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Nomogram for the prediction of crescent formation in IgA nephropathy patients: a retrospective study



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Abstract

Background The 2017 Oxford classification of immunoglobulin A nephropathy (IgAN) recently reported that crescents could predict a worse renal outcome. Early prediction of crescent formation can help physicians determine the appropriate intervention, and thus, improve the outcomes. Therefore, we aimed to establish a nomogram model for the prediction of crescent formation in IgA nephropathy patients.

Methods We retrospectively analyzed 200 cases of biopsy-proven IgAN patients. Least absolute shrinkage and selection operator(LASSO) regression and multivariate logistic regression was applied to screen for influencing factors of crescent formation in IgAN patients. The performance of the proposed nomogram was evaluated based on Harrell's concordance index (C-index), calibration plot, and decision curve analysis.

Results Multivariate logistic analysis showed that urinary protein ≥ 1 g (OR = 3.129, 95%CI = 1.454–6.732), urinary red blood cell (URBC) counts \geq 30/ul (OR = 3.190, 95%CI = 1.590–6.402), mALBU \geq 1500 mg/L(OR = 2.330, 95%CI = 1.008–5.386), eGFR < 60ml/min/1.73m²(OR = 2.295, 95%CI = 1.016–5.187), Serum IgA/C3 ratio \geq 2.59 (OR = 2.505, 95%CI = 1.241–5.057), were independent risk factors for crescent formation. Incorporating these factors, our model achieved well-fitted calibration curves and a good C-index of 0.776 (95%CI [0.711–0.840]) in predicting crescent formation.

Conclusions Our nomogram showed good calibration and was effective in predicting crescent formation risk in IgAN patients.

Keywords IgA nephropathy, Crescent, Prediction, Nomogram

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Introduction

IgA nephropathy (IgAN) is the most common form of primary glomerulonephritis. It has been reported that 25–40% of patients will develop end-stage renal disease (ESRD) within 10–20 years [1]. Due to the clinical and pathological diversity of IgAN,this disease progression and prognosis varies interindividually [2, 3]. The pathological features of IgAN is highly variable, such as renal active lesions(i.e., crescent formation) and chronic glomerular pathology features(i.e., tubular-interstitial fibrosis).

Crescent formation is a common histopathological finding, occurring in approximately 20-60% of IgAN patients [4]. Although the original Oxford study and several subsequent validation studies did not find the predictive value of crescents [5, 6], an increasing number of studies have found that crescents to be an independent predictor of poor renal outcomes in patients with IgAN [7, 8]. Recently, a multicenter study conducted by Haas and colleagues evaluated the effect of crescents on renal function progression [9]. Based on 3,096 IgAN patients from four retrospective studies across the world, crescents were found to be strong predictors of unfavorable prognosis of IgAN. Consequently, cellular and/or fibrocellular crescents (C) were include in the updated Oxford classification of IgAN:C0(no crescents), C1(crescents in <25% of glomeruli), and C2(crescents in \geq 25% of glomeruli) [10].

Considering that the glomerular crescent is highly associated with poor renal prognosis of IgA nephropathy, early diagnosis and intervention of crescent is necessary. Although renal biopsy is the gold standard for the diagnosis of IgAN, it is an invasive operation that is prone to complications such as bleeding and infection, which is not acceptable for some patients [11, 12]. In addition, repeat biopsies in IgAN are still remain rare and it is sometimes difficult to judge the acute and chronic renal lesions of an individual patient, which makes it difficult to use the time-to-event data of the population to guide treatment. Hence, it is clinically necessary to establish a safe and non-invasive diagnostic method in biopsyproven IgAN patients to predict the risk of developing crescents for as an alternative to re-biopsy. However, the individualized prediction of IgA nephropathy with crescent has been rarely reported and should be urgently solved. Therefore, the purpose of this study was to examine influencing factors of crescent formation and construct a nomogram for predicting the incidence of the crescent in IgAN patients.

Methods

Study design and population

This retrospective cohort study included biopsy-proven IgAN patients hospitalized in Guangdong Provincial Hospital of Chinese Medicine between May 2005 and November 2017. The inclusion criteria were: (1) 18 years of age or older; (2) patients with biopsy-proven primary IgAN; (3) With less than eight glomeruli on light microscopy. The exclusion criteria were: (1) insufficient clinical and pathological data; (2) atypical IgA nephropathy, such as crescentic GN; (3) Individuals diagnosed secondary IgAN such as rheumatology disease, Henoch–Schonlein purpura.or liver cirrhosis.

Ethical approval was granted by the Ethics Committee of Guangdong Provincial Hospital of Chinese Medicine (B2016-155-01). Informed consent was exempted since the study only involved analysis of anonymized existing data and records.

Data collection and definition

In this cohort, we recorded demographic variables and laboratory indicators during kidney biopsy, including gender, body mass index(BMI), 24 h urinary protein excretion, immunoglobulin A and complement 3 (C3). eGFR was calculated by the CKD-EPI equation [13]. MAP was equal to diastolic BP+1/3(systolic BP - diastolic BP) [14]. Medications that block the renin-angiotensin system(RAAS) and immunosuppression(mainly steroid therapy, and other drugs) were recorded after kidney biopsy.

The 2017 updated Oxford Classification (MEST-C) was used in this study [10]: mesangial hypercellularity (M):M1 was defined as a mesangial hypercellularity score>0.5. Endocapillary hypercellularity (E): E1 was defined as the presence of endocapillary hypercellularity. Segmental glomerulosclerosis (S):S1 was defined as the presence of segmental glomerulosclerosis. Tubular atrophy/interstitial fibrosis (T):T1 was defined as tubular atrophy/interstitial fibrosis within 26–50% of the cortical area, and T2 was defined as tubular atrophy/interstitial fibrosis in >50% of the cortical area. Crescent(C):C1 was defined as crescents in <1/4 of all glomeruli.

Statistical analysis

Missing data was imputed using multiple imputation with chained equations to generate 5 complete datasets, with results combined across imputed datasets. We have adopted the existing sample size criteria by Riley et al. [15] and pmsampsize R library [16, 17] to calculate the sample size. On the basis of a study by Lee M J et al. [18], it was expected that the prevalence of crescents in IgAN patients is 20%. We estimated a C-index of 0.8 for this prediction model and planned to collect 22 predictor variables. To ensure an accurate estimate of overall risk (model intercept), at least 246 subjects were required. However, since our study was a single-center retrospective study and the incidence of IgAN patients at the

Table 1 Clinical and histological characteristics

Variables	Total (n = 200)	C0 group (n = 118)	C≥1 group (n=82)	P- value
Female ,n(%)	106(53)	59(50)	47(57.3)	0.308
Age (years)	32(26-42)	32.5(25.7-42)	31(26–46)	0.563
MAP (mmHg)	94.67(87.33-104.58)	92.83(86.67-100.67)	97.67(88.33-107.42)	0.009
BMI(kg/m ²)	22(19.82-24.61)	21.79(19.61-24.52)	22.48(20.25-24.80)	0.519
Urinary protein (g/24 h) (%)				< 0.001
<1g	108(54)	80(67.8)	28(34.1)	
≥1g	92(46)	38(32.2)	54(65.9)	
URBC (counts /ul) (%)				0.003
< 30	78(39)	56(47.5)	22(26.8)	
≥30	122(61)	62(52.5)	60(73.2)	
eGFR (ml/min/1.73m ²)(%)				0.017
<60	158(79)	100(84.7)	58(70.7)	
≥60	42(21)	18(15.3)	24(29.3)	
UA(mmol/L)	399.00±107.16	390.09±101.11	411.84±114.72	0.159
TC(mmol/L)	4.73(4.13-5.55)	4.64(4.02-5.35)	5.15(4.22-5.80)	0.029
TG(mmol/L)	1.38(0.91-1.95)	1.25(0.83-1.86)	1.57(0.98-2.09)	0.069
TP(g/L)	66.15(60.2-70.9)	67.4(62.35-71.68)	64.71(57.18-69.93)	0.065
ALB (g/L)	40.05(36.13-42.9)	40.3(37.13-43.13)	38.2(33.1-42.65)	0.039
BUN(mmol/L)	5.15(4.28-6.40)	4.94(4.35-5.95)	5.41(4.15-7.29)	0.072
LDL(mmol/L)	3.01(2.48-3.69)	2.82(2.42-3.55)	3.26(2.63-3.82)	0.016
HDL(mmol/L)	1.19(0.95-1.49)	1.21(0.95-1.45)	1.17(1.01-1.57)	0.915
Serum IgA/C3 (%)				0.024
< 2.59	72(36)	50(42.4)	22(26.8)	
≥2.59	128(64)	68(57.6)	60(73.2)	
lgGU(mg/L)	50.65(21.75-123.25)	33.05(15.78-83.80)	80.40(35.78-207.75)	< 0.001
β2-MgU (mg/L)	0.23(0.22-0.40)	0.23(0.22-0.32)	0.23(0.22-0.73)	0.152
mALBU (mg/L)(%)				< 0.001
< 1500	147(73.5)	101(85.6)	46(56.1)	
≥1500	53(26.5)	17(14.4)	36(43.9)	
a1-MgU (mg/L)	10.02(5.56-20.9)	9.09(5.56-17.33)	12.80(6.29–28.63)	0.005
a2-MgU(mg/L)	2.47(2.37-2.66)	2.41(2.39-2.55)	2.55(2.34-5.32)	0.023
TrfU(mg/L)	29.45(11.25-74.53)	17.50(8.47-48.48)	48.9(20-131.50)	< 0.001
Oxford classification				
M1%	181(90.5)	100(84.7)	81(98.8)	0.001
E1%	33(16.5)	7(5.9)	26(31.7)	< 0.001
S1%	118(59)	63(53.4)	55(67.1)	0.053
T1+2%	60(30)	23(19.5)	37(45.1)	< 0.001
Treatment				
With RAAS blockade	133(66.5)	80(67.8)	53(64.6)	0.641
With immunosuppression	23(11.5)	9(7.6)	14(17.1)	0.039

Values are expressed as mean ± SD, medians (interquartile ranges), or percentages.

MAP, mean arterial pressure; BMI, body mass index; URBC, urinary red blood cell; eGFR, estimated glomerular filtration rate; BUN, Blood Urea Nitrogen; LDL, Lowdensity Lipoprotein; HDL, High-density Lipoprotein; IgGU, urine immunoglobulin G; β2-MgU, urine β2-microglobulin; mALBU, urine micro-albumin; α1-MgU, urine α1-microglobulin; α2-MgU, urine α2-microglobulin; TrfU, Urinary transferrin.

center was not high, the sample size was derived on the basis of the available data. Pearson's test and Chi-square test were used to test the distribution of categorical variables; the Mann-Whitney U test and paired-sample t-test were used for continuous variables. LASSO regression was used to screen for optimal risk factors of crescent formation in IgA patients. Stepwise logistic regression model was used build a prediction model by introducing the features selected in the LASSO regression model and the β regression coefficient, odds ratio (OR), and P-value were calculated. Additionally, the forest plot was drawn to describe the P-value, OR and 95% CI of selected validation visually. A nomogram was constructed to predict the possibility of crescents formation based on statistically significant factors identified by the multivariate logistic regression model. The predictive performance



Fig. 1 Variables selection using the LASSO binary logistic regression model. (a) Optimal parameter (lambda) selection in the LASSO model used 5-fold cross-validation via minimum criteria. The partial likelihood deviance (binomial deviance) curve was plotted versus log (lambda). Dotted vertical lines were drawn at the optimal values by using the minimum criteria and the 1 SE of the minimum criteria. (b) LASSO coefficient profiles of the 22 variables. A coefficient profile plot was produced against the log (lambda) sequence. 5 variables with nonzero coefficients were selected by optimal lambda

of the nomogram was evaluated using Harrell's concordance index (C-index), and calibration with 1000 bootstrap samples was performed to decrease the overfit bias. The predictive accuracy and diagnostic performance of the nomogram were quantified using the areas under the ROC curves. Decision curve analysis (DCA) was used to determine the clinical practicability of nomograms based on the net benefit according to different threshold probabilities in IgAN patients. Statistical analyses were conducted using SPSS 22.0 (IBM Corp, Armonk, NY, USA)

Table 2	Multivariate	logistic rec	ression	analy	vsis
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Variables	Univariable		Multivariable	
	OR (95% CI)	P value	OR (95% CI)	P-
				value
Female	1.343(0.762-2.368)	0.308		
Age (years)	1.013(0.988–1.039)	0.321		
MAP (mmHg)	1.028(1.005-1.052)	0.017		
BMI(kg/m ²)	1.002(0.922-1.088)	0.965		
Urinary protein≥1 g (g/24 h)	4.060(2.233-7.382)	< 0.001	3.129(1.454–6.732)	0.004
URBC counts≥30/ul	2.463(1.342-4.523)	0.004	3.190(1.590-6.402)	0.001
eGFR<60ml/min/1.73m ²	2.299(1.151-4.590)	0.018	2.295(1.016-5.187)	0.046
UA(mmol/L)	1.002(0.999–1.005)	0.159		
TC(mmol/L)	1.151(0.960–1.379)	0.128		
TG(mmol/L)	1.176(0.921–1.501)	0.194		
TP(g/L)	0.979(0.949-1.010)	0.176		
ALB (g/L)	0.962(0.922-1.002)	0.065		
BUN(mmol/L)	1.228(1.063-1.419)	0.005		
LDL(mmol/L)	1.143(0.920-1.420)	0.226		
HDL(mmol/L)	1.024(0.492-2.132)	0.950		
Serum IgA/C3≥2.59	2.005(1.090-3.690)	0.025	2.505(1.241-5.057)	0.010
lgGU(mg/L)	1.005(1.002-1.007)	0.002		
β2-MgU(mg/L)	1.229(1.027-1.471)	0.025		
mALBU≥1500 mg/L	4.650(2.370-9.122)	< 0.001	2.330(1.008-5.386)	0.048
a1-MgU(mg/L)	1.019(1.003-1.035)	0.021		
a2-MgU(mg/L)	1.074(1.006–1.146)	0.032		
TrfU(mg/L)	1.002(0.999-1.005)	0.171		

MAP, mean arterial pressure; BMI, body mass index; URBC, urinary red blood cell; eGFR, estimated glomerular filtration rate; BUN, Blood Urea Nitrogen; LDL, Lowdensity Lipoprotein; HDL, High-density Lipoprotein; IgGU, urine immunoglobulin G; β2-MgU, urine β2-microglobulin; mALBU, urine micro-albumin; α1-MgU, urine α1-microglobulin; α2-MgU, urine α2-microglobulin; TrfU, Urinary transferrin. and the R software (version 3.6.2; https://www.R-project. org). All reported p values were single-sided; statistical significance was considered at p < 0.05.

Results

Characteristics of cohort

All 200 IgA nephropathy patients were divided into the C0 group (118 cases) and C \geq 1 group (82 cases) according to the presence or absence of a crescent. Among these participants, there were 106 females (53%) and 94 males (47%), with a median age of 32 (26-42) years and MAP of 94.67 (87.33-104.58) mmHg. The C≥1 group was characterized by a significantly higher rate of urinary protein ≥ 1 g, URBC counts $\geq 30/ul$, eGFR<60ml/min/1.73m² and serum IgA/C3 ratio \geq 2.59. As for urinary protein components, patients in the $C \ge 1$ group had higher levels of IgGU[80.40(35.78-207.75) vs. 33.05(15.78–83.8), p<0.001], α1-MgU[12.8(6.29–28.63) vs. 9.09(5.56–17.33), p=0.005], α 2-MgU[2.55(2.34–5.32) vs. 2.41(2.39-2.55), p=0.023], TrfU[48.9(20-131.5) vs. 17.5(8.47-48.48), p=0.023], and higher proportion of mALBU≥1500 mg/L(43.9% vs. 14.4%, p<0.001). However, indicators as sex, BMI, TG, UA presented no differences between the two groups (P > 0.05). (Table 1).

As for MEST-C score, patients in the C \geq 1 group had the higher the proportion of M1 (98.8% vs. 84.7%, p=0.001), E1 (31.7% vs. 5.9%, p<0.001), and T1+2 (45.1% vs. 19.5%, p<0.001). Immunosuppression were more commonly used in the C \geq 1 group than in the C0 group (17.1% vs. 7.6%, p=0.039).

Characteristics selection

In terms of clinical features, we used LASSO regression to identify the main variables related to crescent formation in IgAN patients, and a total of 5 variables were screened out of 22 features. These variables included urinary protein \geq 1 g, URBC counts \geq 30/ul, mALBU \geq 1500 mg/L, eGFR<60ml/min/1.73m² and serum IgA/C3 ratio \geq 2.59 (Fig. 1a and b).

Next, candidate predictors variables selected by LASSO regression were included in stepwise logistic regression analysis. The stepwise analysis showed that urinary protein ≥ 1 g (OR=3.129; 95%CI [1.454–6.732]; P=0.004), serum IgA/C3 ratio ≥ 2.59 (OR=2.505; 95%CI [1.241–5.057]; P=0.010), URBC counts $\geq 30/ul$ (OR=3.190; 95%CI [1.590–6.402]; P=0.001), mALBU ≥ 1500 mg/L (OR=2.330; 95%CI [1.008–5.386]; P=0.048) and eGFR<60ml/min/1.73m² (OR=2.295; 95%CI [1.016–5.187]; P=0.046) were associated with incident crescent formation (Table 2). Based on LASSO regression and stepwise logistic regression to exclude overfitting and multicollinearity, five factors were included in the prediction model, and the forest plot summary is shown in Fig. 2.

Development of an individualized prediction model

A model containing the above independent predictors was developed and presented as a nomogram (Fig. 3). The scores of different variables were obtained on the vertical line on the nomogram, after which the total risk score was calculated by adding all the scores of all variables.



Fig. 2 The forest plot of the OR of the selected variables. Forest plot for outcome in LASSO regression model and multivariate logistic regression analysis. URBC counts, Urinary red blood cell counts; eGFR, Estimated glomerular filtration rate; mALBU, urine micro-albumin



Fig. 3 Nomogram for predicting of crescent formation in IgA nephropathy patients. The crescent formation nomogram was developed in the cohort, with five variables, urinary protein, URBC counts, mALBU, eGFR and serum IgA/C3. Abbreviations: URBC counts, Urinary red blood cell counts; eGFR, Estimated glomerular filtration rate; mALBU, urine micro-albumin

The probability of crescent formation could be directly read on the total point axis.

Validation of prediction model

The calibration plot of the model demonstrated that high consistence between prediction of crescent formation and actual observation (Fig. 4). According to ROC curve, the AUC value was 0.776 (95%CI 0.622–0.788) (Fig. 5), indicated that the model has medium discrimination. The C-index of the prediction model was 0.776 (95%CI 0.711–0.840). We further performed internal validation on the nomogram by bootstrapping validation, and calculated revised C-index as 0.755, which indicated that the model had a relatively great predictive discrimination.

Clinical use of the nomogram

In addition to ROC analysis, DCA has been increasingly used to demonstrate the clinical efficacy in clinical models. The DCA for the crescent formation incidence risk nomogram is shown in Fig. 6. The decision curve showed that using the nomogram model to predict the occurrence of crescent formation had a higher net income at a high-risk threshold between 1 and 80%.

Discussion

Crescents are lesions commonly identified by renal biopsy in IgAN patients, which indicate active and severe glomerular injury. Patients with crescents were found to have more clinicopathologic risk factors for poor prognosis of IgAN [19, 20]. In this study, we established a new tool for predicting the risk of crescent formation in IgA nephropathy to guide clinical management. The nomogram is an intuitive graphical prediction model that can be used to predict certain clinical outcomes or rates of adverse events [21]. The C-index of internal verification showed that the model had a good discrimination and calibration capabilities [22]. In addition, we analyzed ROC curves to validate this nomogram model and evaluated the clinical application value of the prediction tool through DCA, which suggested the satisfactory performance of our model. The nomogram suggests that urinary protein ≥ 1 g, URBC counts $\geq 30/$ ul, mALBU≥1500 mg/L, eGFR<60ml/min/1.73m², and serum IgA/C3 ratio≥2.59 may be used as the independent factors that increase the risk of crescent formation.

In this study, we found that urinary protein ≥ 1 g was an independent predictor of crescent formation in patients



Fig. 4 Calibration curve for the crescent formation nomogram prediction in the cohort. The x-axis represents the predicted crescent formation risk. The y-axis represents the actual diagnosed crescent formation. The diagonal dotted line represents a perfect prediction by an ideal model. The solid line represents the performance of the nomogram, of which a closer fit to the diagonal dotted line represents a better prediction

with IgAN, which is similar to some previous studies [23, 24]. Zhang et al. [25] retrospectively analyzed 538 IgAN patients with different proportions of crescents and found that the increasing crescent proportion was associated with an increased amount of urine protein excretion. The above results indicated a synergistic effect between proteinuria and crescent formation. An earlier study found that extensive changes in podocytes were observed during the formation of crescents [26]. Podocytes further induced the proliferation of parietal epithelial cells by adhering to the glomerular basement membrane and parietal basement membrane to form cell crescent, resulting in the damage of the glomerular filtration membrane and proteinuria [27, 28].

Microscopic hematuria is one of the common clinical manifestations of IgAN. Previous studies have found that a large amount of microscopic hematuria could indicate active renal inflammation and thus may be used as a surrogate marker of crescent [29, 30]. Our results indicated that URBC counts \geq 30/ul were an independent risk factor for crescent formation in IgAN patients. Nagai et al. [31] found that the possibility of crescents in the severe hematuria group (URBCs \geq 30/HPF) was significantly increased (OR=4.3, 95%CI 1.7–10.9). In addition, in vitro studies confirmed that glomerular vascular injury and

GBM breaks cause plasma leakage, which triggers cellular and fibrocellular crescent formation [32].

Our results showed that serum IgA/C3 ratio \geq 2.59 is remarkably related to crescent formation. Previous studies have shown that the serum IgA/C3 ratio is an auxiliary diagnostic marker of IgAN, which can predict the severity of clinicopathology and prognosis of IgAN patients [33, 34]. Kawasaki et al. [35] reported that crescents and mesangial hypercellularity score in IgAN patients with a high serum IgA/C3 ratio and strong glomerular C3 staining were higher than those in IgAN patients with a low serum IgA/C3 ratio and weak glomerular C3 staining. It has been speculated that there is complement during the formation of IgAN crescents and that serum IgA/C3 ratio reflects the degree of pathological kidney damage. Moreover, Itami et al. [36] found that the average immunofluorescence scores of renal IgA, MASP2, and kappa of IgAN patients with crescent were significantly higher than those without crescent; it has been suggested that complement participates in the formation of crescents after activation through the MBL pathway. However, this value may only apply to IgAN patients from this cohort and requires external validation in other ethnic groups, or in countries with multi-ethnic populations or different biopsy practices.



Fig. 5 ROC curve for the crescent formation nomogram prediction in the cohort. ROC curve. It showed that the AUC of this model for predicting crescent formation was 0.776. The optimal cut-off value of the ROC curve was 0.385, corresponding to a specificity and sensitivity of 0.622 and 0.788, respectively

In this prediction model, $eGFR < 60 ml/min/1.73 m^2$ is recognized as the important risk factor for crescents formation. Shao et al. [37] performed a meta-analysis to evaluate the clinical and prognostic significance of crescent formation and found that IgAN patients with crescents exhibit lower eGFR levels. Sun et al. [38] reported that an increasing fraction of glomeruli with crescents was associated with a reduced eGFR. The reason for this phenomenon might be that the impact of crescents on nephron depends on the feature and proportion of the crescent [39]. The fading of single nephron glomerular filtration rate was initiated by the gradually increasing counter pressure in Bowman's capsule and may endorse the collapse of the glomerular tuft. Futhermore, once a glomerular crescent involves and obstructs the urinary pole, the entire nephron no longer contributes to total GFR [40].

In addition, Our results demonstrated that IgAN patients have a higher risk of crescent formation when they present with mALBU≥1500 mg/L. Microalbuminuria as an early protein detection index has been widely carried out and applied to the early assessment of renal damage, especially in the early evaluation of diabetic nephropathy and hypertensive kidney damage [41, 42]. A cross-sectional study of Wu et al. [43] found urinary protein components such as urinary ALB, IgG and Trf can reflect severity of pathological lesions in IgAN. Guo et al. [44] further confirmed that crescent is a risk factor to affect prognosis of IgAN patients with microalbuminuria. This strong correlation between them suggests that we should pay attention to IgAN patients with microalbuminuria in our clinical work. Routine screening for microalbuminuria may help to identify those with crescent formation in whom early therapeutic interventions could reduce the risk of developing progressive renal disease.

There are several limitations in the present study. First, the number of cases investigated was small since it was a single-center study and the cohort was not representative of all Chinese patients with IgAN. In the future, we will expand the number of patients in subsequent studies to analyze the prediction model. Although the sample size of this cohort was relatively small, this prediction model still is expected to provide reliable estimates. The reasons are as follows:1) Multivariate regression was carried out only after the independent variables were screened. Since



Fig. 6 Decision curve analysis (DCA) for the crescent formation nomogram. DCA. The y-axis measures the net benefit. The red solid line represents the crescent formation nomogram. The solid grey line represents the assumption that all patients with IgA nephropathy developed crescent formation. The solid black line represents the assumption that none of the patients with IgA nephropathy developed crescent formation. The decision analysis curve showed that the net benefit rate was >0 at the high-risk threshold of 1–80%, which was clinically significant

there were many variables and few samples, we used lasso regression to screen the sample size. 2) The OR values and confidence intervals of the results are relatively normal. 3) The goodness of fit results of the model show that the model modeling is successful(Cox-Snell $R^2=0.218$, $R^2_{Nagelkerke} = 0.293$). Second, the analysis of risk factors did not include all potential factors affecting the formation of crescents, such as drug treatment, inflammatory cytokines, and other factors, were not thoroughly evaluated. Third, the nomogram lacks external validation in other IgAN populations from other regions and countries and may require further external validation by a multicenter sample study.

Conclusions

Based on the five risk factors of urinary protein ≥ 1 g, URBC counts $\geq 30/ul$, mALBU ≥ 1500 mg/L, eGFR < 60 ml/min/1.73m², and serum IgA/C3 ratio ≥ 2.59 , a nomogram model was constructed to predict the risk of crescent formation in IgAN patients. The model had high accuracy, discrimination, and predictive ability, indicating

its potential practicability for high-risk patients' clinical screening and medical intervention.

List of abbreviations

IgAN	IgA nephropathy
ESRD	End-stage renal disease
MAP	Mean arterial pressure
URBC	Urinary red blood cell
eGFR	Estimated glomerular filtration rate
BP	Blood pressure
LDL	Low-density Lipoprotein
HDL	High-density Lipoprotein
lgA	serum immunoglobulin A
C3	serum complement 3
lgGU	urine immunoglobulin G, β2-MgU:urine β2-microglobulin
mALBU	urine micro-albumin
a1-MgU	urine a1-microglobulin
a2-MgU	urine a2-microglobulin
TrfU	Urinary transferrin
М	Mesangial hypercellularity
E	endocapillary hypercellularity
S	Segmental glomerulosclerosis
Т	Tubular atrophy/interstitial fibrosis
С	Crescents
C-index	Harrell's concordance index
AUC	Area under the curve

Acknowledgements

We wish to thank the department of nephrology of Guangdong hospital of traditional Chinese medicine and Beijing University of Chinese Medicine Shenzhen Hospital for their great support of this project.

Authors' contributions

C.Z. and Q.T. made substantial contributions to the study concept and design. Z.L. and L.F. were in charge of the manuscript draft and manuscript revision. H.Z., and X.L. took responsibility for obtaining ethical approval, collecting samples, and confirming data accuracy. Q.L., F.L., L.W., J.M., P.F., and X.L. helped with implementation. All authors discussed, read, revised the manuscript, and gave final approval for publication.

Funding

The study was funded by the Research Project for Practice Development of National TCM Clinical Research Bases (Project no. JDZX2015202). The funder was not involved in the study design, collection, analysis, interpretation of data, the writing of this article or the decision to submit it for publication.

Data availability

The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

Declarations

Competing interests

The authors declare no competing interests.

Ethics approval and consent to participate

Ethical approval was granted by the Ethics Committee of Guangdong Provincial Hospital of Chinese Medicine (B2016-155-01). All methods were carried out in accordance with relevant guidelines and regulations. The Ethics Committee of Guangdong Provincial Hospital of Chinese Medicine informed consent was exempted since the study only involved analysis of anonymized existing data and records (Application Form for Exemption from Signed Consent Application Form, Version : 001/20160620).

Consent for publication

Not applicable.

Received: 15 July 2022 / Accepted: 25 August 2023 Published online: 04 September 2023

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