

# Immunohistochemical Panel for Differentiating Renal Cell Carcinoma with Clear and Papillary Features

Hanan AlSaeid Alshenawy

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**Abstract** Renal cell carcinoma (RCC) in which clear cells with papillary architecture are present is a difficult diagnostic challenge. The most common type, clear cell RCC, only rarely has papillary architecture. The second most common one, papillary RCC, only rarely contains clear cells. However, two recently described less-common types, clear cell papillary and Xp11 translocation RCC characteristically feature both papillary architecture and cells with clear cytoplasm. Accurate diagnosis has both prognostic and therapeutic implications. This study aims to highlight the helpful cytomorphologic and immunohistochemical features of each of these entities to enable reproducible classification. Sixty RCC cases with clear cells and papillary architecture were selected and classified according to The International Society of Urological Pathology (ISUP) Vancouver Classification of Renal Neoplasia and graded according to The International Society of Urological Pathology (ISUP) grading system for renal cell carcinoma then stained for CK7, carbonic anhydrase IX (CA IX),  $\alpha$ -methylacyl-CoA-racemase (AMACR) and TFE-3. The characteristic immunoprofile of Clear RCC is CK7-, AMACR-, CA IX+ and TFE3-, papillary RCC is CK7+, AMACR+, CAIX- and TFE3-, while for clear cell papillary RCC it is CK7+, AMACR-, CAIX+ and TFE3- and lastly Xp11translocation RCC is CK7-, AMACR+, CAIX- and TFE3+. Immunohistochemical staining for CA IX, CK7, AMACR and TFE3 comprises a concise panel for distinguishing RCC with papillary and clear pattern.

**Keywords** Renal cell carcinoma · CK7 · Carbonic anhydrase IX ·  $\alpha$ -methylacyl-CoA-racemase · TFE-3

H. A. Alshenawy (✉)  
Department of Pathology, Faculty of Medicine, Tanta University, 25  
Hamdy Gado street, Tanta, Egypt  
e-mail: hanan\_alshenawy@yahoo.com

## Introduction

Renal epithelial tumors are renal neoplasms arising from renal tubules and can be classified into many major categories based on morphology [1]. Different tumor type appears to have different outcome. With increased understanding of pathogenesis of each type of tumors, new target therapy may be developed [2].

Primary renal cell carcinomas (RCCs) with both papillary architecture and cells with clear cytoplasm may be a difficult diagnostic challenge. The most common RCC, clear renal cell carcinoma, CRCC, which represent about 75 % of the cases, may sometimes have papillary architecture. The second most common RCC, papillary renal cell carcinoma, PRCC which represent about 15 % of the RCC cases, may also contains clear cells [3]. However, 2 recently described but less-common RCCs, clear cell papillary renal cell carcinoma, CPRCC and Xp11 translocation RCC, characteristically feature both papillary architecture and cells with clear cytoplasm. Accurate diagnosis of these distinct entities has both prognostic and therapeutic implications [4]. In most cases, routine cytomorphologic features are sufficient to make the correct diagnosis. In other cases, immunohistochemical markers are needed to establish the correct diagnosis [5].

CPRCC is a recently recognized renal neoplasm, composed of an admixture of cystic, glandular, solid, and papillary components, all lined by cells with clear cytoplasm, usually of low nuclear grade. The nuclei are characteristically located away from the basement membrane to show a “piano-key-like” pattern [3]. The outcome data are limited; however, the available data suggest that this tumor is not aggressive and patients with this type of tumor usually have a good prognosis. Originally they were discovered in a background of end-stage renal disease and acquired cystic kidney disease. However, several studies have reported them in patients with otherwise healthy kidneys [6].

Xp11 translocation RCC initially described in children and young adults. Recently, the term “MiTF/TFE family

translocation-associated carcinoma” has been proposed for tumors that have translocations involving TFE3. TFE3 is transcription factor that belong to the same family of transcription factors that will overexpress nuclear TFE3 at the protein level. These immunohistochemical findings are important given the occurrence of these tumors in the adult population, as they morphologically overlap with CCRCC and PRCC. Recently, cathepsin K, a protein whose expression is mediated by MiTF, has been shown to be sensitive and specific for differentiating translocation-associated carcinomas from CCRCC. From rare cases reported in the literature, these tumors do not appear to respond to immunotherapy [7]. Because these neoplasms have been only recently recognized, outcome data are still premature and good long-term follow-up data are necessary. Published outcome series in adults show a poor prognosis [4].

The treatment paradigm for renal tumors in general and renal cell carcinomas (RCCs) specifically are changing, and these changes are in part driven by tumor classification. Pathologists will play a major role in these new treatment models. Traditionally, RCC has been considered a surgical disease. In some cases, surgery with its associated complications and negative impact on long-term renal function may be very harmful, so follow up after chemo radiotherapy may be used in low grade small tumours [8].

Cytokeratins are a family of intermediate filaments that characterize epithelial differentiation, There have been conflicting results on the expression of CK7 in renal epithelial tumors in the literature as some authors recognized its role in the differentiation of “non-clear cell” RCC from CRCC [9].

The most useful positive immunohistochemical stain in supporting a diagnosis of PRCC is  $\alpha$ -methylacyl-coenzyme A racemase (AMACR), a protein involved in the metabolism of branched chain fatty acids. Subsequently, AMACR was also noted to diffusely label PRCC in a granular cytoplasmic fashion. It is now recognized that AMACR can show positivity in tumors from many different organs and in several different types of renal tumors, including most MiTF/TFE family translocation carcinomas. But AMACR staining has conflicting results in CPRCC as it is often negative [10] but in other studies it is focally or, rarely, diffusely positive [6].

Carbonic anhydrase IX (CA IX) protein, a member of the carbonic anhydrase family, is thought to play a role in the regulation of cell proliferation in response to hypoxic conditions and may be involved in oncogenesis and tumor progression. Previous immunobiochemical studies of malignant and benign renal tissues revealed that CA IX was also highly expressed in RCC, suggesting that CA IX expression may be a useful diagnostic biomarker. Clinical tumor targeting studies with a monoclonal antibody to CA IX have shown that CA IX shows promise as a marker for selecting patients with advanced disease who would benefit from certain specific systemic agents, specifically interleukin-2 (IL-2) [7, 11].

This work aims to highlight the helpful cytomorphologic and immunohistochemical features of each of these entities to enable reproducible classification. We examined the expression of 4 markers in a series of the 4 major renal cell tumors with clear and papillary architecture. In this study, we evaluated the expression of carbonic anhydrase IX (CA IX),  $\alpha$ -methylacyl-CoA-racemase (AMACR), CK7 and TFE-3 for differential diagnosis and subclassification.

## Materials and Methods

### Case Selection and Histopathological Study

A retrospective study was performed on RCC cases selected from January 1, 2010 to April 30, 2014. A total of 250 cases of RCC were removed by nephrectomy either partial or radical and brought to the Department of Pathology, University of Tanta. Representative tissue sections from the surgical specimens were fixed in 10 % buffered formalin and embedded in paraffin. For routine microscopy, 4  $\mu$ -thick sections were stained with Hematoxylin and Eosin (H&E). The clinical sheets for all cases were reviewed. The cases were classified according to The International Society of Urological Pathology (ISUP) Vancouver Classification of Renal Neoplasia [12]. Tumors that fulfill the morphological criteria of clear and papillary renal cell neoplasms were selected. They were 60 cases. Only the selected cases were assessed for size, laterality, multifocality, presence of associated end-stage renal disease (ESRD). Then they assessed histologically for the presence of branched tubular structures, subnuclear vacuoles, acini, thin walled sinusoid-like vessels, ‘secretory’ cells with nuclei aligned at the apical end of the cells, cystic components, character of the stromal compartment, presence of tumor pseudocapsule, and calcification. The selected carcinomas were later reevaluated for morphologic characteristics of those tumors that qualify them in either one of the following categories; CRCC, PRCC, CCPRCC or Xp11 translocation RCC. Specifically, the criteria used for classification of a tumor as a CCPRCC included the following: (1) diffuse cytoplasmic clarity; (2) papillary, tubular or cystic architecture; and (3) characteristic linear arrangement of the nuclei away from the basement membrane [10]. Xp11 translocation RCC cases were confirmed by TFE3 immunostaining positivity. The International Society of Urological Pathology (ISUP) grading system for renal cell carcinoma grading system was applied to access the nucleolar grades [13]. The tumors were staged according to the 2010 UICC/AJCC consensus guidelines [14].

### Immunohistochemical Study and Evaluation

Immunohistochemical staining was performed using the following antibodies: CK7 (OV-TL 12/30, 1:100, DAKO,

Glostrup, Denmark), CA IX (dilution 1:200, mouse monoclonal, Leica), AMACR (13H4, 1:100; DAKO, Glostrup, Denmark) and TFE3 (1:1500, Santa Cruz Biotechnology Inv., Santa Cruz, CA, USA).

Evaluation of the immunohistochemical staining was performed by light microscopy using a 10× objective lens with the selective use of a 20–40 × objective lens for confirmation. The interpretation of immunoreactivity was performed in a semiquantitative manner by analyzing the extent of the staining positivity of the tumor cells. Immunostaining of greater than 10 % of tumor cells was required for scoring as a positive case. The interpretation score was as follows: 0 or negative ≤10 % tumor cell positivity; +1 or weak = 11–25 % tumor cell positivity; +2 or moderate = 26–50 % tumor cell positivity; and +3 or strong >50 % tumor cell positivity [6]. Cytoplasmic and/or membranous expression of CK7 and AMACR were considered positive. Only distinct membranous staining for CA IX and distinct nuclear staining for TFE3 were considered positive [15].

### Statistical Analysis

The clinicopathological and immunohistochemical features were tested for their association with the histological subtype using Student's *t*-test for continuous variables and the chi-square test (or Fisher's exact test) for qualitative variables. All statistical analyses were performed using the SPSS 13.0 software. Statistical significance was considered when *P* value ≤0.05.

## Results

### Clinicopathologic Findings of the Selected Cases

The clinicopathologic findings are listed in Table 1. The selected cases were comprised of 60 cases; 40 men and 20 women (men to women ratio, 2:1) with a mean age of 54 years (range, 40–72 years). A definite positive history of hemodialysis was found in 14 cases end stage renal disease with a duration ranging from 1 to 24 years. The International Society of Urological Pathology (ISUP) grading system was as follows: 24 were grade I, 21 were grade II and 15 were grade III. The tumors pathologic stage was as follows; 38 cases were stage I, 15 were stage II, 7 were stage III and no cases were stage IV.

Overall, of the 60 cases, 28 (47 %) were CRCC, 15 (25 %) PRCC, 8 (13 %) CPRCC and 9 (15 %) Xp11 translocation RCC were identified. The cases were classified according to the published data into these categories according to some variables summarized in Table 2.

CRCC are usually characterized by optically clear cytoplasm and a very well-defined cytoplasmic membrane. The

**Table 1** The clinicopathological variables of the selected cases

Clinicopathological variable		Number of cases
Size	Less than 7 cm	45
	More than 7 cm	15
Laterality	Right	30
	Left	24
	Bilateral	6
Multifocality	Yes	8
	No	52
End stage renal disease	Yes	14
	No	46
Grade	Grade I	24
	Grade II	21
	Grade III	15
Stage	Stage I	38
	Stage II	15
	Stage III	7
	Stage IV	0
Histological types	CRCC	28
	PRCC	15
	CPRCC	8
	Xp11 translocation RCC	9

cells are arranged in nests, alveolar architectures, strands and papillae, or cysts. The stroma contains a very typical and prominent capillary network. Hemorrhage and necrosis are frequent (Fig. 1a).

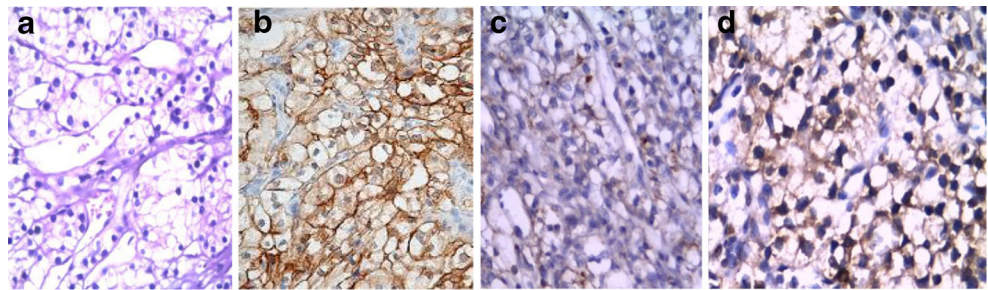
Papillary architecture is the characteristic histology of renal tumors in PRCC. The Delicate fibrovascular stalks lined by small cells with low grade nuclei and scant cytoplasm (Fig. 2a). Psammoma bodies and clear cells are common.

**Table 2** Microscopic finding in the selected cases

Histological features	CRCC	PRCC	CPRCC	Xp11 translocation
Clear cells	Yes	Yes	Yes	Yes
'secretory' cells with nuclei aligned at the apical end of the cells	No	No	Yes	No
Subnuclear vacuoles	No	No	Yes	No
True papillae	No	Yes	Yes	Yes
Tubules	No	No	Yes	Yes
Cysts	Yes	Yes	Yes	Yes
Acini	Yes	No	Yes	Yes
Biologic aggressiveness	Yes	Yes	No	Yes
Capsule	No	Yes	Yes	No
Psammoma bodies	No	Yes	No	No
Leiomyomatous stromal components	Yes	No	Yes	Yes
Thin walled sinusoid-like vessels	Yes	No	No	No



**Fig. 1** A case of CRCC grade I showing clear cells with faint papillary structures (H&E  $\times 100$ , **a**) showing positive membranous staining for CA IX ( $\times 400$ , **b**) and weak positivity for both CK7 ( $\times 200$ , **c**) and AMACR ( $\times 400$ , **d**)



CPRCC were well circumscribed with a well-defined, thin, fibrous capsule and were composed histologically of a mixture of cysts and papillae. Tubular/acinar features were also common. Even in the predominantly cystic areas, the lining cells frequently showed papillary infoldings. The tumors were composed of cuboidal or columnar cells with clear cytoplasm, small hyperchromatic, round or oval nuclei, and inconspicuous nucleoli. A conspicuous nuclear positioning away from the basement membrane with subnuclear vacuoles was noted in all cases examined (Fig. 3a). The cysts, tubules, and acini contained pink secretion. Renal capsular and vascular invasion, mitoses, tumor necrosis, and stromal aggregates of macrophages were not observed. The stromal component formed variably thick bands of fibrous tissue admixed with strands of leiomyomatous tissue inside the tumors

Renal carcinomas with Xp11.2 translocation have a characteristic morphology; they are composed of very large clear cells that form nests, alveoli, and papillae (Fig. 4a) accompanied by an abundance of psammoma bodies. The cells are not very cohesive, which causes alveolar images and pseudo papillae.

#### Correlation Study Between the Studied Cases and Clinicopathological Variables

In correlation between the different pathological types of RCC and the clinicopathological variables, it was found that there is a significant difference between the histological type of RCC and the tumour size as most of the cases of PRCC, Xp11.2 translocation RCC (12/15 and 7/9 respectively) and all the cases of CPRCC were of smaller size ( $\leq 7$  cm) while most of the large sized tumours were CRCC. As regards the laterality, PRCC was the most common bilateral RCC (5/15) as 5 cases

out of the 6 bilateral cases were PRCC with statistical difference. Also the multifocality was only seen in PRCC (8/15) as all the multifocal cases were PRCC with strong statistical difference. End stage renal disease was associated with most of CPRCC (6/8) and some CRCC (6/28) with statistical significance.

The grade was correlated significantly with RCC types as it was low in all the cases of CPRCC (either grade I or II). On the other hands, more than half of the Xp11 translocation RCC cases were of high grade (grade III). The tumour stage was also low in most of the cases of CRCC, PRCC and all the cases of CPRCC with strong significant difference. These data were summarized in Table 3

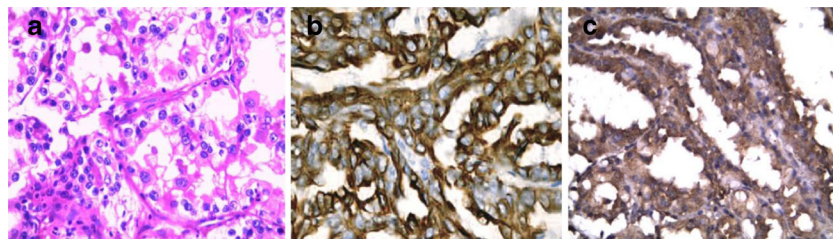
#### Immunohistochemical Findings in the Studied Cases

The immunohistochemical findings are listed in Table 4.

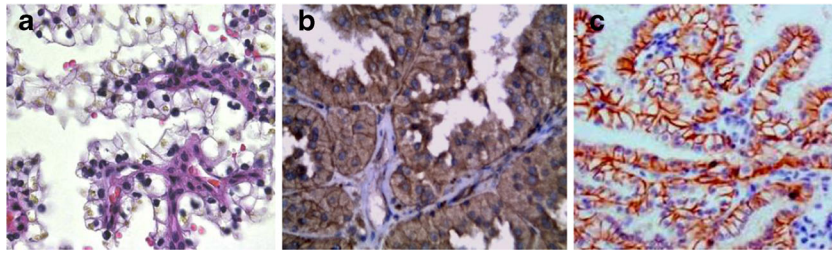
The immunohistochemical profile of CRCC was that all the cases show positive membranous staining for CA IX (18/28 and 10/28 were strongly positive (Fig. 1b) and moderately positive consequently), while they were negative for both CK7 and AMACR, although both may be weekly positive (2/28 (Fig. 1c) and 5/28 were (Fig. 1d) weak positive respectively), especially in higher-grade tumors. TFE3, marker of Xp11 translocation RCC, is consistently negative.

PRCC cases show strong membranous positivity for CK7 (Fig. 2b). Staining with AMACR demonstrates cytoplasmic granular positivity in all the cases (Fig. 2c). The CA IX staining is either completely negative (13/15) or may be weak positive near areas of necrosis (2/15). TFE3 is consistently negative.

CPRCC displays a unique IHC profile overlapping with CRCC and PRCC. CPRCC show positive cytoplasmic



**Fig. 2** A case of PRCC grade II showing prominent papillae with occasional clear cells (H&E  $\times 100$ , **a**). It expresses strong membranous positivity for CK7 ( $\times 200$ , **b**). AMACR demonstrates cytoplasmic granular positivity ( $\times 200$ , **c**)



**Fig. 3** A case of CPRCC grade I show clear cells with low grade nuclei show characteristic linear polarization away from the basement membrane (H&E  $\times 200$ , **a**), it demonstrates positive cytoplasmic

staining for CK7 ( $\times 200$ , **b**), CA-IX diffusely expressed in a membranous distribution in cup-shaped distribution ( $\times 400$ , **c**)

staining for CK7 in all the cases (Fig. 3b); similar to PRCC. AMACR, TFE3 were negative in all the cases examined. The tumor cells also expressed CA-IX diffusely in a membranous distribution (6/8 and 2/8 were strongly and moderately positive consequently); the absence of staining along the luminal borders of the tumor cells was quite characteristic (cup-shaped distribution) (Fig. 3c). TFE3 is consistently negative.

Renal carcinomas with Xp11.2 translocation do not express, or only very weakly express CK7 and CA IX (8/9 and 7/9 were negative for CK7 and CA IX respectively) but they express AMACR (6/9 (Fig. 4b) and 3/9 were strongly and moderately positive respectively). In addition, this group of carcinomas is characterized by nuclear positivity with the TFE3 transcription factor (Fig. 4c).

## Discussion

The World Health Organization classification of renal tumors synthesizes morphological, immunohistochemical, molecular, and clinical data to define distinct entities that are biologically and clinically relevant. Although most epithelial renal tumors can be diagnosed by morphology alone, the diagnosis can be difficult due to the overlap of histological features. In these cases, differentially expressed immunohistochemical markers can be of help [2]. In this study, we performed a 4-antibody panel on RCC cases that show both papillary and clear cell architecture.

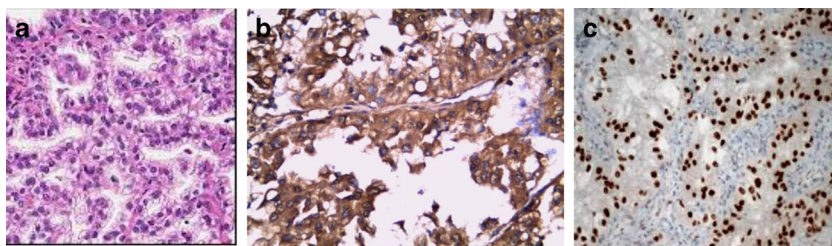
The present study included 60 cases consist of CRCC, PRCC, CPRCC and Xp11.2 translocation RCC. Most of PRCC, Xp11.2 translocation RCC and all CPRCC cases were

of smaller size ( $\leq 7$  cm). PRCC was found to be the most common bilateral and multifocal RCC. End stage renal disease was associated with most of CPRCC but also we observed that CPRCC can occur in otherwise normal kidneys and this was observed in **Michelle et al.** study [10].

CRCC was usually of aggressive architecture as it was characterized by tumor necrosis, mitoses, vascular invasion, and a characteristic network of small, thin-walled sinusoid-like blood vessels, so it was usually presented with higher stage and this was seen in **Brannon et al.** study [16]. PRCC usually has cystic change and papillary architecture. Clear cell change can become quite extensive in some tumors causing morphologic confusion. The cytoplasmic clearing is typically seen in association with tumor necrosis and this was similar to **Aydin et al.** study [17].

CPRCC were well circumscribed with a well-defined fibrous capsule, and were composed of a mixture of cystic, papillary, tubular, and acinar components. The nuclei show characteristic polarization in a linear array away from the basement membrane and of low grade. The stromal component formed variably thick bands of fibroliomyomatous tissue inside the tumors among the epithelial component. These features were observed in previous studies [10, 18].

Xp11.2 translocation RCC has a distinctive pattern which is the presence of both clear cells and papillary architecture. Nuclei tend to be high grade and the cytoplasm may be clear to granular and eosinophilic. Numerous psammomatous calcifications and stromal hyaline nodules are common. Most of our cases were of higher grades and this was similar to **Rajen et al.**, [7].



**Fig. 4** A case of Xp11 translocation grade II shows clear cells and papillary architecture with high grade nuclei (H&E  $\times 200$ , **a**), strongly expressing AMACR in the cytoplasm ( $\times 200$ , **b**), and characteristically show nuclear positivity for TFE3 ( $\times 200$ , **c**)

**Table 3** Correlation study between the studied cases and clinicopathological variables

Histological type	Size		Laterality			Multifocality		ESRD		Grade			Stage			
	≤7 cm n=45	>7 cm n=15	Rt n=30	Lt n=24	Bilateral n=6	Yes n=8	No n=52	Yes n=14	No n=46	I n=24	II n=21	III n=15	I n=38	II n=15	III n=7	IV n=0
CRCC n=28	18	10	15	13	0	0	28	6	22	10	15	3	12	12	4	0
PRCC n=15	12	3	6	4	5	8	7	1	14	6	2	7	13	1	1	0
CPRCC n=8	8	0	3	4	1	0	8	6	2	5	3	0	8	0	0	0
Xp11.2 translocation RCC n=9	7	2	6	3	0	0	9	1	8	3	1	5	5	2	2	0
P value	0.01		0.04			0.003		0.05		0.04			0.008			

The distinction between CRCC, PRCC, CPRCC and Xp11.2 translocation RCC is critical because of different behavior, prognosis and treatment [6].

In the present study, the grade was low in all the cases of CPRCC, and the stage was also low in most of the cases of CRCC, PRCC and all the cases of CPRCC with strong significant difference. On the basis of our results, it appears that CPRCC represents a distinct form of low grade RCC, the recognition of which may be important for prognosis and clinical management no case of CPRCC has behaved aggressively as in previous studies [19]. So immunohistochemical study to differentiate difficult cases may be needed to classify them.

CK7 is a commonly used marker for RCC. Our results showed diffuse positivity for this marker in a membranous pattern, consistent with previous reports [20], confirming its usefulness in differentiating both PRCC and CPRCC from

CRCC and Xp11 translocation RCC which is consistently negative in both.

AMACR is a mitochondrial enzyme, which mediates the oxidation of branched-chain lipids. In this study, it has been shown that this marker is usually positive in PRCC and Xp11translocation RCC with a diffuse strong cytoplasmic staining pattern. Our study showed that AMACR was usually negative or weakly positive in CRCC while CPRCC was negative for this marker, consistent with prior reports [2, 3].

CA IX is a hypoxia-induced protein and is predominantly reported to be positive in CRCC [2, 21]. In this study, CPRCC is largely positive for this marker, a result that concurs with previous observations [11]. CA IX is useful for the differentiation of CRCC and CPRCC from PRCC with clear cell changes. CRCC and CPRCC are usually positive for this marker, whereas the PRCC is either completely negative or may be weakly positive near areas of necrosis. However, the pattern of expression of CA IX in CPRCC differs from that seen in CRCC in that most cells in clear-cell papillary RCC lack labeling on the luminal aspect. This pattern of staining was not observed in any of the CRCCs evaluated in this study and this was in agree with **Stephen et al.**, [15] and **Tickoo and Reuter** [22]. Xp11 translocation RCC diagnosis is supported by TFE3 positivity.

In summary, The characteristic immunoprofile of CRCC is CK7-, AMACR-, CA IX+ and TFE3-, for PRCC it is CK7+, AMACR+, CA IX- and TFE3-, while for CPRCC it is CK7+, AMACR-, CA IX+ and TFE3- and lastly Xp11translocation RCC is CK7-, AMACR+, CA IX- and TFE3 + .

To conclude, immunohistochemical staining for CA IX, CK7, AMACR and TFE3 comprises a concise panel for distinguishing RCC with papillary and clear pattern. Our marker panel is a clear advancement in terms of immunohistochemistry application for RCC subtype differentiation when papillary and clear cells are the predominant architecture as this is very crucial for further prognosis and targeted therapy because of the different behavior of each type.

**Table 4** Immunohistochemical profile of the studied cases

Studied marker	CRCC N=28	PRCC N=15	CPRCC N=8	Xp11 translocation N=9	
CK7	Negative	26	0	0	8
	+	2	0	0	1
	++	0	0	0	0
	+++	0	15	8	0
AMACR	Negative	23	0	8	0
	+	5	0	0	0
	++	0	0	0	3
	+++	0	15	0	6
CA IX	Negative	0	13	0	7
	+	0	2	0	2
	++	10	0	2	0
	+++	18	0	6	0
TFE3	Negative	28	15	8	9
	+	0	0	0	0
	++	0	0	0	0
	+++	0	0	0	0



**Conflict of Interest** The author declares that there is no conflict of interest.

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