Cyclooxygenase-2 and p53 Expression as Prognostic Indicators in Conventional Renal Cell Carcinoma

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The aim of this study was to investigate the relationship of cyclooxygenase (COX)-2 and p53 expression with prognosis in patients with conventional renal cell carcinoma (RCC). Formalin-fixed, paraffin-embedded tissue sections of conventional RCC from 92 patients, who had undergone radical nephrectomy, were examined for COX-2 and p53 expression by immunohistochemistry and compared with clinicopathological variables. The COX-2 expression significantly correlated only with tumor size (p=0.049), whereas the p53 expression profoundly correlated with the TNM stage (p=0.024), M stage (p=0.001), and metastasis (synchronous or metachronous; p=0.004). The COX-2 overexpression did not significantly associate with p53 positivity (p=0.821). The survival rate of patients correlated with the p53 expression (p < 0.0001) but not with the COX-2 expression (p=0.7506). Multivariate analyses indicated that tumor size, M stage, and p53 expression were independent prognostic factors for cancer-specific survival. The COX-2 expression was not an independent factor. These results show that the increased expression of p53 was associated with metastasis and a worse prognosis in conventional RCC, which suggests that p53 might have played an important role in the progression of conventional RCC. The increased expression of COX-2 was associated only with tumor size, but may not be an important prognostic factor in conventional RCC. No association was observed between COX-2 overexpression and p53 positivity in conventional RCC.

Key Words: Cyclooxygenase-2, p53, prognosis, renal cell carcinoma

INTRODUCTION

The renal cell carcinoma (RCC) is the most com-

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mon malignant tumor of the adult kidney, and its incidence is increasing. To date, tumor stage and grade have been considered the most important prognostic parameters for patients with RCC. However, in many cases, these parameters were insufficient to predict the clinical behavior of RCC tumors. Therefore, it is important to identify additional indicators of the biological aggressiveness of RCC.

Cyclooxygenase (COX) catalyzes the synthesis of prostaglandins from arachidonic acid. There are two isoforms of COX: COX-1 and COX-2. While COX-1 is constitutively expressed, COX-2 is induced by various stimuli such as the cytokines, growth factors, and tumor promoters.^{4,5} The COX-2 levels have been shown to increase in several types of human cancers like the ones which develop in the gastrointestinal tract, breast, cervix, lung, prostate, and bladder. 6-12 This suggests that the COX-2 may play an important role in the cancer progression by affecting tumor cell proliferation, resistance to apoptosis, angiogenesis, immunosuppression, and tumor invasion.7,13-15 However, the relationship between the COX-2 expression and RCC remains unclear. Although the COX-2 is highly expressed in canine RCC, ¹⁶ it is down-regulated in RCC in the Eker (TSC2 gene mutant) rat model.¹⁷ The COX-2 is highly expressed in human RCC, but the results of its correlation with tumor stage and grade have been contradictory. 18,19

Apoptosis is a critical pathway that selectively allows certain cells with damaged DNA to undergo cell death. Abnormalities in this pathway may lead to uncontrolled cellular proliferation and, ultimately, to carcinogenesis. p53 is a known

inducer of apoptosis, and it is the most common genetic mutant found in cancers. There are wide variations in the reported incidence of p53 related mutations in RCC, and its prognostic significance for this type of cancer remains controversial. Some researchers have reported that p53 mutations in the case of RCC may be used as a prognostic factor, but others disagreed. Recent evidences have suggested that COX-2 may also inhibit apoptosis. Some

In the present study, the expressions of COX-2 and p53 was investigated, and their relationships to clinicopathological variables in patients with conventional RCC were evaluated.

MATERIALS AND METHODS

Patients and specimens

Formalin-fixed, paraffin-embedded, archival surgical specimens that had been obtained from 92 patients (71 men and 21 women; mean age, 54.8 years; range, 26 - 81 years) who had received a diagnosis of primary conventional RCC were studied. All patients underwent radical nephrectomy at our institution between January 1995 and September 2002. During the lymphadenectomy, only the renal hilar lymph nodes were routinely removed. Whenever an enlarged lymph node was observed during surgery, a node dissection clearing the ipsilateral great vessel and the renal hilum was performed. Tumors were staged according to the 1997 revised TNM system, 24 and were graded according to Fuhrman's nuclear grading system. 25

Immunohistochemistry

Paraffin-embedded blocks were sectioned at 4µm thickness, deparaffinized, and rehydrated.
After microwave pretreatment in citrate buffer
(pH 6.0) for antigen retrieval, slides were immersed in 3% hydrogen peroxide for 10 minutes
to block the endogenous peroxidase activity. The
sections were blocked for 10 minutes with blocking reagent using the Cap-PlusTM Detection Kit
(ZYMED, San Francisco, CA, USA). Sections were
then incubated manually with goat monoclonal
antibody to COX-2 (diluted 1:200; Santa Cruz

Biotechnology, Santa Cruz, CA, USA) for 1 hour at 37°C or with mouse monoclonal antibody to p53 (diluted 1:20; Novo Castra, Newcastle, UK) overnight at 4°C in a humidified chamber. After a second incubation with a biotinylated secondary antibody, slides were incubated with peroxidaseconjugated streptavidin (DAKO LSAB+ Kit; DAKO Corp., Carpinteria, CA, USA) for COX-2 or the Cap-PlusTM Detection Kit (ZYMED) for p53. Reaction products were visualized by immersing the slides in diaminobenzidine tetrachloride and then counterstaining them with Mayer's hematoxylin. The positive staining of smooth muscle cells from blood vessels provided an internal positive control for COX-2 immunostaining. The colon carcinoma samples with known p53 mutations and documented accumulation of p53 by immunohistochemistry were used as positive control for p53. Negative controls were obtained by omitting the primary antibody.

Evaluation of immunohistochemical staining

The immunostaining was independently evaluated by two pathologists who were unaware of the clinical data. The COX-2 expression was based on the presence of the cytoplasmic staining, which was semi-quantitatively estimated according to the methods described by Sinicrope et al.,26 with minor modifications. On the basis of the percentages of immunopositive cells, the data were subdivided into five categories as follows: (0) \leq 10%; (1) 11 - 25%; (2) 26 - 50%; (3) 51 - 75%; and (4) > 75% positive cells. The immunointensity was also subclassified into four categories: (0) negative; (1) weak; (2) moderate (same intensity of smooth muscle cells); and (3) strong (Fig. 1). The immunoreactive scores for each case were generated by multiplying the values of the two parameters, which were then stratified into three groups: weak (scores 0-4), moderate (scores 5-8), and strong (scores 9-12) COX-2 expression for the survival analysis. The p53 immunostaining was considered to be positive when 10% or more of the tumor cell nuclei were stained (Fig. 2).

Statistical analysis

Either an independent-sample t-test or a chi-

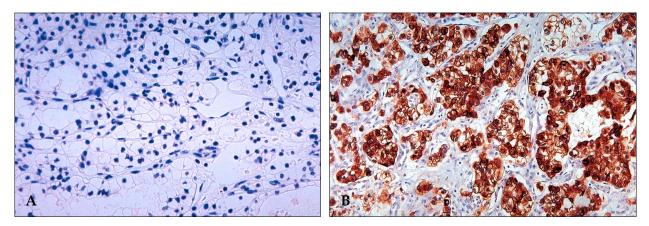


Fig. 1. Immunohistochemical staining for COX-2. (A) COX-2 immunostaining was not seen (COX-2 intensity, 0). (B) Almost all cancer cell cytoplasms strongly stained for COX-2 (COX-2 intensity, 3). Original magnification, × 400.

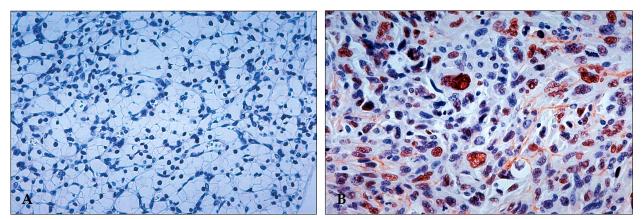


Fig. 2. Immunohistochemical staining for p53. (A) Tumor cells showed no nuclear staining for p53 (p53 negative). (B) A strong nuclear expression of p53 was seen (p53 positive). Original magnification, × 400.

square test was used to analyze the correlation between COX-2 or p53 expression and various clinicopathological variables. The relationship between COX-2 and p53 expression was evaluated by the Mann-Whitney U-test. The paired t-test or McNemar test was used for the correlation of the primary and metastatic sites in the expression of the COX-2 or p53. The survival calculations were illustrated with Kaplan-Meier curves, and univariate and multivariate analyses were performed using the log-rank test or the Cox proportional-hazards regression model. The values of p<0.05 were considered to be statistically significant in all of the analyses.

RESULTS

The clinicopathological characteristics of the 92 patients are summarized in Table 1. Sixty-seven patients were disease-free at a median follow-up of 36 months (range, 12 - 96 months); the other 25 patients had either synchronous metastases (n=12) at the time of nephrectomy or metachronous metastases (n=13) a median of 23 months (range, 4 - 72 months) after nephrectomy. Fourteen patients died during the follow-up period, 7 with synchronous metastases and 7 with metachronous metastases. For 18 of the 25 patients with metastases, their metastases were confirmed histopathologically.

Of the 92 sections, the COX-2 expression (im-

Table 1. Clinicopathological Data of 92 Patients with Conventional Renal Cell Carcinoma

Characteristics	No. (%)
TNM stage	
I	43 (46.7)
II	11 (12.0)
III	25 (27.2)
IV	13 (14.1)
T stage	
T1	48 (52.1)
T2	11 (12.0)
Т3	31 (33.7)
T4	2 (2.2)
N stage	
N0	91 (98.9)
N1	1 (1.1)
M stage	
M0	80 (87.0)
M1	12 (13.0)
Grade	
1	5 (5.4)
2	14 (15.2)
3	61 (66.3)
4	12 (13.0)
Metastasis (synchrono	us or metachronous)
No	67 (72.8)
Yes	25 (27.2)

munoreactive scores) was weak in 17 (18.5%), moderate in 21 (22.8%), and strong in 54 (58.7%). Positive immunostaining for p53 was observed in 11 (12.0%) of the 92 sections.

Because only 1 patient had lymph node metastasis, the analysis according to the lymph node status was not performed. The COX-2 expression significantly correlated with the tumor size (p= 0.049) but not with the TNM stage (p=0.541), T stage (p=0.368), M stage (p=0.394), grade (p=0.658), or metastasis (synchronous or metachronous; p= 0.564). The p53 expression profoundly correlated with the TNM stage (p=0.024), M stage (p=0.001), and metastasis (p=0.004) but not with the tumor

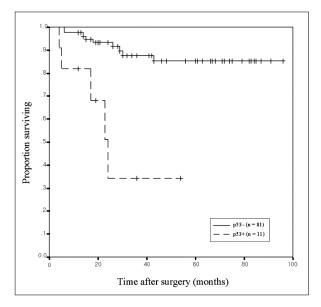


Fig. 3. Kaplan-Meier cancer-specific survival curves according to the p53 expression. The survival rate of patients with p53-positive tumors was significantly lower than that of patients with p53-negative tumors (p < 0.0001).

size (p=0.777), T stage (p=0.480), or grade (p=0.829) (Table 2).

The COX-2 overexpression did not significantly associate with p53 positivity (p=0.821). The expressions of COX-2 (p=0.476) and p53 (p=0.125) were not significantly different between the primary and metastatic sites. A Kaplan-Meier survival curve showed that the survival rate of patients with p53-positive tumors was significantly lower than that of patients with p53-negative tumors (p<0.0001) (Fig. 3). There was no difference in the survival rates according to the level of the COX-2 expression (p=0.7506) (Fig. 4).

The univariate analyses identified tumor size, TNM stage, T stage, M stage, and p53 expression as significant prognostic factors for cancer-specific survival, whereas the multivariate analyses indicated that tumor size (p=0.002), M stage (p<0.001), and p53 expression (p=0.016) were independent prognostic factors. The COX-2 expression was not an independent factor (Table 3).

DISCUSSION

An elevated COX-2 expression has been ob-

Table 2. Relationship between COX-2 or p53 Expression and Clinicopathological Variables in 92 Patients with Conventional Renal Cell Carcinoma

Variables	No. of patients	COX-2 immunoreactive scores	р	No. (%) of p53-positive tumors	p
Tumor size					
$\leq 7 \text{ cm}$	62	2.29 ± 0.80	0.040*	7 (11.3)	0.777
> 7 cm	30	2.63 ± 0.72	0.049*	4 (13.3)	
TNM stage					
Low (I + II)	54	2.44 ± 0.77	0.544	3 (5.6)	0.024*
High (III + IV)	38	2.34 ± 0.81	0.541	8 (21.1)	
T stage					
Low (T1 + T2)	59	2.46 ± 0.75	0.040	6 (10.2)	0.480
High (T3 + T4)	33	2.30 ± 0.85	0.368	5 (15.2)	
M stage					
M0	80	2.38 ± 0.80	0.204	6 (7.5)	0.001*
M1	12	2.58 ± 0.67	0.394	5 (41.7)	
Grade					
Low (G1 + G2)	19	2.47 ± 0.84	0.450	2 (10.5)	0.829
High (G3 + G4)	73	2.38 ± 0.78	0.658	9 (12.3)	
Metastasis					
No	67	2.37 ± 0.79	0.544	4 (6.0)	0.004:
Yes	25	2.48 ± 0.77	0.564	7 (28.0)	0.004*

Metastasis, synchronous or metachronous metastasis.

Table 3. Univariate and Multivariate Survival Analysis of 92 Patients with Conventional Renal Cell Carcinoma

77 - 11	Univariate	Multivariate		
Variables	p	Hazards ratio	p	
Tumor size	0.0003	13.510	0.002	
TNM Stage	0.0008	1.267	0.863	
T stage	0.0146	0.536	0.590	
M stage	<0.0001	18.267	<0.001	
Grade	0.1237	2.365	0.429	
COX-2	0.7506	0.778	0.573	
p53	<0.0001	5.432	0.016	

^{*}statistically significant

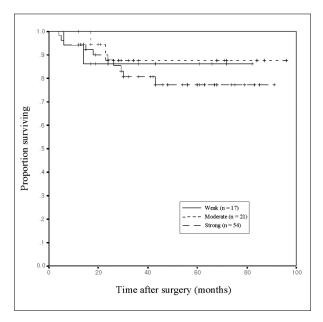


Fig. 4. Kaplan-Meier cancer-specific survival curves according to the COX-2 expression. There was no difference in survival rates according to the level of the COX-2 expression (*p*=0.7506). Weak, scores 0-4; moderate, scores 5-8; strong, scores 9-12.

served in several types of human cancers, and the overexpression of the COX-2 has been shown to correlate with carcinogenesis and poor prognostic outcome. However, past studies in RCC have been limited, and the relationship between the COX-2 expression and RCC remains unclear.

Khan et al.¹⁶ showed that COX-2 was overexpressed in 2 of 3 dogs with spontaneously developed RCC, which suggested that COX-2 might play a role in the modulation of neoplastic cell growth. Conversely, Okamoto et al.¹⁷ found that COX-2 mRNA expression was down-regulated in 8 of 10 spontaneously developed RCCs of Eker (TSC2 gene mutant) rats and that it was not down-regulated, but showed almost the same intensity as in the normal kidney in the other two rats.

With regard to the human RCC, Hara et al.²⁷ showed that COX-2 was overexpressed in RCC but that the expression levels did not correlate with cancer grade. In RCC, the immunoreactive signals in granular cell subtypes were stronger than those in clear cell subtypes. Miyata et al.¹⁸ found that increased COX-2 expression was observed in 70 (53.4%) of 131 patients with RCC.

Among the patients, 84% had conventional RCC. The COX-2 expression significantly associated with various clinicopathological variables, including the T stage, N stage, M stage, and tumor grade. It was correlated with the Ki-67 labeling index, microvessel density, and MMP-2 expression but not with the apoptotic index. The COX-2 expression was identified as an independent risk factor for large tumor size (> 7 cm) but not an independent prognostic factor for cancer-specific survival, which suggested that the COX-2 is not a significant prognostic factor in RCC.

In the present study, only the conventional RCC was studied in order to exclude the influence of the histologic type. The COX-2 expression was moderate or strong in 75 (81.5%) of the 92 RCC samples. The COX-2 expression significantly correlated with the tumor size but not with the TNM stage, T stage, M stage, grade, or metastasis (synchronous or metachronous). There was no difference in the survival rates according to the level of the COX-2 expression. The COX-2 expression was not an independent prognostic factor for cancer-specific survival according to our multivariate analysis. The results show that COX-2 is not an important prognostic factor in conventional RCC. These results are in agreement with those of Yoshimura et al.¹⁹ who reported that COX-2 overexpression was detected in RCC tissues but was not associated with tumor stage or grade. The findings of the present study are also in keeping with the results of Miyata et al. 18 that the COX-2 expression was associated with tumor size but was not an independent prognostic factor for cancer-specific survival.

Recent evidences have suggested that COX-2 contributes to the inhibition of apoptosis.¹³ The possible mechanism of the COX-2 inhibition of apoptosis involves the generation of prostaglandin products or the removal of the substrate arachidonic acid via COX-2 catalytic activity.^{28,29}

p53 is a known inducer of apoptosis, and its prognostic significance for RCC remains controversial. In the current study, the p53 expression significantly correlated with the TNM stage, M stage, and metastasis (synchronous or metachronous). The survival rate of patients with p53-positive tumors was significantly lower than that of patients with p53-negative tumors. The p53

expression, together with tumor size and M stage, was an independent prognostic factor for cancerspecific survival according to the results of the multivariate analysis conducted in this study. However, p53 positivity did not significantly associate with the COX-2 overexpression, and that was in harmony with the results of Miyata et al. ¹⁸ that the COX-2 expression did not correlate with the apoptotic index.

In conclusion, the increased expression of p53 was associated with metastasis and a worse prognosis in the conventional RCC, which suggested that p53 might have played an important role in the progression of the conventional RCC. However, the increased expression of the COX-2 was only associated with tumor size, but may not be an important prognostic factor in the conventional RCC. There was no association between the COX-2 overexpression and p53 positivity in the conventional RCC.

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