ORIGINAL RESEARCH

Combination of Hematology Indicators and Oncological Characteristics as a New Promising Prognostic Factor in Localized Clear Cell Renal Cell Carcinoma

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Purpose: This study aimed to construct a predictive model for recurrence and metastasis in patients with localized clear cell renal cell carcinoma (ccRCC) based on multiple preoperative blood indexes and oncological characteristics.

Patients and Methods: Overall, 442 patients with localized ccRCC between 2013 and 2015 were included. Using least absolute shrinkage and selection operator (LASSO) Cox regression analysis, the top three risk factors from the peripheral blood indicators were screened to construct a risk score, and a prognostic model was established. Harrell's concordance index (C-index) was applied to evaluate the predictive accuracy of the model for predicting disease-free survival (DFS) in ccRCC.

Results: Out of 38 blood indexes, the top three predictors were fibrinogen (FIB), C-reactive protein (CRP) and neutrophil-lymphocyte ratio (NLR). The FIB-CRP-NLR (FCN) score (hazard ratio [HR]: 1.86, 95% confidence interval [CI]: 1.21–2.9, P = 0.005) was an independent prognostic factor in multivariate analysis. Furthermore, the FIB-CRP-NLR -T-Grade (FCNTG) risk model combining FCN score, T stage and Furhman grade achieved a higher prognostic accuracy (mean C-index, 0.728) than both the FCN score alone (mean C-index, 0.675) and the stage, size, grade, and necrosis (SSIGN) score (mean C-index, 0.686) in the validation cohort.

Conclusion: The FCN score combining peripheral blood indicators of inflammation and coagulation is an independent prognostic marker of ccRCC. The FCNTG model, which systemically incorporates preoperative blood indexes to oncological characteristics, shows its advantages of convenience and high prediction efficiency.

Keywords: clear cell renal cell carcinoma, prognosis, fibrinogen, C-reactive protein, neutrophil-lymphocyte ratio

Introduction

Renal cell carcinoma (RCC) is the third most common urological cancer, with an estimated more than 70,000 new cases and 14,000 deaths per year.¹ Although most patients with localized clear cell renal cell carcinoma (ccRCC) present as slow-growing tumors, one-quarter of patients experience disease recurrence and metastasis after complete surgical resection.² Clinical staging with tools such as the University of Los Angeles Integrated Staging System (UISS) and the Mayo Clinic Stage, Size, Grade, and Necrosis (SSIGN) score are valuable for

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Figure I Study flowchart. Abbreviation: ccRCC, clear cell renal cell carcinoma.

prognostication.³ However, these models did not take into account of the cancer-related inflammation and coagulation factors, which comprising an essential component of the tumor microenvironment.^{4,5} To date, some studies have evaluated the prognostic role of pre-treatment blood parameters including C-reactive protein

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Variables	Overall	Training Cohort	Validation Cohort	P value
	(n=442)	(n=221)	(n=221)	
Age (years)				0.6
< 60	314 (71.0%)	154 (69.7%)	160 (72.4%)	
≥ 60	128 (29.0%)	67 (30.3%)	61 (27.6%)	
Sex				0.833
Male	315 (71.3%)	156 (70.6%)	159 (71.9%)	
Female	127 (28.7%)	65 (29.4%)	62 (28.1%)	
Diabetes status				0.883
Yes	52 (11.8%)	27 (12.2%)	25 (11.3%)	
No	390 (88.2%)	194 (87.8%)	196 (88.7%)	
BMI (kg/m ²)				0.249
< 25	191 (43.2%)	89 (40.3%)	102 (46.2%)	
≥ 25	251 (56.8%)	I 32 (59.7%)	119 (53.8%)	
Laterality				0.069
Left	200 (45.2%)	110 (49.8%)	90 (40.7%)	
Right	242 (54.8%)	111 (50.2%)	131 (59.3%)	
Tumor size (cm)				0.157
≤ 7	416 (94.1%)	212 (95.9%)	204 (92.3%)	
> 7	26 (5.9%)	9 (4.1%)	17 (7.7%)	

(Continued)

Table I (Continued).

Variables	Overall	Training Cohort	Validation Cohort	P value
	(n=442)	(n=221)	(n=221)	
T stage				0.658
Tla-2b	390 (88.2%)	197 (89.1%)	193 (87.3%)	
T3a	52 (11.8%)	24 (10.9%)	28 (12.7%)	
Fuhrman grade				0.681
Low (GI–2)	305 (69.0%)	150 (67.9%)	155 (70.1%)	
High (G3–4)	137 (31.0%)	71 (32.1%)	66 (29.9%)	
FIB (g/L)				0.119
FIB < 2.77	173 (39.1%)	78 (35.3%)	95 (43.0%)	
FIB ≥ 2.77	269 (60.9%)	143 (64.7%)	126 (57.0%)	
CRP (mg/dL)				0.612
CRP < 0.2	298 (67.4%)	152 (68.8%)	146 (66.1%)	0.012
CRP ≥ 0.2	144 (32.6%)	69 (31.2%)	75 (33.9%)	
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NLR		01 (41 29/)	00 (20 0%)	0.846
NLR < 1.68	179 (40.5%)	91 (41.2%)	88 (39.8%)	
NLR ≥ 1.68	263 (59.5%)	130 (58.8%)	133 (60.2%)	
Basophils(×10 ⁹ /L)				0.065
Mean (SD)	0.0381 (0.0257)	0.0358 (0.0208)	0.0403 (0.0297)	
Median (IQR)	0.030 (0.02, 0.05)	0.030 (0.02, 0.04)	0.040 (0.02, 0.28)	
Albumin (g/L)				0.262
Mean (SD)	43.86 (4.07)	44.07 (3.65)	43.64 (4.44)	
Median (IQR)	44.1 (42.0, 46.1)	44.3 (42.3, 46.1)	43.9 (41.5, 46.1)	
PLR				0.489
Mean (SD)	124.0 (50.5)	122.4 (48.7)	126.7 (52.2)	
Median (IQR)	114.6 (90.5, 145.7)	111.5 (90.9, 144.4)	117.1 (90.3, 145.8)	
LMR				0.309
Mean (SD)	4.59 (1.75)	4.67 (1.75)	4.50 (1.74)	0.001
Median (IQR)	4.44 (3.37, 5.56)	4.56 (3.41, 5.61)	4.21 (3.33, 5.48)	
mGPS				
0	397 (89.8%)	200 (90.5%)	197 (89.1%)	0.753
	39 (8.8%)	18 (8.1%)	21 (9.5%)	0.737
2	6 (1.4%)	3 (1.4%)	3 (1.4%)	I
SIM score				
0	49 (11.1%)	25 (11.3%)	24 (10.9%)	1
I	280 (63.3%)	139 (62.9%)	141 (63.8%)	0.765
2	104 (23.5%)	53 (24.0%)	51 (23.1%)	0.646
3	9 (2.0%)	4 (1.8%)	5 (2.3%)	I
SSIGN score				0.871
≤ 4	400 (90.5%)	201 (91.0%)	199 (90.0%)	
> 4	42 (9.5%)	20 (9.0%)	22 (10.0%)	
ТММ				0.658
I–II	390 (88.2%)	197 (89.1%)	193 (87.3%)	0.000
	52 (11.8%)	24 (10.9%)	28 (12.7%)	

Abbreviations: BMI, body mass index; IQR, interquartile range; SD, standard deviation; PLR, platelet-lymphocyte ratio; LMR, lymphocyte-monocyte ratio; mGPS, modified Glasgow prognostic score; SIM score, systemic inflammatory marker score.



Figure 2 Screening of blood indicators using LASSO regression analysis. The top three important prognostic markers screened by LASSO regression out of 38 blood indexes in the training cohort were FIB, CRP and NLR.

Abbreviations: LASSO, least absolute shrinkage and selection operator; FIB, fibrinogen; CRP, C-reactive protein; NLR, neutrophil-lymphocyte ratio.

(CRP), neutrophil-lymphocyte ratio (NLR), fibrinogen (FIB) and erythrocyte sedimentation rate (ESR).^{6–10} Nevertheless, these studies evaluated only one or two indexes, and the mechanism was unclear.

We have reported that the plasma concentrations of FIB, CRP and neutrophils were positively correlated with the circulating tumor cells (CTCs), which was a prerequisite for recurrence and metastasis.¹¹ In this study, we evaluated the preoperative blood indexes to explore a risk classification for predicting disease-free survival (DFS) in patients with stage I–III ccRCC.

Patients and Methods Patients

This retrospective cohort study collected 712 patients with stage I–III ccRCC who underwent partial or radical nephrectomy at the Cancer Institute and Hospital of the Chinese Academy of Medical Sciences (CAMS) from March 2013 to March 2015. The clinical data, including general demographic factors, preoperative peripheral blood indicators, and clinicopathological parameters were collected.

The inclusion criteria of the study were as follows: (1) no primary cancer of any other organs before the operation; (2) no chronic inflammatory allergic disease; (3) no history of taking anticoagulants such as for cardiovascular or cerebrovascular thrombosis; (4) exact pathological diagnosis of ccRCC; (5) complete resection of the tumor, which was defined as a negative incisal margin; (6) complete clinicopathological characteristics and follow-up data; and (7) no evidence of extrarenal metastasis. Finally, 442 patients with stage I–III ccRCC were enrolled in this study (Figure 1). The study protocol was approved by the Ethics Committee of the Cancer Institute and Hospital of the CAMS, and all patients provided informed consent.

Clinicopathological Data

The clinicopathologic parameters were investigated retrospectively including age, sex, the clinical and histopathological characteristics and preoperative peripheral blood

Table 2 Association of FCN Score with Tumor Pathological Characteristic	s, TNM and SSIGN Score in the Training Cohort
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Variables	FCN Score = 0 (n=37)	FCN Score = 1 (n=75)	FCN Score = 2 (n=62)	FCN Score = 3 (n=47)	P value
< 60	30 (81.1%)	53 (70.7%)	39 (62.9%)	32 (68.1%)	
≥ 60	7 (18.9%)	22 (29.3%)	23 (37.1%)	15 (31.9%)	
Sex					0.495
Male	30 (81.1%)	51 (68.0%)	43 (69.4%)	32 (68.1%)	
Female	7 (18.9%)	24 (32.0%)	19 (30.6%)	15 (31.9%)	
BMI (kg/m ²)					0.755
< 25	15 (40.5%)	33 (44.0%)	25 (40.3%)	16 (34.0%)	
≥ 25	22 (59.5%)	42 (56.0%)	37 (59.7%)	31 (66.0%)	
Laterality					0.55
Left	18 (48.6%)	37 (49.3%)	35 (56.5%)	20 (42.6%)	
Right	19 (51.4%)	38 (50.7%)	27 (43.5%)	27 (57.4%)	
Tumor size(cm)					0.04
≤ 7	37 (100%)	74 (98.7%)	59 (95.2%)	42 (89.4%)	
> 7	0 (0%)	I (I.3%)	3 (4.8%)	5 (10.6%)	
T stage					0.019
TIa-2b	35 (94.6%)	69 (92.0%)	57 (91.9%)	36 (76.6%)	
T3a	2 (5.4%)	6 (8.0%)	5 (8.1%)	11 (23.4%)	
Fuhrman grade					0.233
Low (G1–2)	29 (78.4%)	51 (68.0%)	43 (69.4%)	27 (57.4%)	
High (G3–4)	8 (21.6%)	24 (32.0%)	19 (30.6%)	20 (42.6%)	
FIB (g/L)					<0.001
FIB < 2.77	37 (100%)	38 (50.7%)	3 (4.8%)	0 (0%)	
FIB ≥ 2.77	0 (0%)	37 (49.3%)	59 (95.2%)	47 (100%)	
CRP (mg/dL)					<0.001
CRP < 0.2	37 (100%)	70 (93.3%)	45 (72.6%)	0 (0%)	
CRP ≥ 0.2	0 (0%)	5 (6.7%)	17 (27.4%)	47 (100%)	
NLR					<0.001
NLR < 1.68	37 (100%)	41 (54.7%)	13 (21.0%)	0 (0%)	
NLR ≥ 1.68	0 (0%)	34 (45.3%)	49 (79.0%)	47 (100%)	
SSIGN score					0.002
≤ 4	36 (97.3%)	71 (94.7%)	58 (93.5%)	36 (76.6%)	
> 4	I (2.7%)	4 (5.3%)	4 (6.5%)	11 (23.4%)	
ТММ					0.019
I–II	35 (94.6%)	69 (92.0%)	57 (91.9%)	36 (76.6%)	
III	2 (5.4%)	6 (8.0%)	5 (8.1%)	11 (23.4%)	

indexes (Table 1). The pathological characteristics were evaluated according to the 2010 Fuhrman grading system, and the 2010 American Joint Committee on Cancer TNM classification.¹² Additionally, peripheral blood samples were obtained within 2 weeks before the operation in our hospital.

Follow-Up

All patients were followed up regularly. Follow-up assessments were routinely performed in intervals of 3–6 months for the first 2 years and then once a year thereafter, in accordance with the standard follow-up procedures of our institution. Physical examination,



Figure 3 Multivariate Cox regression analysis in the training cohort. The FCN score and T stage were independent risk factors for DFS in patients with ccRCC. Abbreviations: FCN score, FIB-CRP-NLR score; SIM score, systemic inflammatory marker score; mGPS, modified Glasgow prognostic score; BMI, body mass index.



Figure 4 The FIB-CRP-NLR-T-Grade (FCNTG) risk model combining FCN score, T stage and Furhman grade.

routine laboratory tests, and imaging screenings, such as chest X-ray and abdominal computed tomography (CT), were performed in every follow-up. The endpoint was the DFS time, which was calculated as the interval between surgery and the last follow-up or the date of recurrence, metastasis or death.

Statistical Analysis

The dataset was split into training and validation cohorts with repeated random sampling until there was no significant difference (P value > 0.05) between the two cohorts with respect to all variables (Table 1). The P value was calculated using Welch's *t*-test for

continuous variables and χ^2 test or Fisher's exact test for categorical variables.

The blood indexes were selected by least absolute shrinkage and selection operator (LASSO) Cox regression (R software and "glmnet" package). Then, the optimal cut-off values were determined by X-tile. Furthermore, we evaluated the prognostic accuracy of the risk model using Harrell's concordance index (C-index), which is appropriate for censored data.¹³ All statistical tests were two-sided, a *P* value < 0.05 was statistically significant. Both the multivariable Cox regression model and the C-index were completed with R version 3.6.2. and mean C-index calculation using Stata 14.0 (Stata Corp. Texas, USA).

Results

Patient Demographics

The median follow-up time (interquartile range) was 58.7 months (range 51.7 to 65.1 months). Overall, 58 (13.1%) patients had tumor recurrence, metastasis or death, including 36 patients with distant metastasis, 15 patients with local recurrence and 7 patients died of other causes. In all, 35 (7.9%) patients died due to various reasons during the follow-up period. The time to progress (TTP) which was calculated as the interval between surgery and the date of recurrence or metastasis. The 5-year TTP rates were 88.3% (95% confidence interval [CI]: 84.9–91.1%), and the 5-year DFS rates was 86.9% (95% CI: 83.3–89.7%).

FCN Score, and the Relationship Between FCN Score and Clinical Characteristics

The top three important prognostic markers screened by LASSO Cox regression out of 38 blood indexes were FIB, CRP, and NLR (Figure 2). The optimal cutoff values for FIB, CRP and NLR were 2.77 g/L, 0.2 mg/dL and 1.68, respectively. Thus, patients with low levels of all three indicators, FIB < 2.77 g/L, CRP < 0.2 mg/dL, and NLR < 1.68, were scored as 0. Patients were scored 1, 2, and 3 on the basis of these three indicators, respectively. Higher FCN score was significantly associated with larger tumor size (P = 0.04), more advanced T stage (P = 0.019) and higher SSIGN score (P = 0.002) (Table 2).

Prognostic Risk Model Combining FCN Score, T Stage and Furhman Grade

The multivariate Cox regression analysis in the training cohort further showed that FCN score (hazard ratio

[HR]: 1.86, 95% CI: 1.21–2.9, P = 0.005) and T stage (HR: 1.53, 95% CI: 1.23–1.9, P < 0.001) were independent risk factors for DFS in patients with ccRCC (Figure 3). Additionally, the basophils count, body mass index (BMI), diabetes status, systemic inflammatory marker (SIM) score and modified Glasgow prognostic score (mGPS) calculated according to the cutoffs in the literature,^{14–16} were unindependent risk factors in multivariate analysis (P > 0.05). Therefore, the FIB-CRP-NLR-T-Grade (FCNTG) risk model combining FCN score, T stage and Furhman grade was constructed (Figure 4).

Comparison of Predictive Accuracy Between the FCNTG Model and the FCN Score Alone

The predictive power for prognosis of DFS between the FCNTG model and other variables were compared. The mean C-index of the FCNTG model was 0.728 (95% CI: 0.625–0.834), which was significantly higher than that of the single FCN score (mean C-index: 0.675, 95% CI: 0.574–0.776) (Table 3; Figure 5A).

Comparison of Predictive Accuracy Between the FCNTG Model and the SSIGN Score

As shown in Table 3, the C-index of the FCNTG model for predicting DFS in the validation cohort was 0.728 (95% CI, 0.625–0.834), which was significantly higher than that of the SSIGN score (0.686, 95% CI: 0.582–0.789). The calibration curve showed good agreement between prediction and observation in the probability of 3-year DFS (Figure 5B) and 5-year DFS (Figure 5C). In the overall data, the 5-year DFS rates of patients with low-, intermediate-, and high-risk kidney cancer grouped by the

Table 3 Concordance Index Analysis of the Prognostic Accuracy
of FCNTG and Other Variables for DFS in Indicated Sets

C-Index (95% CI)	Training Cohort	Validation Cohort
	(n=221)	(n=221)
FCNTG	0.791 (0.724–0.857)	0.728 (0.625–0.834)
FCN score	0.723 (0.650–0.795)	0.675 (0.574–0.776)
SSIGN	0.685 (0.595–0.775)	0.686 (0.582–0.789)

Abbreviations: CI, confidence interval; C-index, concordance index; SSIGN, stage, size, grade, and necrosis.



Figure 5 (A) Compare the FCNTG model with other variables using C-index. The prediction accuracy of the FCNTG model was higher than that of the single FCN score, and the FCNTG model showed superiority in assessing risk of recurrence compared to the SSIGN score. (B) The calibration curve for predicting patient DFS at 3 years in the validation cohort. (C) The calibration curve for predicting patient DFS at 5 years in the validation cohort. Abbreviations: FCNTG, FIB-CRP-NLR-T-Grade; C-index, Harrell's concordance index; DFS, disease-free survival.



Strata + Low-risk + Intermediate-risk + High-risk

Figure 6 Kaplan-Meier analysis in the overall data. The 5-year DFS rates of patients with low-, intermediate-, and high-risk renal carcinoma grouped by the FCNTG model were 97%, 84% and 61%, respectively (*P* < 0.0001).

Abbreviations: FCNTG, FIB-CRP-NLR-T-Grade; DFS, disease-free survival.

FCNTG model were 97%, 84% and 61%, respectively (P < 0.0001) (Figure 6).

Discussion

As malignant tumors are the result of genetic mutations, it is necessary to comprehensively consider the immune status of the body and the biological characteristics of the tumor to improve the accuracy of prognostic predictions for malignant tumors.^{17,18} The commonly used SSIGN score and UISS exclude the hematological indicators that reflect the state of the body, which may lead to their limited predictive effectiveness.¹⁹ Some studies have reported the predictive value of hematology indicators, such as basophil, SIM score, and mGPS which is the incorporation of CRP and albumin levels.^{14,16,20-22} Furthermore, The FIB, D-dimers, and ESR are all related to the poor prognosis of patients with RCC.7,9,23,24 However, it is unclear which indicator or set of indicators provide the greatest accuracy for prognosis, and the mechanism has not been clarified.

Machine learning methods can detect more effective prognostic factors that are difficult to identify from the complex combination of multiple parameters.²⁵ Therefore, in this study, we applied a Cox proportional hazards model using LASSO penalization to screen 38 blood indexes; this

approach favored the selection of blood indicators with powerful prognostic value, and the three strongest blood indexes, FIB, CRP, and NLR, were all related to coagulation and inflammation status. It has been suggested that FIB, an extracellular matrix component, promotes the growth of cancer cells in the tumor microenvironment by binding to fibroblast growth factor 2 and vascular endothelial growth factor.²⁶ The CRP concentration may be elevated owing to hepatic CRP synthesis, which is stimulated by cancer cellderived inflammatory cytokines.²⁷ Neutrophils may also contribute to tumor development and progression by providing an adequate tumor microenvironment via the production of cytokines and chemokines.²⁸ Our previous research also pointed out that the abnormality of inflammation and coagulation indexes is correlated with viable CTCs, which is the premise of tumor recurrence and metastasis, and the neutrophil extracellular traps (NETs) induced by circulating neutrophils could protect the survival of circulating tumor cells, thus promoting postoperative recurrence and metastasis in patients with RCC.¹¹ Therefore, it is reasonable to construct a prognostic model for local renal cell carcinoma based on the combination of hematological indicators FIB, CRP, and NLR.

In this study, although the basophils count, BMI, diabetes status, SIM score and mGPS were positively

associated with the risk of recurrence and metastasis in multivariate Cox regression analysis, but these indicators were not independent prognostic risk factors. Finally, we constructed an FCNTG model to predict the prognosis of patients with localized ccRCC which combined peripheral blood indicators of inflammation and coagulation and oncology staging information. The predictive effectiveness of the FCNTG model (C-index, 0.728) was higher than that of the SSIGN score (C-index, 0.686), and the 5-year DFS rates of the patients with low-, intermediate-, and high-risk kidney cancer grouped by the FCNTG model were 97%, 84%, and 61%, respectively. Therefore, the FCNTG model combining peripheral blood markers and tumor stage was consistent with the idea that integrating multiple markers may provide higher accuracy,²⁹ and the model showed the advantages of convenience and high prediction efficiency.

The limitations of this study are as follows: first, the inherent bias associated with its retrospective design. Second, the number of cases is relatively small. In the future, we hope to accumulate more cases and carry out multicenter and prospective studies to further verify the predictive performance of the model.

In conclusion, we found the FIB, CRP, and NLR indexes could reflect the antitumor status of patients with kidney cancer. Furthermore, the FCNTG model, which combines inflammatory and coagulation indicators with tumor stage and grade information, improves the accuracy of prognostic predictions for patients with ccRCC.

Abbreviations

RCC, renal cell carcinoma; ccRCC, clear cell renal cell carcinoma; BMI, body mass index; CI, confidence interval; C-index, concordance index; SSIGN, stage, size, grade, and necrosis; DFS, disease-free survival; CTCs, circulating tumor cells; WBC, white blood cell count; Hb, hemoglobin; PLT, platelet count; PDW, platelet distribution width; PCT, platelet hematocrit, HCT, hematocrit; NEUT, neutrophil count; NETU%, neutrophil percentage; LYMPH, lymphocyte count; MONO, monocyte count; MONO%, monocyte percentage; NLR, neutrophil-lymphocyte ratio; EOS, eosinophils count; BAS, basophils count; AST, aspartate aminotransferase; ALT, alanine aminotransferase; ALP, alkaline phosphatase; GGT, glutamyl transferase; LDH, lactate dehydrogenase; TBIL, total bilirubin; TP, total protein; Alb, albumin; URIC, uric acid; BUN, blood urea nitrogen; CRE, creatinine; IgA, immunoglobulin A; IgG, immunoglobulin G; IgM, immunoglobulin M; CRP, C-reactive protein; PT, prothrombin time; APTT, activated partial thromboplastin time; FIB, fibrinogen; PLR, platelet-lymphocyte ratio; FAR, fibrinogen-albumin ratio; ESR, erythrocyte sedimentation rate; LMR, lymphocyte-monocyte ratio; SIM score, systemic inflammatory marker score; mGPS, modified Glasgow prognostic score.

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Disclosure

The authors report no conflicts of interest in this work.

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