and INV were also observed in natural human bladder carcinogenesis (14,15).

The biological history of bladder cancer shows that DNA aneuploidy is a reflection of genomic instability and chromosomal derangement, and a high aneuploidy level was observed in human advanced bladder cancer (17). *TP53* genetic alterations are associated with tumour stage and grade, and there is a significant association with patient outcome (16,18,19). A high proliferative index is now accepted as a prognostic factor in human bladder cancer improving identification of patients who are at increased risk for disease progression after radical cystectomy (20).

In the experimental orthotopic model described herein, we observed the entire spectrum of lesions that is described in the dual-track pathway of human bladder carcinogenesis: the papillary and non-papillary pathways (21). We also observed that the variation in the frequency of genetic alterations was similar between the rodent (rat) and human cancer models. Thus, the rate of DNA aneuploidy, p53 immunoeexpression and Ki-67 labelling index was higher in more aggressive lesions. However, a biological profile similar to that in human invasive tumours was observed in early tumour stages in rats suggesting that this murine model is a beneficial model with which to study the invasive carcinogenesis pathway. William and colleagues also observed that induced murine tumours exhibited more similarities in gene expression to human muscle invasive tumours than non-invasive tumours. Thus, rodent (rat) tumours provide an accurate mechanistic study of genes putatively involved in invasive and metastatic bladder cancer (7).

We also observed a high frequency of biological alterations in high-risk non-muscle invasive tumours, similar to those that occur in human counterparts. This result indicates that this model is also suitable for the study of these particular tumours.

The present and previous results obtained by our group (10,11) emphasize that we now have a technically suitable and highly reproducible model of bladder urothelial carcinogenesis. This model resembles human disease both histologically and in biological behaviour, particularly for high-risk non-muscle invasive and invasive urothelial bladder cancer. Therefore, it is a reliable model with which to identify new molecular targets and evaluate the feasibility and tumour response of new cytotoxic drugs or drug combinations in the field of urologic oncology.

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