

Development and Validation of a Prediction Rule Using the Oxford Classification in IgA Nephropathy

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Summary

Background and objectives The risk assessment for developing ESRD remains limited in patients with IgA nephropathy (IgAN). The aim of this study was to develop and validate a prediction rule for estimating the individual risk of ESRD in patients with IgAN.

Design, setting, participants, & measurements A total of 698 patients with IgAN diagnosed by renal biopsy at Kyushu University Hospital (derivation cohort) between 1982 and 2010 were retrospectively followed. The Oxford classification was used to evaluate the pathologic lesions. The risk factors for developing ESRD were evaluated using a Cox proportional hazard model with a stepwise backward elimination method. The prediction rule was verified using data from 702 patients diagnosed at Japanese Red Cross Fukuoka Hospital (validation cohort) between 1979 and 2002.

Results In the derivation cohort, 73 patients developed ESRD during the median 4.7-year follow-up. The final prediction model included proteinuria (hazard ratio [HR], 1.30; 95% confidence interval [95% CI], 1.16 to 1.45, every 1 g/24 hours), estimated GFR (HR, 0.84; 95% CI, 0.74 to 0.96, every 10 ml/min per 1.73 m²), mesangial proliferation (HR, 1.85; 95% CI, 1.10 to 3.11), segmental sclerosis (HR, 3.21; 95% CI, 1.37 to 7.51), and interstitial fibrosis/tubular atrophy (T1: HR, 5.30; 95% CI, 2.63 to 10.7; T2: HR, 20.5; 95% CI, 9.05 to 46.5) as independent risk factors for developing ESRD. To create a prediction rule, the score for each variable was weighted by the regression coefficients calculated using the relevant Cox model. The incidence of ESRD increased linearly with increases in the total risk scores (*P* for trend <0.001). Furthermore, the prediction rule demonstrated good discrimination (c-statistic=0.89) and calibration (Hosmer-Lemeshow test, *P*=0.78) in the validation cohort.

Conclusions This study developed and validated a new prediction rule using clinical measures and the Oxford classification for developing ESRD in patients with IgAN.

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Introduction

IgA nephropathy (IgAN) is the most common cause of primary GN in the world (1). Many epidemiologic studies have identified risk factors for poor kidney prognosis in patients with IgAN, including age (2), sex (3), hypertension (3,4), decreased estimated GFR (eGFR) (2–5), proteinuria (2,3,6–8), and pathologic severity (2,9,10). In 2009, the Oxford classification of IgAN was proposed by an international working group (11,12). They identified four types of lesions as specific pathologic measures or features associated with the development of ESRD and/or halving of the eGFR: the mesangial hypercellularity score (M), endocapillary hypercellularity (E), segmental glomerulosclerosis (S), and tubular atrophy/interstitial fibrosis (T) (11). Recent findings have suggested the usefulness of this classification system for predicting kidney prognosis (13–16). However, the prediction ability of the Oxford classification itself for individual kidney prognosis may be limited because of the difficulty in estimating individuals' absolute risks for future ESRD.

The accurate prediction of kidney prognosis in individual cases is important for determining the therapeutic strategy. Therefore, some investigators have attempted to develop a risk prediction rule for estimating the risk of ESRD in individual patients (2,8,17). However, few studies have evaluated the internal or external validity of their risk prediction tools. Furthermore, these studies have not addressed the extent to which each pathologic measure and clinical variable contributed to the risk prediction of kidney prognosis. Herein, we developed a new clinical and pathologic risk prediction rule using the Oxford classification to identify the subgroup of Japanese patients with IgAN at high risk of developing ESRD, and we verified the external validity of the score in an independent cohort.

Materials and Methods

Study Population

Derivation Cohort. A total of 846 patients with primary IgAN, but not Henoch-Schönlein purpura,

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were diagnosed by kidney biopsy between January 1982 and December 2010 in six hospitals in the northern region of Kyushu Island in Japan (the Kyushu University Hospital, Hamanomachi Hospital, Munakata Medical Association Hospital, Japan Seamen's Relief Association Moji Hospital, Karatsu Red Cross Hospital, and Hakujuji Hospital). Among them, we excluded 85 patients whose biopsy specimens contained <10 glomeruli, 50 patients without available clinical measures, and 13 patients with IgA deposition secondary to such conditions as systemic lupus erythematosus or those with diabetes mellitus. Finally, 698 patients were enrolled in this study, as a derivation cohort. The patients were followed until December 31, 2012.

Validation Cohort. Seven hundred ninety-four patients with primary IgAN underwent biopsy between October 1979 and September 2002 and were followed for at least 1 year at Japanese Red Cross Fukuoka Hospital. Ninety-two biopsy specimens that contained <10 glomeruli were excluded. The remaining 702 patients were included in the study as a validation cohort (18).

Pathologic Measures

Pathologic lesions were evaluated according to the Oxford classification (11). The mesangial hypercellularity score (M) was defined as M0 if the score was ≤ 0.5 and M1 if the score was > 0.5 ; endocapillary hypercellularity (E) and segmental glomerulosclerosis (S) were expressed as E0 and S0, respectively, if absent and E1 and S1 if present. Tuft adhesions were regarded as S1 lesions. Tubular atrophy/interstitial fibrosis (T) was semiquantitatively classified according to the percentage of the cortical area involved in the tubular atrophy or interstitial fibrosis: T0 for 0%–25%, T1 for 26%–50%, and T2 for $> 50\%$.

Clinical Measures

Clinical measures were obtained from medical records at the time of the renal biopsy. They included age, sex, BP, cholesterol levels, triglycerides, serum creatinine, and 24-hour urinary protein excretion or urinary protein-to-creatinine ratio. Hypertension was defined as BP $\geq 140/90$ mmHg and/or current use of antihypertensive agents. Dyslipidemia was defined as serum total cholesterol level ≥ 220 mg/dl and/or serum triglyceride level ≥ 150 mg/dl. The categorical classifications for eGFR and urinary protein excretion were defined using traditional cutoffs for clinical significance (19). Total cholesterol concentrations and triglycerides were determined enzymatically. Serum creatinine was measured by the Jaffe method until April 1988 and by the enzymatic method from May 1988 onward at Kyushu University. At the other participating institutions, serum creatinine was measured by the Jaffe method until December 2000 and by the enzymatic method from January 2001. Serum creatinine values measured by the Jaffe method were converted to values for the enzymatic method by subtracting 0.207 mg/dl (20). The eGFR was calculated using the Schwartz formula in patients younger than age 18 years and the following formula in patients older than age 18 (21–23):

$$eGFR \text{ (ml/min per } 1.73m^2) = 194 \times Cr^{-1.094} \times Age^{-0.287} \\ \text{(if female, } \times 0.739).$$

The formula for adults proposed by Matsuo *et al.* (21) is accepted as a more accurate tool for the Japanese population than the previously reported one.

Renal Outcome

The primary endpoint was ESRD, which was defined as the initiation of renal replacement therapy (hemodialysis, peritoneal dialysis, or renal transplantation). The renal outcomes were surveyed by medical records or by telephone consultation with the clinics and hospitals the patients visited or with the patients themselves.

Statistical Analyses

In the derivation cohort, we performed univariate analyses to estimate the hazard ratios (HRs) with 95% confidence intervals (95% CIs) for each risk factor for the development of ESRD using a Cox proportional hazards model. For these analyses, patients were censored at the date of their death or at the end of follow-up for those still alive. To build the risk prediction model, we selected the independent risk factors for the development of ESRD using a multivariate Cox proportional hazards model with stepwise backward elimination with $P < 0.05$ for remaining variables; in this selection, the clinically or biologically plausible risk factors for ESRD were included as initial candidate variables: age, sex, hypertension, dyslipidemia, urinary protein excretion, eGFR, and pathologic variables based on the Oxford classification. The categories of eGFR 30–59 ml/min per $1.73 m^2$ and urinary protein excretion 0.5–1.0 g/24 hours were included in the final multivariable-adjusted model, which were defined using traditional cutoffs for clinical significance (19), although their β coefficients were not statistically significant. To generate a simple integer-based point score for each variable, we assigned scores by dividing regression coefficients by the value of the smallest coefficient in the model and rounding up to the nearest integer (24). The predicted 5-year absolute risk of incident ESRD according to the total risk score was calculated using relevant Cox models with the baseline survivor function.

For internal and external validation of the prediction rule, we examined the discrimination ability of the models, which was reported as the c-statistic (*i.e.*, area under the curve by logistic regression models with binary outcomes), as well as the calibration of the model by means of the Hosmer-Lemeshow chi-squared statistic in both the derivation and validation cohorts (25). For the Hosmer-Lemeshow chi-squared statistic, the observed 5-year absolute risk for the development of ESRD was calculated using the Kaplan-Meier method. The c-statistics of the different models in the derivation cohort and those of the risk prediction rule in the two independent cohorts were compared using the method of Henley and McNeil (26,27).

The incidence rates of ESRD according to the total risk score were calculated with the person-year method. The risk of incident ESRD per 5-point increment in the total risk score was estimated using the Cox model including the total risk score as a continuous variable. Statistical analyses were conducted using SAS software, version 9.2 (SAS Institute, Inc., Cary, NC), and MedCalc software, version 12.3.0.0 (bvba, Mariakerke, Belgium). A two-tailed P value < 0.05 was considered to represent statistically significant findings.

Ethical Considerations

This study was conducted with the approval of the Kyushu University Institutional Review Board for Clinical Research. The ethics committee of all participating institutions granted approval to waive requirement for written, informed consent because of the retrospective nature of the present study.

Results

As shown in Table 1, the mean age of patients in the derivation cohort ($n=698$) was 36.1 years, and 48.7% of patients were men. Patients in the validation cohort ($n=702$) had a mean age of 33.1 years; 42.2% of these patients were men. The derivation cohort was significantly older, and the proportion of men was significantly higher than in the validation cohort. The median follow-up times after renal biopsy were 4.7 years and 5.1 years in the derivation and validation cohort, respectively; the difference in follow-up period was not statistically significant. The proportion of heavy urinary protein excretion and the prevalences of Oxford classification grades E1, S1, or T2 were higher in the validation cohort than in the derivation cohort.

Development of the Risk Prediction Model for Kidney Prognosis in the Derivation Cohort

During the follow-up period, 73 patients (10.5%) in the derivation cohort experienced ESRD. Seven variables (hypertension, dyslipidemia, urinary protein excretion, eGFR, and the pathologic measures of the Oxford classification [M, S, and T]) were significantly associated with a higher risk of incident ESRD in univariate analysis (Table 2). Neither age nor sex was a significant risk factor for ESRD. In the multivariate analysis with stepwise backward elimination, two clinical variables—urinary protein excretion (HR, 1.30; 95% CI, 1.16 to 1.45, every 1 g/24 hour) and eGFR (HR, 0.84; 95% CI, 0.74 to 0.96, every 10 ml/min per 1.73 m^2)—and three pathologic measures—M (HR, 1.85; 95% CI, 1.10 to 3.11), S (HR, 3.21; 95% CI, 1.37 to 7.51), and T (T1: HR, 5.30; 95% CI, 2.63 to 10.7; T2: HR, 20.5; 95% CI, 9.05 to 46.5)—were selected as independent risk factors for incident ESRD. The goodness of fit of the developed risk prediction model was very good on the basis of the *c*-statistic (0.87; 95% CI, 0.82 to 0.92) (Figure 1) and the Hosmer-Lemeshow test (chi-squared statistic with 8 d.f.=4.65; $P=0.79$) (Supplemental Figure 1). In a

Table 1. Baseline characteristics of the patients in the derivation and validation cohorts

Characteristics	Derivation Cohort ($n=698$)	Validation Cohort ($n=702$)	<i>P</i> Value
Age (yr)	36.1±15.4	33.1±14.2	<0.001
Men (%)	48.7	42.2	0.01
Follow-up (yr)	4.7±1.7–9.3	5.1±3.0–8.6	0.22
Systolic BP (mmHg)	125.8±17.7	125.9±19.3	0.93
Diastolic BP (mmHg)	74.8±13.2	75.2±14.2	0.57
Hypertension ^a (%)	25.2	26.4	0.62
Serum total cholesterol (mg/dl)	202.9±49.5	198.3±45.1	0.07
Serum triglycerides (mg/dl)	127.2±80.4	123.2±94.7	0.40
Urinary protein excretion levels (%)			
<0.5 g/24 hr	38.0	36.6	<0.001
0.5–1.0 g/24 hr	23.4	16.8	
1.0–3.5 g/24 hr	31.1	34.6	
≥3.5 g/24 hr	7.5	12.0	
Estimated GFR (%)			
≥60 ml/min per 1.73 m^2	72.5	74.7	0.73
30–59 ml/min per 1.73 m^2	22.2	20.8	
15–29 ml/min per 1.73 m^2	4.3	3.4	
<15 ml/min per 1.73 m^2	1.0	1.1	
Pathologic measures (Oxford classification) (%)			
Mesangial hypercellularity score			
M0 (≤0.5 of glomeruli)	87.8	87.8	0.99
M1 (>0.5 of glomeruli)	12.2	12.2	
Endocapillary hypercellularity			
E0 (absence)	64.3	58.1	0.02
E1 (presence)	35.7	41.9	
Segmental glomerulosclerosis			
S0 (absence)	29.7	20.8	<0.001
S1 (presence)	70.3	79.2	
Tubular atrophy/interstitial fibrosis			
T0 (≤25%)	78.9	70.7	<0.001
T1 (26%–50%)	14.0	17.5	
T2 (>50%)	7.0	11.8	

Values given are means ± SD or percentages.

^aHypertension was defined as BP≥140/90 mmHg and/or current treatment with antihypertensive agents.

Table 2. Unadjusted and multivariate-adjusted hazard ratios for the development of ESRD in the derivation cohort

Variables	Patients (n)	Events (n)	Unadjusted		Multivariate-Adjusted ^a	
			HR (95% CI)	P Value	HR (95% CI)	P Value
Age						
<30 yr	307	25	1.00 (reference)		Not selected	
30–59 yr	326	43	1.23 (0.75 to 2.02)	0.41		
≥60 yr	65	5	1.20 (0.46 to 3.16)	0.71		
Sex						
Women	358	38	1.00 (reference)		Not selected	
Men	340	35	1.19 (0.75 to 1.89)	0.46		
Hypertension^b						
No	520	43	1.00 (reference)		Not selected	
Yes	175	29	1.89 (1.18 to 3.03)	0.01		
Dyslipidemia^c						
No	369	20	1.00 (reference)		Not selected	
Yes	314	52	3.44 (2.05 to 5.76)	<0.001		
Urinary protein excretion						
<0.5 g/24 hr	266	2	1.00 (reference)		1.00 (reference)	
0.5–1.0 g/24 hr	163	8	5.85 (1.24 to 27.6)	<0.001	3.32 (0.69 to 15.9)	0.13
1.0–3.5 g/24 hr	217	47	25.3 (6.1 to 104.0)	<0.001	6.26 (1.46 to 26.9)	0.01
≥3.5 g/24 hr	52	16	71.4 (16.4 to 313.9)	<0.001	16.2 (3.47 to 75.2)	<0.001
Estimated GFR						
≥60 ml/min per 1.73 m ²	506	25	1.00 (reference)		1.00 (reference)	
30–59 ml/min per 1.73 m ²	155	26	3.96 (2.30 to 6.94)	<0.001	1.37 (0.71 to 2.64)	0.35
15–29 ml/min per 1.73 m ²	30	16	30.6 (15.4 to 60.7)	<0.001	3.27 (1.45 to 7.40)	0.003
<15 ml/min per 1.73 m ²	7	6	115.9 (45.0 to 298.6)	<0.001	13.8 (4.54 to 42.0)	<0.001
Pathologic measures (Oxford classification)						
Mesangial hypercellularity score						
M0 (≤0.5 of glomeruli)	613	47	1.00 (reference)		1.00 (reference)	
M1 (>0.5 of glomeruli)	85	26	5.48 (3.38 to 8.90)	<0.001	1.78 (1.05 to 3.01)	0.03
Endocapillary hypercellularity						
E0 (absence)	449	49	1.00 (reference)		Not selected	
E1 (presence)	249	24	1.33 (0.81 to 2.19)	0.26		
Segmental glomerulosclerosis						
S0 (absence)	207	7	1.00 (reference)		1.00 (reference)	
S1 (presence)	491	66	5.25 (2.37 to 11.7)	<0.001	3.61 (1.54 to 8.47)	0.003
Tubular atrophy/interstitial fibrosis						
T0 (≤25%)	551	20	1.00 (reference)		1.00 (reference)	
T1 (26%–50%)	98	24	9.95 (5.39 to 18.4)	<0.001	5.27 (2.63 to 10.6)	<0.001
T2 (>50%)	49	29	57.2 (29.8 to 109.8)	<0.001	19.7 (8.47 to 45.7)	<0.001

HR, hazard ratio; CI, confidence interval.

^aVariables were selected by using a Cox proportional hazard model and a stepwise backward method with $P < 0.05$ for remaining variables to determine the risk factors for the development of ESRD. Neither age nor sex was a significant risk factor for the development of ESRD in univariate analysis.^bHypertension was defined as BP ≥ 140/90 mmHg and/or current treatment with antihypertensive agents.^cDyslipidemia was defined as serum total cholesterol ≥ 220 mg/dl and/or serum triglyceride ≥ 150 mg/dl.

sensitivity analysis, we compared the risk prediction ability of the models that included only the clinical measures or only the pathologic measures with that of the model that included both the clinical and pathologic measures by means of c-statistics (Figure 1). The results showed that the model including both the clinical and pathologic measures had a significantly higher c-statistic value than the other two models (both $P < 0.01$).

Because the recruitment period for this study was long, there was the possibility that differences in the treatment (e.g., use of renin-angiotensin inhibitors) over time modified the effect of each risk factor on the renal prognosis. However, there was no evidence of significant heterogeneity in the effect of any risk factor on renal prognosis between 1982–1996 and 1997–2010 (all P values for heterogeneity > 0.13).

Making a Score-Based Prediction Rule

A score-based prediction rule containing five variables selected in multivariate analysis was made using the regression coefficients obtained from the relevant Cox model (Table 2). A regression coefficient of 0.31, which was the smallest value among the variables included, corresponded to 1 point (Table 3). The incidence rate of ESRD increased linearly with increases in the total risk score (P for trend < 0.001) (Figure 2A). Every 1-point increment in the total risk score was associated with a 1.33-fold (95% CI, 1.18- to 1.50-fold) increased risk of ESRD. The predicted 5-year absolute risks of ESRD per 1-point increment in the total prediction rule are shown in Table 4. The prediction rule showed the model's excellent discrimination ability for incident ESRD with a c-statistic of 0.87 (95% CI, 0.82 to 0.92) (Supplemental Table 1) and a good calibration in the Hosmer-Lemeshow test (chi-squared statistic with 8 d.f.=2.80; $P=0.95$) (Figure 3A).

External Validation of the Prediction Rule in the Validation Cohort

The prediction rule was externally validated in a population (the validation cohort) independent from the derivation cohort. During the follow-up period, 85 patients (12.1%) experienced ESRD in the validation cohorts. In the validation cohort, the incidence of ESRD also increased linearly with an increase in the total risk score (P for trend < 0.001) (Figure 2B). The prediction rule had very good discrimination (c-statistic=0.89; 95% CI, 0.86 to 0.93). There was no evidence of a significant difference in the c-statistics between the derivation and the validation cohorts ($P=0.39$) (Supplemental Table 1). Furthermore, the prediction rule also showed good calibration in the validation cohort (chi-squared statistic with 8 d.f.=4.75; $P=0.78$) (Figure 3B).

Subgroup Analysis by Sex or Age

We also conducted analyses stratified by age and sex in both cohorts (Supplemental Table 1). There was no evidence of a significant difference in c-statistics between the derivation and the validation cohorts, regardless of age and sex (all $P > 0.25$). Further, the prediction rule had good discrimination for ESRD (c-statistic=0.86; 95% CI, 0.81 to 0.92) and was well calibrated ($P=0.61$) when analysis was performed ($n=650$) excluding those younger than 18 years old.

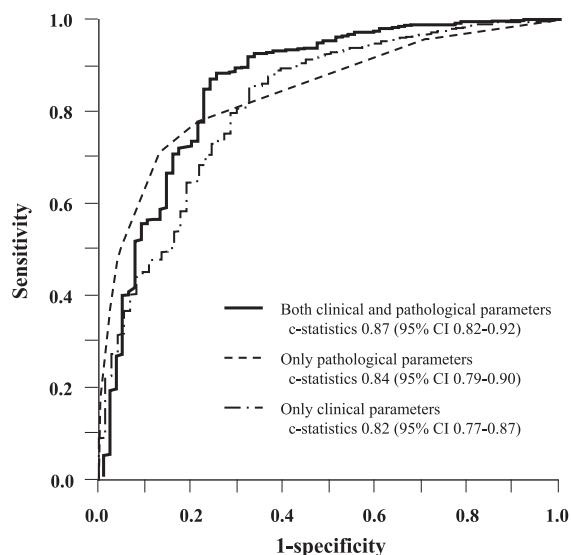


Figure 1. | Comparisons of c-statistics among the risk prediction models using only clinical measures, only pathologic measures, or both clinical and pathologic measures in the derivation cohort. The risk prediction model with only clinical measures included urinary protein excretion and estimated GFR. The model with only pathologic measures included mesangial score (M), segmental glomerulosclerosis or adhesion (S), and the severity of tubulointerstitial disease (T). The model with the clinical and pathologic measures included all of the above-mentioned clinical and pathologic measures. CI, confidence interval.

Table 3. Risk scores for the development of ESRD

Variables	Scores
Urinary protein excretion	
<0.5 g/24 hr	0 points
0.5–1.0 g/24 hr	4 points
1.0–3.5 g/24 hr	6 points
≥3.5 g/24 hr	9 points
Estimated GFR	
≥60 ml/min per 1.73 m ²	0 points
30–59 ml/min per 1.73 m ²	1 point
15–29 ml/min per 1.73 m ²	4 points
<15 ml/min per 1.73 m ²	9 points
Pathologic measures (Oxford classification)	
Mesangial hypercellularity score	
M0 (≤0.5 of glomeruli)	0 points
M1 (>0.5 of glomeruli)	2 points
Segmental glomerulosclerosis	
S0 (absence)	0 points
S1 (presence)	4 points
Tubular atrophy/interstitial fibrosis	
T0 (≤25%)	0 points
T1 (26%–50%)	6 points
T2 (>50%)	10 points
Maximum total risk scores	34 points

Discussion

In this study, we developed and validated a new prediction rule consisting of five variables: proteinuria; eGFR; and the Oxford histologic grades M, S, and T. Of note, our

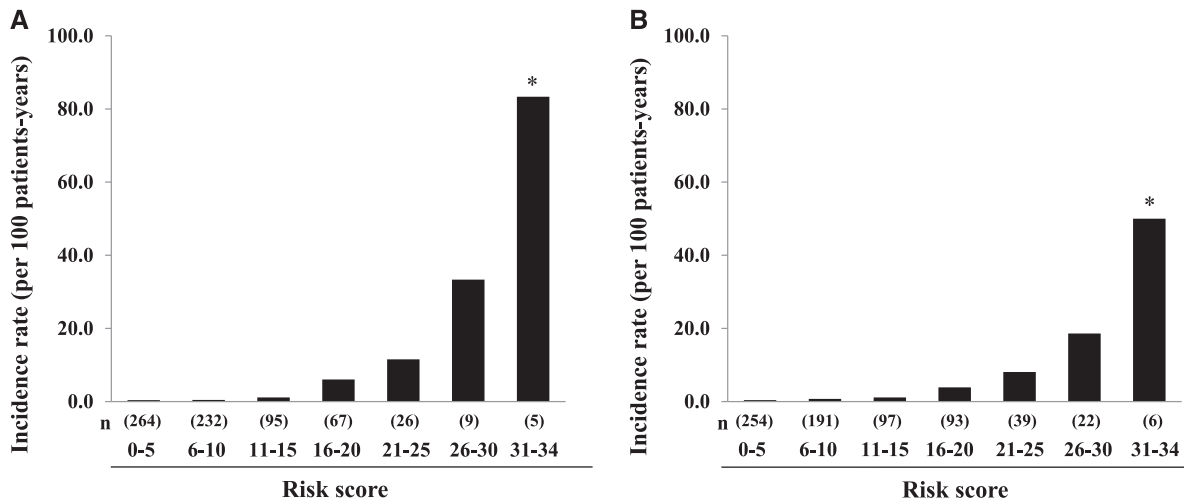


Figure 2. | The incidence rate of ESRD by 5-point increments of total risk score in the derivation cohort (A) and the validation cohort (B). *P for trend <0.001.

Total Risk Score (Points)	Predicted 5-Year Absolute Risk (%)
0	0.07
1	0.10
2	0.13
3	0.18
4	0.24
5	0.33
6	0.45
7	0.60
8	0.81
9	1.09
10	1.47
11	1.98
12	2.66
13	3.58
14	4.80
15	6.42
16	8.57
17	11.40
18	15.10
19	19.80
20	25.70
21	33.10
22	41.90
23	51.90
24	62.80
25	73.70
26	83.50
27	91.20
28	96.20
29	98.80
30	99.70
≥31	100.00

ing that both clinical and pathologic measures should be taken into account in the risk assessment for kidney prognosis. Furthermore, our prediction rule showed good discriminations of future ESRD and the incidence of ESRD estimated by the prediction model exhibited a good fit with the observed incidence rate of ESRD in both the derivation and validation cohorts. These results suggest that our prediction rule is statistically valid and accurate for the risk assessment of ESRD among Japanese patients with IgAN. We believe that this score will be useful for determining the initial therapeutic strategies of patients with IgAN.

Several risk scores for developing ESRD in patients with IgAN have been reported previously (2,8,17). However, these studies had several limitations. The study of Goto *et al.*, in which 2283 Japanese patients were followed up for a median of 7.3 years, had the limitations inherent in a mail-based survey: namely, the authors could not collect complete data for about a quarter of the participants or check the quality of the datasets (2,28). Berthoux *et al.* made a risk score from the data of 332 French patients for a median 11.3-year follow-up (8), but their score evaluated histologic lesions using their own original pathologic classification system (29,30), which is not generally accepted. Xie *et al.* developed a risk score for predicting ESRD in 619 Chinese patients followed for an average of 3.4 years and showed that their score had a better predictive performance on their dataset than previously published risk scores (17). However, their score has not been verified in an independent validation cohort. Furthermore, none of these previously published risk scores used an internationally accepted pathologic grading system, such as the Oxford classification (12,14,31). To our knowledge, our study is the first investigation attempting to develop a risk score using the Oxford classification and to verify its external validity in an independent cohort.

sensitivity analysis revealed that the model with both clinical and pathologic measures had a statistically better fit than the model with either type of measure alone, suggest-

In the present study, we identified five variables as risk factors for ESRD by multivariate analysis—proteinuria, eGFR, and the Oxford pathologic measures—which is consistent with many prior investigations (1–7,9,10,32–34). In

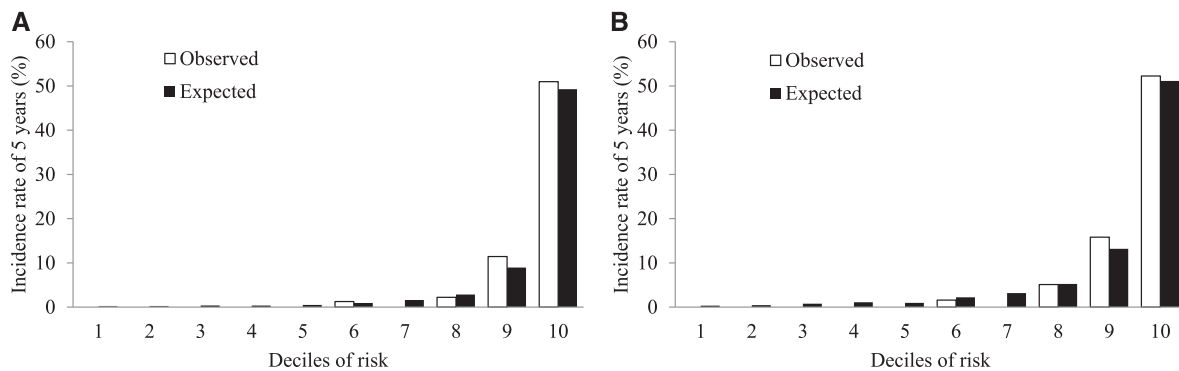


Figure 3. | Observed and predicted 5-year absolute risk for the development of ESRD by deciles of risk in the derivation cohort (A) and the validation cohort (B). Hosmer-Lemeshow chi-squared statistic=2.80, d.f.=8, $P=0.95$ for the derivation cohort and Hosmer-Lemeshow chi-squared statistic=4.75, d.f.=8, $P=0.78$ for the validation cohort.

particular, increased proteinuria and decreased eGFR have been well acknowledged as major risk factors for ESRD among patients with IgAN (3,5,6,9,35). In this study, M, S, and T were independently associated with renal outcome. T has been established as a significant prognostic factor associated with renal outcome in other studies, whereas the effects of M and S on renal prognosis have been inconsistent among studies (11,15,16,36). Some reports showed that M or S were not both selected as risk factors for future ESRD in multivariate analysis (15,16,36). These inconsistencies may be due to insufficient statistical power resulting from the small sample sizes of the studies. Like most previous studies, the present study failed to reveal a significant association of E with renal prognosis (15,18,36). Therefore, the variables included in our prediction rule would seem to be reasonable for application to research or clinical studies.

Several limitations of this study should be noted. First, one-time measurement of each risk factor may have caused the misclassification of study patients. Such misclassification would weaken the association found in this study, biasing the results toward the null hypothesis. Specifically, a previous study found that average BP is a more important predictor of renal prognosis than a single measurement at the time of biopsy or first visit (6,37). Therefore, the lack of information on average BP during the follow-up period would tend toward the underestimation of the effect of BP on the renal prognosis, which may be why hypertension was not statistically selected as a risk factor for ESRD in this study. Second, we could not consider the effect of therapeutic intervention on the kidney prognosis. Patients with heavy proteinuria and advanced pathologic findings were likely to receive aggressive treatment (e.g., steroid and cytotoxic agents), which may have shifted the effect of these risk factors on the risk of ESRD downward. Finally, the generalizability of our prediction rule for populations of other races may be limited. It has been reported that the renal survival rates in Japanese patients with IgAN resemble those in Europe and China (35,38,39). However, the patients in the present study showed only 10% of ESRD, which reflected the relatively “low-risk” features of our cohort compared with other published cohorts. In fact, most of our patients had

relatively low-grade proteinuria (<1 g/d) and low-risk pathologic features, and fewer had hypertension and advanced CKD stage than those in previous studies (6,8,17,37). Thus, one should take care in applying the risk scores developed in the present study to a traditionally high-risk population of patients with IgAN. It may be necessary to evaluate the validity of our prediction rule in other populations. However, we believe that this prediction rule may also provide useful risk assessments for ESRD in patients with IgAN in populations of other races or ethnic groups.

In conclusion, we have developed a new prediction rule for developing ESRD in patients with IgAN and verified its validity in an independent cohort. This prediction rule should provide a useful guide to estimate the individual risk for ESRD in patients with IgAN and may be effective at identifying those at high risk for the future development of ESRD. Further investigations will be needed to fully confirm that therapeutic interventions based on this prediction rule can reduce the burden of patients with IgAN.

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Disclosures

None.

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Supplementary table 1. Comparisons of the *c*-statistics of the prediction rule between the subgroups stratified by age or sex

Subgroups	Derivation cohort	Validation cohort	<i>P</i> value ^a
	(<i>n</i> = 698)	(<i>n</i> = 702)	
	<i>c</i> -statistics (95% CI)	<i>c</i> -statistics (95% CI)	
Overall population	0.87 (0.82 to 0.92)	0.89 (0.86 to 0.93)	0.39
Subgroup stratified by age			
Younger (<30 years)	0.86 (0.77 to 0.95)	0.90 (0.85 to 0.96)	0.41
Older (≥30 years)	0.87 (0.81 to 0.93)	0.90 (0.85 to 0.93)	0.56
Subgroup stratified by sex			
Men	0.88 (0.82 to 0.95)	0.88 (0.83 to 0.93)	0.90
Women	0.87 (0.81 to 0.94)	0.92 (0.88 to 0.96)	0.25

Abbreviation: CI, confidence interval.

^a*P* value for the differences in *c*-statistics between the derivation and the validation cohorts.

Supplemental Figure 1

Observed and predicted development of end-stage kidney disease by deciles of risk in the derivation cohorts (A) and the derivation cohorts.

Hosmer-Lemeshow χ^2 -statistic = 4.65, d.f. = 8, $P = 0.79$.

Supplementary figure 1

