Investigation of tumour suppressor protein p53 in renal cell carcinoma patients

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Background. Investigation of p53 immunoreactivity in formalin-fixed paraffin-embedded tissues of normal renal tissue and renal cell carcinoma with respect to histopathologic subtype and nuclear grade of RCC.

Methods. 42 tissue sections of RCC and 5 samples of normal renal tissue were stained for p53 expression using immunohistochemical assay. The results were analyzed in relation to nuclear grade and histopathologic subtype.

Results. In total, p53 expression was found to be 4 to 5 times higher (30.8%) in other types of RCC than in the clear-cell type of RCC (6.9%). Further, there was no statistically significant difference in p53 overexpression among the histopathologic subtypes (P > 0.05, P = 0.063). No association was found between the expression of p53 and nuclear grade (P > 0.05, P = 0.17). Interestingly, our study also showed weak cytoplasmic positivity in renal tubular epithelium.

Conclusion. Our findings suggest that p53 might play an important role in tumour development or progression and it might be used as a new predictor and therapeutic target for RCC.

Key words: renal cell carcinoma, expression of p53, immunohistochemistry

INTRODUCTION

Renal cell carcinoma (RCC) accounts for approximately 85% of all kidney cancers. There are at least five subtypes of RCC currently recognized and the most common form is clear cell cancer, which accounts for roughly 70% to 80% of cases. This tumour is characterized by exceptionally high resistance to radiation and chemotherapy, which can be explained in part by naturally high levels of expression of multidrug transporters and elevated activity of the glutathione system in RCC progenitors, kidney tubular epithelial cells. However, treatment resistance of tumour cells can also be associated with suppression of apoptotic mechanisms of cellular response to stress, such as the p53 tumour suppressor pathway.

The tumour suppressor gene p53 is located on chromosome 17p13.1 that encodes a nuclear phosphoprotein of 53kDa (ref.3). It has been implicated in controlling a checkpoint during the G1 phase of the cell cycle, inducing cell cycle arrest and apoptosis in the presence of damaged DNA (ref.4). Wild-type p53 is usually not detectable by routine immunohistochemistry because of its short half-life. In contrast, the mutant p53 has a longer half-life, accumulates in the tissue, and can be easily detected in the cell nucleus. P53 appears to be mutated in about 50% of many human cancers, while in RCC, there is a low incidence of p53 mutations. P53 mutation has been reported in 3-33% of patients with RCC (ref.7). Although p53 wild type remains wild type in the majority of RCCs, this does not mean, as we know from other tumour types, that it is functional. P53 function can be repressed by some other mechanisms, which may involve overexpression of natural negative regulators of p53, such as MDM2 or MDMX (ref.8), or loss of positive regulators, such as Arf (ref.9), or by viral proteins, such as E6 of the human papilloma virus10. There have been a number of controversial studies on the association of p53 expression and nuclear grade. While some investigators have found no association, a strong relationship has been demonstrated by others and it is regarded as a potential marker in determining the prognosis of patients with RCC (ref.12).

In our study, we investigated p53 immunoreactivity in formalin-fixed paraffin-embedded tissues of normal renal tissue and renal cell carcinoma. To detect this protein, we used monoclonal antibody. Next, we conducted this study to evaluate the relationship between the overexpression of p53 tumour suppressor protein and histopathologic subtype of RCC. We also compared p53 immunoreactivity with the nuclear grade of our set of RCC as there is a dearth of published data on this connection.

MATERIALS AND METHODS

Patients
In the study, we used 42 samples of renal cell carcinoma and 5 samples of normal renal tissue. The samples were obtained from the Department of Pathology, Pasteur Faculty Hospital, Košice, Slovak Republic.
The renal cell carcinoma samples were divided according to histopathological type into 2 groups: 1\textsuperscript{st} group – conventional type of RCC (clear cell type) = 29 samples and 2\textsuperscript{nd} group – other types of RCC (5 papillary, 3 chromophobe, 1 sarcomatoid, 1 multilocular cystic and 3 unclassified types) = 13 samples. Patients and tumour characteristics are summarized in Table 1. All samples were immunohistochemically analyzed for wild and mutant type p53.

**Antibody**

We used the following primary monoclonal antibody: mouse anti-p53, clone DO7 (BioGenex Laboratories, Inc.), which recognizes an epitope residing between amino acids 32 to 79. This antibody is known to detect wild-type and mutated p53 in paraffin-embedded histological tissue and it is the most used antibody for detecting p53 in many diagnostic histopathological laboratories.

**Immunohistochemical detection of p53:**

After paraffin removal, sections were finally washed in phosphate-buffered saline containing 0.05% Tween-20 (PBS-Tw), pH 7.6. Endogenous peroxidase activity was blocked by 0.3% H\textsubscript{2}O\textsubscript{2} in methanol for 30 min at room temperature. The slides were pre-treated in a water bath in sodium citrate buffer 0.1 M (pH 6.0) in the microwave 2\times 5 min for antigen retrieval. The p53 staining procedure continued by blocking nonspecific staining with milk buffer (5% dry milk in TRIS buffer) for 30 min at room temperature. The next step was primary antibody which was applied overnight in a humidified chamber at 4˚C. After rinsing in PBS-Tw (3\times 5 min) the sections were subsequently incubated with the secondary antibody: prediluted biotinylated horse antibody (Vector Laboratories, USA) for 30 min at room temperature. The sections were visualized with DAB (3.3’- diaminobenzidine tetrahydrochloride) at a concentration of 0.5 mg/ml in Tris buffer (pH 7.6) and 0.015% H\textsubscript{2}O\textsubscript{2}. Slides were stream-rinsed with tap water, counterstained with hematoxylin for 2 min, washed in tap water, dried, mounted and coverslipped.

Sections processed with the omission of primary antibody served as negative control of immunohistochemical procedure.

**Immunohistochemical analysis of p53**

Immunostaining was assessed by two independent observers blinded to patient characteristics.

Expression of p53 was evaluated separately using the following scale: 3+ = high level (91-100% of positive cells), 2+ = medium level (11-90% of positive cells), 1+ = low level (up to 10% of positive cells), – = negative cells (0% of positive cells). For statistical analysis as positive were considered only samples with high level [3+] and medium level [2+] proteins expression. Samples scored as [1+] or [–] were considered negative.

**RESULTS**

**Immunohistochemical analysis of p53**

We analysed p53 in 42 samples of RCC and 5 samples of normal renal tissue using mouse anti-p53, clone DO7 antibody. P53 immunostaining of renal cell carcinoma samples according to subgroups (RCC-conventional type and RCC-other types) and nuclear grade were compared with respect to possible differences in p53 immunoreactivity using a two tailed Fisher’s Exact Test and the Chi-square test. Other types of RCC (papillary, chromophobe, sarcomatoid, multilocular and unclassified) were evaluated together because the number of samples in each group was too small for separate statistical analysis. \( P<0.05 \) was considered to be significant.

Fig. 2. RCC – conventional type: negative expression of p53. Note complete absence of p53 immunostaining in carcinoma cells of RCC (A). Positive expression of p53 in the nuclei of tumour cells. Immunopositive nuclei are represented by brown colour (B). Magnification: A – 40x, B – 40x.

Fig. 3. RCC – other type: tumour cells show negative expression of p53 (A). Positive expression of p53 in the nuclei of tumour cell. P53 positive nuclei show brown staining (B). Magnification: A – 40x, B – 40x.
tissue samples revealed nuclear localisation with non-restriction to any particular cell type (Fig. 2b, Fig. 3b).

In normal renal tissue specimens, staining of p53 showed weak cytoplasmic positivity in renal tubular epithelium (Fig. 1a,b).

In the first group (RCC: clear-cell), only one of the 29 samples expressed p53 at a very high level, in 1/29 samples medium expression level was detected, 8/29 samples showed low expression and 19/29 were p53 negative (Fig. 2a). Taken together, 6.9% of conventional and/or clear cell carcinoma cases showed p53 positivity and the rest (93.1%) were considered negative. Some differences were obtained in p53 expression in our set of second group (RCC – other types). Strongly positive expression was detected in one of 13 cases, medium positive were 3/13 samples, weakly positive expression had no cases and 9 tissue samples were p53 negative (Fig. 3a). In total, p53 expression was found to be four to five times higher in other types of RCC than in the clear-cell type of RCC.

Comparison of p53 expression with histopathologic subtypes

We found no statistically significant difference in p53 overexpression among the different histopathologic subtypes ($P>0.05$, $P=0.063$). For more details concerning the expression of p53 (see Table 2).

Comparison of p53 expression with nuclear grade of RCC

Nuclear grade, a clinically relevant predictor parameter, was determined and compared with p53 expression. The comparison of this parameter was evaluated in 42 clinical samples of all histopathological types of RCC.

No statistically significant difference in p53 expression in renal cell carcinomas was found with respect to the nuclear grade ($P>0.05$, $P=0.17$). The exact numbers of tissue tumour samples and the statistical analysis are shown in Table 3.

DISCUSSION

The determination of p53 status by immunohistochemistry is a widely accepted tool in surgical pathological evaluation. However, the role of p53 overexpression in RCC is still controversial. The reported results regarding both the rate of immunoreactive tumours and the impact of p53 overexpression on patient’s prognosis are inconsistent.

These divergences may be the result of low patient numbers on the one hand and different patterns of stage, nuclear grade and histopathologic subtype dependent on specimen selection on the other. Additionally, technical differences, including the use of various antibodies and staining procedures, make comparison of data difficult. Interestingly, our study also showed significant differences in p53 expression between normal and tumour tissue.

Some reports suggest that p53 is not detectable immunohistochemically in normal human tissues due to half-life of wild-type p53 (ref.19). Despite this fact, our immunostaining for p53 revealed positive reaction in our set of normal human kidney. Pillai et al.20 also observed cytoplasm staining of p53 in epithelial cells of proximal tubules of normal human kidney. Our results confirm their findings and in addition we used the same monoclonal antibody. The accumulation of p53 in normal tissues may be related to a regulatory defect mechanism between p53 and MDM2 proteins or may result from overexpression of wild-type p53. From this it follows that, the accumulation of p53 and its immunohistochemical detection

### Table 2.

<table>
<thead>
<tr>
<th>Quantity of p53 expression</th>
<th>RCC: clear cell</th>
<th>RCC: other type</th>
</tr>
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<tbody>
<tr>
<td>[3+] 1 (3.45%)</td>
<td>1 (7.7%)</td>
<td></td>
</tr>
<tr>
<td>[2+] 1 (3.45%)</td>
<td>3 (23.1%)</td>
<td></td>
</tr>
<tr>
<td>[1+] 8 (27.6%)</td>
<td>0 (0%)</td>
<td></td>
</tr>
<tr>
<td>[-] 19 (65.5%)</td>
<td>9 (69.2%)</td>
<td></td>
</tr>
<tr>
<td>Number of positive samples</td>
<td>2 (6.9%)</td>
<td></td>
</tr>
<tr>
<td>Number of negative samples</td>
<td>27 (93.1%)</td>
<td></td>
</tr>
</tbody>
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Fisher’s Exact Test (two tailed) $P>0.05$

### Table 3.

<table>
<thead>
<tr>
<th>p53 (number)</th>
<th>nuclear grade 1</th>
<th>nuclear grade 2</th>
<th>nuclear grade 3</th>
<th>unknown</th>
<th>$P$ value</th>
</tr>
</thead>
<tbody>
<tr>
<td>p53 positive</td>
<td>3 (50%)</td>
<td>1 (16.7%)</td>
<td>0 (0%)</td>
<td>2 (33.3%)</td>
<td>$P&gt;0.05$</td>
</tr>
<tr>
<td>p53 negative</td>
<td>9 (25%)</td>
<td>19 (52.8%)</td>
<td>4 (11.1%)</td>
<td>4 (11.1%)</td>
<td></td>
</tr>
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in the cytoplasm is a marker for p53 dysfunction\textsuperscript{21}. The proper interpretation of p53 cytoplasm immunopositivity remains an unresolved question.

Several relatively small studies have suggested that loss of p53 function, inferred from immunohistochemical detection in clinical samples, is relatively rare (25%, 16%, 2%) (ref.\textsuperscript{14,12,23}). Two other important articles strongly suggest that loss of p53 function is a critical event in the evolution of RCC (ref.\textsuperscript{12,26}). In our study with a similar number of patients, the overexpression of p53 was seen in 6.9% and in 30.8% of conventional and other types of RCC, respectively. A large, more recent study (\(n=97\)) concluded that p53 expression is present in 36% of cases\textsuperscript{27}. In another even larger study of 246 RCC primary and metastatic samples, p53 overexpression was detected in 29.5% of cases\textsuperscript{28}. This study also found a statistically significant difference in metastasis-free survival between p53-positive and p53-negative tumours for RCC (\(P=0.0005\)) and that p53 was an independent prognostic indicator (\(P=0.01\)). We did not find any statistical difference in nuclear p53 overexpression among histopathologic subtypes in our set of RCC. The same result was obtained by Uhlman who investigated 175 RCCs (ref.\textsuperscript{19}). On the other hand, the latter two studies described a statistically significant difference in p53 overexpression among histopathologic subtypes\textsuperscript{24,25}. The difference in p53 overexpression might reflect alternative pathways of tumourigenesis among different subtypes, possibly related to subtype-specific genetic changes. Increased p53 protein expression was accompanied by higher grade of tumour in several studies\textsuperscript{12,16}. However, in our series of samples, no relation was found between nuclear grade and increased p53 protein expression.

Taken together, all these studies suggest that, as with other cancers, one effective way that tumours can lose p53 function is through mutation. The most common in RCC carcinogenesis is the loss of 19p. Others are gain of 1q, loss of 16q, loss of 10q and loss of 17. Apart from the above example, with the loss of chromosome 17, which is the site where p53 is located, a reciprocal translocation between chromosomes 10 and 17 has been also document\textsuperscript{ed}\textsuperscript{26}. In studies of cell lines derived from renal cell carcinoma, loss of 17p heterozygosity has been documented at a higher frequency (48%) than has been inferred from the above clinical studies. This presumably reflects the selection for loss of p53 function in cells that are better able to adapt to growth ex vivo\textsuperscript{27}. It has been recently proposed that p53 is rarely mutated in renal cancer because it was already inactivated in a novel dominant manner that was neither MDM2 nor p14\textsuperscript{ARF} dependent\textsuperscript{28}. Nevertheless, Warburton et al.\textsuperscript{29} showed that p53 function is essentially "normal" and is regulated by MDM2 to a "normal" degree and is to a lesser extent regulated by p14\textsuperscript{ARF}. Binding of MDM2 to p53 can block p53 transcriptional activity by preventing it from interacting with the transcriptional machinery\textsuperscript{28}. MDM2 also causes p53 degradation by targeting it for destruction by the 26S proteasome. The latter effect is caused by the ability of MDM2 to act as an E3 ubiquitin ligase with specificity for p53 (among other targets) (ref.\textsuperscript{30}). Identification of these RCC-specific p53-inhibitory factors is the next step in this work. We hope that it will be helpful for the development of a new effective treatment of this notoriously difficult disease.

**ACKNOWLEDGEMENTS**

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