# Expression of CK19, CD105 and CD146 are associated with early metastasis in patients with renal cell carcinoma

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Abstract. The present study was designed in order to explore the association between the early metastasis of renal cell carcinoma (RCC) and biological markers of tumor cells. A total of 200 patients with RCC, who received a nephrectomy between January 2015 and October 2015, were enrolled in the present study, while 100 healthy patients served as controls. The expression of cytokeratin 19 (CK19), endoglin (CD105) and cluster of differentiation 146 (CD146) were detected using immunohistochemical staining and western blotting. All three markers were highly expressed in tumor tissues compared with adjacent normal tissues. Subsequently, an enzyme-linked immunosorbent assay was used to detect the differential expression of CK19, CD105 and CD146. The results revealed that there was a statistically significant difference in the expression of CK19 and CD105 between the two groups (P<0.05), whereas CD146 did not exhibit a statistically significant difference. The results of further experiments revealed no significant difference between four time points (Q1, 1 day pre-operation; Q2, 1 day post-operation; Q3, 1 week post-operation; and Q4, 1 month post-operation). Then, subgroup analysis was performed based on whether patients were circulating tumor cell (CTC)-positive or not, and the difference between the Q1 time point and other three time points (Q2-4). The results revealed no difference between the CTC-positive and -negative groups, and no difference between the time points Q1 and Q2. However, the expression of CK19 and CD105 exhibited a significant difference between CTC-positive and CTC-negative groups according to the difference between the time points Q1 and Q3. Furthermore, on the basis of the difference between Q1 and Q4, the expression of CK19, CD105 and

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*Key words:* renal cell carcinoma, cytokeratin 19, endoglin, cluster of differentiation 146, circulating tumor cell, metastasis

CD146 were significantly different (P<0.05). Taken together, the results suggested that CK19, CD105 and CD146 markers of peripheral blood may be considered to be effective tools to evaluate the early metastasis in a CTC-positive condition. CK19, CD105 and CD146 may be useful for CTC in evaluating the prognosis of patients with RCC, although a larger sample size is necessary for further investigation.

#### Introduction

Despite substantial improvement in the survival rates of patients with cancer, cancer remains the most common mortality-associated disease in the United States in 2015 (1). Renal cell carcinoma (RCC) is a lethal malignant tumor type in the urinary system, accounting for 3% of all malignancies globally in 2015 (2). In China, the incidence of this malignancy is notably increasing. Nephrectomy remains the first-line therapy in patients with a solid tumor (3). However, the frequent metastasis greatly impedes the curative effect following kidney resection. Nearly 30% of patients with RCC progress to metastatic disease subsequent to radical nephrectomy surgery (4). Numerous independent risk factors, including nodule number, tumor stage [American Joint Committee on Cancer (AJCC, 2010)] (5) and grade, were identified to have a close correlation with the recurrence and metastasis of patients with RCC (6). Nevertheless, these parameters are not absolutely accurate. Hence, the introduction of novelty predictive and prognostic biomarkers will aid oncologists in performing preventative treatment and individual therapy in patients with RCC.

Cytokeratin 19 (CK19) is an epithelial cytoskeleton marker and is highly expressed in epithelial tumor types (7). One previous study has revealed that CK19 may be an appropriate marker for the detection of cancer cells (8). Furthermore, it may function as a prognostic indicator for patients with cancer (9). Endoglin (CD105) was first reported over two decades ago and identified as an accessory receptor for transforming growth factor- $\beta$  (TGF- $\beta$ ) (10,11). It is notably upregulated in endothelial cells and has been considered to be a biomarker for tumor angiogenesis (12). The role of CD105 in the early metastasis of a tumor has gained attention, including in squamous cell carcinoma and pancreatic ductal adenocarcinoma (13,14). In RCC, the function of CD105 remains controversial (15,16). Cluster of differentiation 146 (CD146) is a 113 kD membrane glycoprotein which was first reported in malignant melanoma (17). One previous study has reported that CD146 is only expressed in blood vessels and smooth muscle cells (18). Subsequent research has demonstrated that CD146 has multiple functions in development, immunity and angiogenesis (17). CD146 is additionally implicated in the development of the kidney and the nervous system (19,20). However, to the best of our knowledge, there are no previous reports on the association between CD146 expression and RCC metastasis following surgical treatment.

Circulating tumor cells (CTCs) are cells that exist in the blood that have antigenic and/or genetic traits of a specific tumor type (21). They have been used to indicate a poor prognosis in a variety of cancer types, including breast cancer, prostate cancer, colorectal cancer and lung cancer (22-25). The detection of CTCs is easy to implement via peripheral blood samples and may be conducive to improving cancer prognostication and treatment (26).

In the present study, the aim was to explore the association between CK19, CD105 and CD146 and the early metastasis of RCC following surgery, and to evaluate whether the expression of CK19, CD105 and CD146 is associated with the early metastasis of patients with CTC-positive RCC.

#### Materials and methods

*Ethics statement*. The present study was approved by the Ethical Committee of Second Affiliated Hospital of Medical School, Xi'an Jiaotong University (Xi'an, China). The procedures in the present study followed the principles of the Declaration of Helsinki. Written informed consent was obtained from all patients prior to the study.

Patient selection. The present retrospective study recruited 200 RCC cases between January 2015 and October 2015. Of these, there were 100 male and 100 female patients with RCC at stage I, II and III (AJCC, 2010) (5) of the disease (from 70, 122 and 8 patients, respectively) who were accepted for surgical treatment at the Department of Urology, Second Affiliated Hospital of Medical School, Xi'an Jiaotong University (Xi'an, China). Inclusion criteria were as follows: i) Pathological diagnosis confirmed RCC; ii) clinical stage cT1-3 (AJCC, 2010) (5), ready for nephrectomy; iii) distant metastasis were not identified prior kidney resection. Exclusion criteria were as follows: i) Patients with RCC with other malignancies; ii) patients who had undergone chemotherapy or radiotherapy prior to surgery. A total of 100 patients with no sign of tumor in pathology were additionally included as the control group. The mean age of the patients with RCC and without were 60 years (age range from 40 to 79 years) and 61.36 years (age range from 39 to 75 years), respectively.

Blood sampling and tumor tissues collecting. Once informed consent was obtained from all patients, 10 ml peripheral venous blood samples were collected at four different times (Q1, 1 day pre-operation; Q2, 1 day post-operation; Q3, 1 week post-operation; and Q4, 1 month post-operation), transferred to the laboratory and used for further experiments. Tumor tissues and adjacent normal tissues were resected during surgery, collected and stored at -80°C for subsequent experiments. *Hematoxylin-eosin (H&E) staining.* All surgical specimens were collected and transferred to the laboratory. Subsequently, the samples were fixed using 4% paraformaldehyde for 48 h at room temperature. Then, the samples were embedded in paraffin and submitted to H&E staining at room temperature for 48 h.

Immunohistochemistry. Tumor samples (4  $\mu$ m thick) were fixed with 4% paraformaldehyde at room temperature for 48 h and embedded in paraffin. Following deparaffinization and rehydration as follows: Ethanol for 15 min 3 times, 95% ethanol for 2 h twice, 90% ethanol for 1 h, 80% ethanol for 1 h and 70% ethanol for 30 min, the CD105 antigen was retrieved using a citric acid buffer (pH 6.0; Boster Biological Technology, Pleasanton, CA, USA). Subsequently, the samples were blocked using endogenous peroxidase for 30 min at room temperature. Following washing with potassium-free phosphate-buffered saline (PBS) buffer 3 times, the samples were incubated with monoclonal antibodies against CK19 (cat no. 13092), CD105 (cat no. 14606) and CD146 (cat no. 13475) (Cell Signaling Technology, Inc., Danvers, MA, USA; dilution 1:100) overnight at 4°C. Subsequent to rewarming, the samples were incubated with the corresponding secondary antibodies [anti-mouse IgG (cat no. BA1051) and anti-rabbit IgG (cat no. BA1055) horseradish peroxidase-conjugated secondary antibodies; Boster Biological Technology; dilution 1:1,000] for 1 h at room temperature. Then the sections were stained using Mayer's hematoxylin for 5 min at room temperature, followed by dehydration (70% ethanol for 30 min, 80% ethanol for 1 h, 90% ethanol for 1 h, 95% ethanol for 2 h twice and ethanol for 15 min 3 times) and transparency. Finally, the sections were mounted with neutral resins and visualized using a light microscope (x20 and x40).

Western blotting. Briefly, total proteins were washed using PBS buffer 3 times and extracted using RIPA lysis buffer (50 mM Tris-HCl pH 7.4, 150 mM NaCl, 0.1% Triton X-100, 1% PMSF) for 10 min at 4°C. Following centrifugation for 15 min at 14,000 x g at 4°C. Protein concentration was determined using a Bicinchoninic Acid Assay kit (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany). Proteins were separated using sodium dodecyl sulfate-polyacrylamide gel electrophoresis (10%) and transferred to polyvinylidene difluoride membranes. Then, the membranes were blocked with 5% nonfat milk for 1 h at room temperature and incubated with primary antibodies against CK19 (cat no. 13092), CD105 (cat no. 14606) and CD146 (cat no. 13475) (Cell Signaling Technology, Inc.; dilution 1:1,000) overnight at 4°C. The membranes were washed 3 times using tris-buffered saline with Tween-20 buffer (10 mM Tris-HCl pH 7.4, 150 mM NaCl and 0.1% Tween-20) and incubated with horseradish peroxidase-conjugated secondary antibodies [anti-mouse IgG (cat no. BA1051) and anti-rabbit IgG (cat no. BA1055); Boster Biological Technology; dilution 1:1,000] at room temperature for 1 h. Finally, the protein bands were visualized using an enhanced chemiluminescence substrate.

*Enzyme-linked immunosorbent assay (ELISA).* Blood samples (5 ml) were collected from patients and controls. Following centrifugation at 4°C at 1,200 x g for 10 min, the serum levels of CK19, CD105 and CD146 were determined using an ELISA kit (R&D Systems, Inc., Minneapolis, MN, USA) according

Table I. Clinical characteristics of	patients with RCC.
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Variables	Frequency	Percent (%)
Age (years)		
Median age (range)	60 (40-79)	
Sex		
Female	100	50.0
Male	100	50.0
Tumor grade		
I	70	35.0
II	122	61.0
III	8	4.0
TNM stage		
T1aN0M0	72	36.0
T1bN0M0	98	49.0
T2aN0M0	23	11.5
T3aN0M0	7	3.5
Pathological type		
Clear cell RCC	166	83.0
Papillary RCC	34	17.0
Surgical		
NSS	135	67.5
RN	65	32.5
CTC		
Positive	90	45.0
Negative	110	55.0

RCC, renal cell carcinoma; T, tumor; N, node; M, metastasis; NSS, nephron-sparing surgery; RN, radical nephrectomy; CTC, circulating tumor cells.

to the manufacturer's protocol. Each sample was detected in duplicate.

*CTC analysis*. The CellSearch system (Janssen Diagnostics LLC, Raritan, NJ, USA) was used for the detection and evaluation of CTCs according to the manufacturer's protocol. A blood volume of 7.5 ml was prepared for each sample analysis.

Statistical analysis. All analyses were calculated using SPSS 19.0 software (IBM Corp., Armonk, NY, USA). The results are represented as the mean  $\pm$  standard deviation. A paired Student's t-test was used for the analysis of numerical data. P<0.05 was considered to indicate a statistically significant difference.

## Results

*Patient characteristics*. The present study analyzed 200 patients with RCC who underwent nephrectomy at the Second Affiliated Hospital of Medical School, Xi'an Jiaotong University between January 2015 and October 2015. An additional 100 patients without RCC were recruited as controls. The mean age of the patient and control groups were 60 years (ranging from 40 to 79 years) and 61.36 years (ranging from

Table II. CK19, CD105 and CD146 expression between patient and control groups.

Variables	Patients group	Control group	P-value
CK19	3.29±1.89	2.73±1.01	0.038
CD105	6.07±3.18	$4.90 \pm 1.91$	0.014
CD146	204.90±100.63	217.18±80.26	0.471

CK19, cytokeratin 19; CD105, endoglin; CD146, cluster of differentiation 146.

Table III. Expression of CK19, CD105 and CD146 in patients with a CTC-positive condition according to the difference between the time points of Q1 and Q3.

Variables	CTC-positive group	Control group	P-value
CK19	0.667±0.877	-0.176±0.266	0.010
CD105	$0.760 \pm 0.849$	-0.446±0.346	0.001
CD146	19.495±34.640	-19.390±113.124	0.337

CTC, circulating tumor cells; CK19, cytokeratin 19; CD105, endoglin; CD146, cluster of differentiation 146.

39 to 75 years), respectively. All patients had a non-metastatic form of the disease (stage I-III) (Table I).

CK19, CD105 and CD146 expression. Firstly, H&E staining was used to confirm whether the resected tissues were from the solid tumor. The results revealed notable tumor cell clusters different from the adjacent epithelial cells (Fig. 1A). Subsequently, the expression of CK19, CD105 and CD146 in tumor tissues and adjacent normal tissues were detected using immunohistochemistry staining. Positive CK19 staining was observed in the cytoplasm of the tumor cells of 200 patients (Fig. 1B). The staining of the tumor cells of all patients was strong and cytoplasmic. The distribution of CD105 and CD146 staining were also located in the cytoplasm of tumor cells (Fig. 1B). To further explore the differential expression of CK19, CD105 and CD146 between tumor and normal tissues, western blotting was performed. The results demonstrated that CK19, CD105 and CD146 are highly expressed in tumor tissues compared with their corresponding adjacent normal tissues (Fig. 1C).

Association between the early metastasis of patients with RCC and the expression of CK19, CD105 and CD146. The application of ELISA to analyze biomarker expression in the peripheral blood of patients with a tumor is an emerging concept in the management of cancer (25). Thus, the expression of CK19, CD105 and CD146 were detected at four different time points in patient and control groups using ELISA (Q1, 1 day pre-operation; Q2, 1 day post-operation; Q3, 1 week post-operation; and Q4, 1 month post-operation). The results revealed that there is a significant difference between the

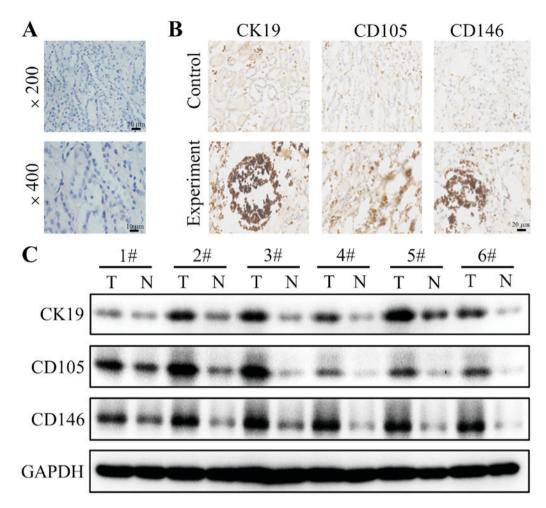


Figure 1. CK19, CD105 and CD146 expression in renal cell carcinoma tissues. (A) Representative tumor samples with hematoxylin-eosin staining. (B) Representative sections of CK19, CD105 and CD146 in tumor tissues compared with adjacent normal tissues, which were detected using immunohis-tochemical staining. (C) Representative blots extracted from tumor tissues and normal tissues were subjected to western blotting for CK19, CD105 and CD146. GAPDH was used as a loading control. Each number corresponds to a different patient. Representative blots of three experiments are shown. CK19, cytokeratin 19; CD105, endoglin; CD146, cluster of differentiation 146.

Table IV. Expression of CK19, CD105 and CD146 in patients with a CTC-positive condition according to the difference between the time points of Q1 and Q4.

Variables	CTC positive group	Control group	P-value
CK19	1.082±1.555	-0.026±0.163	0.038
CD105	$1.403 \pm 2.395$	-0.201±0.120	0.049
CD146	57.943±56.742	-28.270±79.113	0.015

CTC, circulating tumor cells; CK19, cytokeratin 19; CD105, endoglin; CD146, cluster of differentiation 146. Table V. Expression of CK19, CD105 and CD146 in patients with a CTC-negative condition according to the difference between the time points of Q1 and Q4.

Variables	CTC negative group	Control group	P-value
CK19	0.100±1.448	-0.026±0.163	0.788
CD105	-0.471±0.566	-0.201±0.120	0.156
CD146	-28.790±44.910	-28.270±79.113	0.98

CTC, circulating tumor cells; CK19, cytokeratin 19; CD105, endoglin; CD146, cluster of differentiation 146.

expression of CK19, CD105 between the patient and control groups (P<0.05), but not for CD146 (Table II). Then, the differential expressions of CK19, CD105 and CD146 between all four time points were investigated through pairwise comparison. The results indicated no statistically significant differences (data not shown).

CTCs have been identified as a predictor of early metastasis in patients with cancer (26). Based on whether patients were CTC-positive or CTC-negative, subgroup analysis was performed. The results demonstrated that the expression of CK19, CD105 and CD146 was not significantly different between CTCs-positive and CTC-negative groups or between four time points using pairwise comparison (data not shown). These results are in accordance with previous results in the present study. Under a CTC-positive condition, the expression of CK19, CD105 and CD146 in patient and control groups was detected and compared according to the difference between the time point Q1 and the other three time points (Q2-4). The results of the present study suggested that the three biomarkers had no statistical difference in the patient group compared with the control group based on the difference between the Q2 and Q1 time points (data not shown). A significant difference between the difference of expression of CK19 and CD105 between the Q1 and Q3 time points of CTC-positive patient and non-RCC control groups was identified (P<0.05; Table III). However, no significant difference was identified with CD146. On the basis of the difference between Q1 and Q4, the expressions of all three markers exhibited a statistically significant difference (P<0.05; Table IV). Nevertheless, no significant difference was identified in a CTC-negative condition (P>0.05; Table V).

## Discussion

Tendency to invade and transfer is one of the obstacles in the treatment of RCC. Clinical parameters including Fuhrman stage may not entirely reflect prognosis and early metastasis (27). Patients may have a recurrence or metastasis with a low-grade and early-stage tumor following surgery. Thus, oncologists and clinicians produced insight into alternative or complementary indicators, expecting to predict disease progression and early prognosis more accurately (28). The aim of the present study was to explore peripheral blood biomarkers (CK19, CD105 and CD146) for the early metastasis of RCC following surgery.

CK19 is a characteristic intermediate filament of epithelial cells, which has gained attention as a marker for micro-metastasis (29). It has been reported that CK 19 may be detected in the tumor cells of patients with breast cancer (30). Although additional validation studies are essential, the present study demonstrated that CK19 is highly expressed in tumor tissues compared with that in normal tissues, which were confirmed using immunohistochemical staining and western blotting. CD105, a co-receptor of TGF- $\beta$ , is implicated in tumor proliferation, migration and differentiation (31). It has been reported to be a marker of tumor neovascularization, and to provide prognostic data (31). It has been demonstrated that the high expression of CD105 is closely correlated with a more favorable prognosis compared with lower expression (32). However, other studies have demonstrated that the high CD105 expression is positively associated with a poorer progression-free survival (33). In the present study, CD105 expression in tumor tissues was greater compared with that in normal tissues, which was consistent with the latter. CD146 was revealed to participate in cancer progression. It may enhance cancer migration and invasion in melanoma, gallbladder adenocarcinoma and breast cancer (34-36). Furthermore, CD105 had been considered to be an indicator of early metastasis including in gastric cancer and lung cancer (37). However, the expression of CD146 was greater in the intermediate/high grade of oral mucoepidermoid carcinoma compared with that in patients with local recurrence and distant metastasis (38). To the best of our knowledge, the role of CD146 in RCC cells had not yet been explored. In the present study, CD146 expression in tumor tissues was revealed to be higher compared with that in normal tissues, which corresponded with the results of a previous study (37).

In the subsequent experiment, the results indicated that the expressions of CK19 and CD105 had statistically significant differences between two groups (P<0.05), but no such significant difference was identified for CD146. Further experiments revealed no significant difference between four time points (data not shown). Furthermore, CTCs were examined, a crucial blood biomarker in cancer (26). CTC clusters have been reported to be present in patients with metastatic colorectal cancer and prostate cancer, and have a positive correlation with decreased survival in lung cancer and breast cancer (39,40). Furthermore, the CTC clusters may indicate a poor outcome in breast cancer via detection using the CellSearch system (41). In view of the indicated role of CTCs on tumor prognosis and early metastasis, subgroup analysis was performed based on whether the patients were CTC positive or not, and the difference between the point of Q1 and other three points (Q2-4). Similar to the results presented before, the results revealed no statistically significant difference between the CTC-positive or -negative condition, and in the difference between the time points Q1 and Q2 (data not shown). Nevertheless, the expression of CK19 and CD105 exhibited a significant difference with a CTC-positive condition according to the difference between the time points Q1 and Q3 (P<0.05). Furthermore, on the basis of the difference between Q1 and Q4, the expression of CK19, CD105 and CD146 were significantly different (P<0.05). The results concluded that CK19, CD105 and CD146 could not be used individually to predict the early metastasis of patients with RCC subsequent to surgery. The expressions of CK19, CD105, and CD146 at Q2 (1 day post-operation) may not accurately reflect the levels of biomarkers following surgery, due to the state of stress caused by surgical trauma, which resulted in no statistical difference between the CTC-positive condition in the difference between the point of Q1 and Q2. But at the Q3 time point (1 week post-operation), patients had recovered from the surgery. The expression of CK19 and CD105 exhibited a significant difference with a CTC-positive condition based on the difference between Q1 and Q3 (P<0.05), although CD146 did not. Furthermore, a statistically significant difference was present in the expression of CK19, CD105 and CD146 on the basis of the difference between Q1 and Q4 (P<0.05), indicating that CK19, CD105 and CD146 may be novel predictors for the early metastasis of CTC-positive condition in the difference between Q1 and Q3/Q4. Additionally, CK19, CD105 and CD146 may be auxiliary indicators for CTCs to estimate the prognosis of patients with RCC.

Altogether, the results suggest that CK19, CD105 and CD146 markers of peripheral blood may be considered to be effective tools to evaluate the early metastasis of patients with a CTC-positive condition between Q1 and Q3/Q4. CK19, CD105 and CD146 may be useful for CTCs to evaluate the prognosis of patients with RCC, although more widespread studies with larger scale samples are required in order to verify the results of the present study. As a nonsurgical strategy, CK19, CD105 and CD146 may be the potential targets for treatment of RCC.

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## **Competing interests**

The authors declare that they have no competing interests.

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