

A retrospective cohort study of clinical characteristics and outcomes of type 2 diabetic patients with kidney disease

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Background: Type 2 diabetes mellitus (T2DM) with chronic kidney disease (CKD) poses a serious health threat and becomes a new challenge. T2DM patients with CKD fall into three categories, diabetic nephropathy (DN), non-diabetic kidney disease (NDKD), and diabetic nephropathy plus non-diabetic kidney disease (DN+NDKD), according to kidney biopsy. The purpose of our study was to compare the clinical characteristics and kidney outcomes of DN, NDKD, and DN+NDKD patients.

Methods: Data on clinical characteristics, pathological findings, and prognosis were collected from June 2016 to July 2022 in patients with previously diagnosed T2DM and confirmed DN and or NDKD by kidney biopsy at Tongji Hospital in Wuhan, China. The endpoint was defined as kidney transplantation, dialysis, or a twofold increase in serum creatinine.

Results: In our 6-year retrospective cohort research, a total of 268 diabetic patients were admitted and categorized into 3 groups by kidney biopsy. The 268 patients were assigned to DN (n=74), NDKD (n=109), and DN+NDKD (n=85) groups. The most frequent NDKD was membranous nephropathy (MN) (n=45, 41.28%). Hypertensive nephropathy was the most common subtype in the DN+NDKD group (n=34, 40%). A total of 34 patients (12.7%) reached the endpoint. The difference between the Kaplan-Meier survival curves of the DN, NDKD, and DN+NDKD groups was significant ($p < 0.05$). Multifactorial analysis showed that increased SBP [HR(95% CI): 1.018(1.002-1.035), $p=0.025$], lower Hb [HR(95% CI): 0.979(0.961-0.997), $p=0.023$], higher glycosylated hemoglobin [HR(95% CI): 1.338(1.080-1.658), $p=0.008$] and reduced serum ALB [HR(95% CI): 0.952(0.910-0.996), $p=0.032$] were risk factors for outcomes in the T2DM patients with CKD.

Conclusions: This research based on a Chinese cohort demonstrated that the risk of endpoint events differed among DN, NDKD, and DN+NDKD patients. In T2DM patients with CKD, DN patients were more often responsible for the kidney endpoints. Increased SBP, higher glycosylated hemoglobin, lower Hb, and decreased serum ALB may be correlated with adverse kidney outcomes in T2DM patients.

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14

15 Abstract

16 **Background:** Type 2 diabetes mellitus(T2DM) with chronic kidney disease (CKD) poses a
17 serious health threat and becomes a new challenge. T2DM patients with CKD fall into three
18 categories, diabetic nephropathy (DN), non-diabetic kidney disease (NDKD), and diabetic
19 nephropathy plus non-diabetic kidney disease (DN+NDKD), according to kidney biopsy. The
20 purpose of our study was to compare the clinical characteristics and kidney outcomes of DN,
21 NDKD, and DN+NDKD patients.

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27 admitted and categorized into 3 groups by kidney biopsy. The 268 patients were assigned to DN
28 (n=74), NDKD (n=109), and DN+NDKD (n=85) groups. The most frequent NDKD was
29 membranous nephropathy (MN) (n=45,41.28%). Hypertensive nephropathy was the most
30 common subtype in the DN+NDKD group (n=34,40%). A total of 34 patients (12.7%) reached
31 the endpoint. The difference between the Kaplan-Meier survival curves of the DN, NDKD, and
32 DN+NDKD groups was significant ($p < 0.05$). Multifactorial analysis showed that increased
33 SBP[HR(95% CI):1.018(1.002-1.035), $p=0.025$], lower Hb[HR(95% CI): 0.979(0.961-0.997),
34 $p=0.023$], higher glycosylated hemoglobin [HR(95% CI): 1.338(1.080-1.658), $p=0.008$] and
35 reduced serum ALB[HR(95% CI):0.952(0.910-0.996), $p=0.032$] were risk factors for outcomes in
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37 **Conclusions:** This research based on a Chinese cohort demonstrated that the risk of endpoint
38 events differed among DN, NDKD, and DN+NDKD patients. In T2DM patients with CKD, DN
39 patients were more often responsible for the kidney endpoints. Increased SBP, higher
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41 kidney outcomes in T2DM patients.

42 **Keywords:** Diabetic nephropathy; Non-diabetic kidney disease; Type 2 diabetes mellitus;
43 kidney biopsy.

44

45 Introduction

46 More than 500 million people around the world, accounting for over 10.5% of the global adult
47 population are affected by diabetes mellitus (Sun et al. 2022). Type 2 diabetes mellitus(T2DM)
48 comprises the majority of cases. There has been an increase in the population with T2DM from
49 1990 to 2019 universally in a systematic analysis of T2DM (Ye et al. 2023). The incidence of
50 T2DM among Chinese adults was 12.4%, higher than the world estimate reported by Wang, L.
51 et al (Wang et al. 2021).

52 Chronic exposure to hyperglycaemia affects the microvasculature in multiple organs, including
53 the kidney, the ocular, the peripheral nervous systems and so on (Barrett et al. 2017). Based on
54 the pathological diagnosis, T2DM patients with chronic kidney disease (CKD) can be classified
55 into diabetic nephropathy (DN), non-diabetic kidney disease (NDKD), and diabetic nephropathy
56 plus non-diabetic kidney disease (DN+NDKD) (Anders et al. 2018). DN affects approximately
57 one-quarter of the diabetic population, which is the primary etiology of end stage renal disease
58 (ESRD) (Faselis et al. 2020). In China, the prevalence of DN was nearly one-fifth of patients
59 with T2DM (21.8%) (Zhang et al. 2020). The prevalence of DN has remained stable while the
60 prevalence of NDKD in T2DM fluctuated greatly. The prevalence of NDKD ranged from 6.5% to
61 94%, with an average of 41.3% (Zhang et al. 2022). Part of the reason for the difference in
62 prevalence is the discrepancy in clinical practice. This is a reflection of the wide range of
63 considerations by clinicians before a patient undergoes a renal biopsy. NDKD can be either a
64 solitary disease or a coexistence with DN. The diagnosis of NDKD is important since the
65 complete reversal of NDKD is achievable through accurate diagnosis and prompt treatment.
66 The pathological feature and clinical characteristics of T2DM with CKD are likely to change
67 under the conditions of aging population, increasing incidence of infections and malignancies,
68 and the environmental pollution (Prasad et al. 2023). Our understanding of the pathophysiologic
69 mechanisms of T2DM with CKD has progressed as we continue to refine our classification of
70 the pathologic types of T2DM with CKD. The commonly reported variables of NDKD, including
71 DM shorter duration, lower glycosylated hemoglobin, absence of retinopathy, lower blood
72 pressure, hematuria, higher proteinuria, higher hemoglobin, and lower serum creatinine were
73 considered as the risk factors for kidney function progression in previous studies (Horvatic et al.
74 2014; Jing et al. 2021; Prasad et al. 2023). To our knowledge, the etiology and demographic
75 data is limited in South China, and there have been few studies comparing the prognosis of
76 T2DM patients with CKD based on the classification of DN, NDKD, and DN+NDKD.

77 Therefore, it is imperative to reassess CKD in T2DM and know the spectrum of T2DM with CKD

78 considering the huge burden of T2DM and diabetes-related kidney diseases in China. Our study
79 used the cohort in our center to further evaluate the differences in prognosis among DN, NDKD
80 and DN+NDKD patients. The endpoint was defined as kidney transplantation, dialysis or a
81 twofold increase in serum creatinine. Thus, it will be possible to clarify whether patients with DN
82 have a worse prognosis than patients with NDKD and to investigate prognostic risk factors. By
83 managing the associated risk factors, our research is expected to provide preventive or
84 therapeutic interventions for T2DM patients with CKD. Effective prevention can reduce the
85 disease burden in patients with CKD and improve their quality of life and prognosis.

86

87 **Materials & Methods**

88 Study design and patients

89 Patients with previously diagnosed T2DM with CKD by kidney biopsy were enrolled from 1 June
90 2016 to 31 July 2022, at Tongji Hospital, Wuhan, China in this retrospective study. The inclusion
91 criteria for this study were as followed: (i) age >18 years; (ii) clinical diagnosis of T2DM; (iii)
92 underwent kidney biopsy. The exclusion criteria included the items below: (i)ESRD diagnosed
93 before kidney biopsy or estimated glomerular filtration rate (eGFR) <15 ml/min per 1.73 m²
94 (exclusion:7); (ii)patients with other types of diabetes mellitus or combined malignancy
95 (exclusion:18); (iii)severe clinical data deficit (exclusion:22); (iii)kidney transplantation, acute
96 kidney injury and urinary tract infection (exclusion:0). The endpoint was defined as kidney
97 transplantation, dialysis or a twofold increase in serum creatinine. This study adhered to the
98 tenets of the Declaration of Helsinki declaration. Informed consent was waived by the Ethics
99 Review Board of Tongji Hospital, Tongji Medical College, Huazhong University of Science and
100 Technology(No.TJ-IRB20210929).

101

102 Data acquisition

103 From the electronic medical record, we extracted demographic data (age and sex), blood
104 pressure values (systolic blood pressure [SBP] and diastolic blood pressure [DBP]), and
105 medication history, with all original examination dates obtained from patients' initial admissions.
106 All original examination dates were derived from patients initial admission. Clinical data included
107 hemoglobin (Hb), serum albumin(ALB), 24h proteinuria, eGFR, serum creatinine (Scr),
108 hemoglobin A1C(HbA1C), immunoglobulin G (IgG), immunoglobulin A (IgA), immunoglobulin M
109 (IgM), complement 3 (C3), and complement 4 (C4).

110

111 Pathological examination

112 The kidney puncture tissues were examined by light microscopy, immunofluorescence and
113 electron microscopy. The pathological diagnosis of DN was based on the 2010 version of the
114 pathologic classification of diabetic nephropathy (Tervaert et al. 2010). A diagnosis of DN is
115 confirmed by one of the following conditions: class I, glomerular basement membrane
116 thickening; class II, mild (IIa) or severe (IIb) mesangial expansion; Class III, nodular sclerosis
117 (Kimmelstiel-Wilson lesions): at least one glomerulus with nodular increase in mesangial matrix
118 (Kimmelstiel-Wilson); Class IV, more than 50% global glomerulosclerosis. Light microscopy,

119 immunofluorescence, and electron microscopy were used to diagnose NDKD based on
120 characteristic changes. Two experienced independent pathologists reviewed all biopsy
121 specimens.

122

123 Statistical analysis

124 SPSS (version 25.0, IBM, US) software and R software (version 4.2.2, <https://www.r-project.org>)
125 were used to analyze the full analysis set. Data were presented as median (interquartile range,
126 IQR) or mean (standard deviation, SD) after normality tests on continuous variables, and as
127 numbers and percentages on categorical variables. Missing values were imputed by predictive
128 mean of closest points. The one-way ANOVA test, the Kruskal-Wallis test, and the χ^2 test were
129 used to assessed differences. Endpoints were defined as dialysis, death, or twofold increase in
130 serum creatinine. Kaplan-Meier analysis and Cox regression analysis were utilized to perform
131 time-to-event analysis. Kaplan-Meier survival curves were plotted for patients in the DN, NDKD,
132 and DN+NDKD groups. The results were compared using log-rank tests. Relevant risk factors
133 and covariates with p values <0.1 were included in Cox regression proportional risk models.
134 Covariates included indicators that met the requirements or were clinically significant after
135 univariate analysis. Variables were entered into the Cox model through backward entry method.
136 The validity was determined by testing the chi-square value of the Cox model.

137

138 Results

139 Baseline characteristics of three groups

140 From a cohort of 315 T2DM patients underwent kidney biopsy between 2016 and 2022, we
141 excluded 47 non-compliant patients after applying exclusion criteria. The remaining 268 patients
142 included 74 in the DN group, 109 in the NDKD group, and 85 in the DN+NDKD group, or
143 27.61%, 40.67%, and 31.72% of the cohort, respectively (Fig.1).

144 Table 1 presents the baseline characteristics of our study. 180(67.2%) of the 268 patients with
145 inclusion criteria were male and 88(32.84%) were female. The median (\pm IQR) age of all those
146 included in the criteria was 52.50 \pm 15 years, varying from 26 to 73 years. The median age of the
147 DN group was 50 years and it was 54 years (IQR 15) in the NDKD group, and 53 years (IQR
148 12) in the DN+NDKD group. We followed patients for an average of 16.44 months. The mean
149 duration of DM was 60 months, in order of NDKD, DN+NDKD, DN, from shortest to longest.
150 Immunosuppressants and glucocorticoids were most commonly used in the NDKD group.
151 Insulin was the predominant treatment in the DN group. Statistically meaningful differences
152 were observed in the three groups with regard to gender ($p=0.001$), family history of diabetes
153 ($p=0.021$), duration of T2DM ($p=0.007$), diabetic retinopathy ($p<0.001$), red blood cell count
154 ($p=0.043$), urinary sediment red blood cell count ($p=0.004$), glycosylated hemoglobin ($p=0.002$),
155 HDL ($p=0.043$), and C3 ($p=0.003$) (table 1).

156

157 Pathological characteristics of kidney alterations in T2DM patients

158 Typical DN pathologic images are shown (Fig 2). The most prevalent pathological type in the

159 NDKD group was membranous nephropathy (n=45). Other subtypes within the NDKD category
160 were IgA nephropathy(n=26), hypertensive nephropathy (n=10), Henoch-Schoenlein purpura
161 nephritis(n=5), obesity-related glomerulopathy(n=4), focal segmental glomerulosclerosis(n=3),
162 light chain deposition disease (LCDD) (n=1), kidney amyloidosis(n=1), tubulointerstitial
163 nephritis(n=2), thrombotic microangiopathy(n=2), hepatitis B virus-related nephropathy(n=1),
164 sclerosing glomerulonephritis(n=3), minimal change disease(n=2), proliferative glomerular
165 lesions(n=4). Hypertensive nephropathy (n=34) was the dominant subtype, followed by IgA
166 nephropathy (n=15) in the DN+NDKD group (table 2, supplementary fig 1).

167

168 The comparison of the cumulative incidence of endpoints in T2DM with CKD patients
169 Our average follow-up in this cohort was 16.44 months. The study's endpoints were all-cause
170 death, kidney transplantation, dialysis, and a twofold increase in serum creatinine. For an
171 overall endpoint frequency of 12.7%, a total of 34 patients met the endpoint. After analyzing the
172 incidence of the endpoints, our study found that the number of patients with endpoints in the
173 three groups were 13, 9, 12, with proportions of 17.57%, 8.26%, 14.12%. Endpoint incidence
174 was notably greater in the DN group compared to the other groups ($p<0.05$) (Fig 3). The median
175 survival time remained at 52.0 months for NDKD and 34.5 months for DN. The median survival
176 time of DN+NDKD group can't be estimated as there were few endpoints in this group and most
177 survival times correspond to survival probabilities greater than 0.5. One-year survival rate of
178 kidney in each group were 88.8%, 97.4%, 87.7% in the DN, NDKD and DN+NDKD group.

179

180 Prognostic factors for endpoints

181 The proportional hazards (PH) assumption tests were conducted for the variables in the
182 endpoints. The test results indicated that all variables satisfied the PH assumption. A
183 multivariate Cox proportional hazards regression model included baseline variables that were
184 deemed clinically relevant or univariately associated with the outcomes. The final model was
185 simplified by careful selection of variables based on the number of events available (table 3).
186 Lower serum ALB[HR(95%CI):0.685(0.559-0.839), $p<0.001$], 24h proteinuria[HR(95%CI):
187 0.999(0.999-1.000), $p=0.006$], and increased SBP[HR(95%CI):1.047(1.006-1.089), $p=0.024$]
188 and age[HR(95%CI): 0.890(0.802-0.988), $p=0.028$] were determined to be important
189 contributors to adverse kidney outcomes in the DN group by multivariate Cox regression
190 analysis (table 4). NDKD patients with higher 24h proteinuria[HR(95%CI): 1.000(1.000-1.001),
191 $p=0.019$] and decreased C3 [HR(95%CI):0.001(0.000-0.356), $p=0.021$], were at increased risk
192 for adverse kidney effects (Table 4). Multivariate Cox regression results showed that serum
193 ALB[HR(95%CI):0.828(0.724-0.947), $p=0.006$], Scr[HR(95%CI): 1.011(1.005-1.018), $p=0.001$],
194 IgM[HR(95%CI): 13.708(3.611-52.034), $p<0.001$], SBP[HR(95%CI): 1.050(1.007-1.094),
195 $p=0.021$], age[HR(95%CI): 0.851(0.756-0.958), $p=0.008$] were significant risk indicators for the
196 endpoint event in the cohort of the DN+NDKD group (Table 4). T2DM patients with CKD
197 showed that SBP [HR(95%CI): 1.018(1.002-1.035), $p=0.025$], Hb[HR(95%CI): 0.979(0.961-
198 0.997), $p=0.023$], ALB[HR(95%CI): 0.952(0.910-0.996), $P=0.032$], glycosylated hemoglobin
199 [HR(95%CI): 1.338(1.080-1.658), $p=0.008$], were independent indicators of risk for the adverse

200 kidney outcomes.

201

202 **Discussion**

203 T2DM with CKD patients were divided into three groups in this study according to kidney biopsy.
204 Our results found that 40.67% of biopsied T2DM patients were diagnosed with NDKD and the
205 incidence of DN+NDKD was more than one-third (31.72%) of T2DM patients. Previous study
206 showed that the prevalence of NDKD averaged 41.3%(Zhang et al. 2022) and prevalence in the
207 DN+NDKD group varied from 4.7% to 19.72% (Fontana et al. 2021; Liu et al. 2016; Shadab et
208 al. 2022). The above study demonstrates that a high proportion of T2DM patients with CKD still
209 have NDKD, and that there is a great heterogeneity in the prevalence.

210 MN was the most prevalent with 41.28 %, followed by IgA nephropathy with 23.85 % in our
211 study, consistent with the findings reported by Wang (Wang et al. 2019). But some researchers
212 conclude that the major pathologic subtype of NDKD is IgA nephropathy (Byun et al. 2013; Zhou
213 et al. 2008). Regional and ethnic differences, as well as the mechanism of kidney pathologic
214 diagnosis, may contribute to the pathologic distribution of the NDKD group.

215 Progression of T2DM, poor glycemic control, DR, deterioration of kidney function, hematuria,
216 hypertension can guide to differentiation between DN and NDKD in many studies (Li et al. 2020;
217 Popa et al. 2021; Saini et al. 2021), which were consistent with our findings. Pallayova et al.
218 found that a strong predictor of NDKD was low serum HbA1c level (Pallayova et al. 2015). The
219 ratio of glycated albumin to HbA1c, according to Wang, was better biopsy-proven DN indicators
220 than HbA1c (Wang et al. 2017). DN and DR, as the two most important microvascular diseases
221 of T2DM, share many pathophysiologic and pathologic similarities. DR was closely correlated
222 with DN (\pm NDKD), and the absence of DR was a highly predictive of NDKD (Lin et al. 2018),
223 while Kritmetapak et al. found that in a multivariate analysis, DR was not an independent
224 predictor (Kritmetapak et al. 2018) and the association between DN and DR is not exactly
225 parallel conducted by Li, M. et al (Li et al. 2021). Usually, lack of DR is predictive of NDKD, but
226 does not exclude DN.

227 The hemoglobin levels in the DN patients were markedly lower as opposed to the NDKD
228 patients. In the primal stages of kidney disease, studies have revealed that CKD patients with
229 T2DM may become anemic (Xie et al. 2023). A recent cohort study in Japan showed that serum
230 Hb concentration, reflecting the onset of kidney fibrosis, may be useful in predicting the
231 development of DN (Yamanouchi et al. 2022). Ito, K. considered that because of severe
232 interstitial fibrosis and tubular atrophy, DN is associated with anemia and anemia may aid in
233 clinical differentiation between isolated DN and NDKD (Ito et al. 2021). Furthermore,
234 erythrocytes deformability and lifespan are also reduced by chronic inflammation and advanced
235 glycation end products (Tsai & Tarng 2019).

236 In our study, HDL levels differed at baseline levels, but did not affect the prognosis.

237 Nevertheless low HDL-C and high TG levels, in an Italian study, were considered independent
238 risk factors for DN prognosis over 4-year period (Russo et al. 2016). The cause of high TG and
239 low HDL-C may be caused by metabolic syndrome, and may result from underlying insulin

240 resistance. Multiple aspects of kidney function, including kidney hemodynamics and tubular
241 function, are adversely affected by insulin resistance (Artunc et al. 2016).

242 The pathological classification of CKD with T2DM, in our results, was significantly associated
243 with kidney prognosis. Sun et al have also shown that DN patients had relatively poorer
244 outcomes than NDKD (Sun et al. 2023). DN patients have a faster progression to ESRD than
245 other CKD etiologies, requiring earlier kidney replacement therapy, which results in a significant
246 health and economic burden.

247 To further investigate potential predictors of the kidney endpoint of T2DM patients, we
248 conducted the multivariate Cox regression analyses in T2DM patients with CKD. We found that
249 lower serum ALB, elevated SBP, glycosylated hemoglobin and Hb were independent risk
250 factors for the endpoint of all patients. In addition, we explored factors affecting kidney
251 prognosis through subgroup analyses, including those mentioned above. ALB, 24h proteinuria,
252 SBP and age were the most powerful risk factors for adverse kidney outcomes of the DN group
253 in our analysis. These factors largely correspond to the risk factors traditionally linked to DN.

254 Hypoalbuminemia may reflect multiple diseases: cirrhosis, malnutrition, kidney diseases and
255 chronic inflammation (Aldebeyan et al. 2017; Efremova et al. 2023; Sheinenzon et al. 2021).

256 Therefore, hypoalbuminemia may influence the progression of CKD through the mechanisms
257 described above. In Japanese patients with CKD, there was a negative and non-linear
258 relationship between ALB and the decline in kidney prognosis (Cheng et al. 2023). Moreover,
259 hypertension, identified as an independent predictor of microvascular complications (Asghar et
260 al. 2023), induces oxidative stress and inflammation in the kidney (Lopes de Faria et al. 2011).

261 With the exception of age, most of these risk factors are controllable, which is particularly critical
262 for the management of DN. It is known that typical lesions of diabetic nephropathy include
263 glomerular hyperfiltration and podocyte injury. Among the mechanisms of podocyte injury are
264 lipotoxicity, oxidative stress, mitochondrial damage, and autophagy (Li et al. 2023b; Nagata
265 2016). In fact, the molecular mechanism of DN is complex and many pathways are involved in
266 DN development and progression in a hyperglycemic environment including polyol,
267 hexosamine, PKC, and AGE pathways. These indicators may be involved in the progression of
268 diabetic nephropathy through the mechanisms described above.

269 Surprisingly, C3 was identified as putative risk features for the endpoint in the NDKD group. It is
270 universally acknowledged that C3 is an important part of the complement system and has three
271 distinct modes of activation: classic, lectin, and alternative. Jiayi Li's study found that MN
272 patients with 24h proteinuria over 0.75g or serum albumin below 35g/l had persistent low serum
273 C3(Li et al. 2023a). Previous studies have also suggested that low serum C3 predicted poor
274 kidney outcomes (Tsai et al. 2019). Besides, Rajasekaran et al. noted that complement markers
275 in kidney biopsies of IgAN patients were related to disease activity and predicted poor kidney
276 prognosis. With regard to the composition of the pathology types in the NDKD and DN+NDKD
277 group, we supposed that the mechanisms of MN and IgAN have complement involvement,
278 which influences the prognosis of the NDKD group and differences in the composition of
279 pathologic types between the two groups also led to different prognostic factors in the NDKD
280 and DN+NDKD groups. That, to some extent, could give some explanation of why lower C3

281 predicts ESKD in NDKD group.
282 Additionally, ALB, Scr, IgM, SBP and age were possible risk elements for the outcome in
283 patients with DN+NDKD. Beside the traditional risk factors, we noted the appearance of IgM.
284 Prior research had found that IgM may cause damage through activation of the glomerular
285 thylakoid complement cascade mediated by the classical immune complex (Mubarak & Kazi
286 2012). Al Romaili, D. M. found that IgM deposition in minimal change disease(MCD) showed
287 statistical association with CKD and IgM may play a role in MCD (Al Romaili et al. 2019). While
288 there was only one case of this type in the DN+NDKD group we studied. Further studies are
289 needed to verify the causal relationship between elevated IgM levels and kidney prognosis.
290 Taken together, these indicators were associated with declining kidney function.
291 Predictive models for diabetes-related kidney disease have been developed by many
292 researchers. But most of the models are not applicable to the Chinese population due to patient
293 populations, and study methodology. Riphagen et al. chose two clinical end points: development
294 of (micro)albuminuria and progressive kidney function loss (Riphagen et al. 2015). The inclusion
295 population of Anderson et al. included some patients without diabetes mellitus (Anderson et al.
296 2021). Meanwhile, Chen, S. et al. focused on analyzing the risk factors for three different
297 endpoint events by constructing Cox regression models (Chen et al. 2022). Strengths of the
298 present study are that it focused on subgroup analysis of T2DM populations, explored different
299 prognostic factors for DN, NDKD and DN+NDKD, and established three group different
300 prediction models for Chinese populations.
301 There are some limitations of this study and the analysis of the results may be biased. We
302 analyzed risk factors affecting prognosis using only a single-center cohort of individuals from
303 China. Because the epidemiology of T2DM patients with CKD shows significant global variation,
304 it may affect the generality of the application, but it may be useful to physicians in the region in
305 their daily practice. We hope to follow up with a multi-center, large sample size study. Next, in
306 our cohort, there was insufficient follow-up time for some patients, but it is emphasized that the
307 majority of patients enrolled in our study were not newly diagnosed with diabetes at the start of
308 the follow-up period. This aspect partially mitigated the limitations of our relatively short follow-
309 up duration. Additionally, one of the inevitable problems with clinical retrospective studies is the
310 presence of bias: exclusion of patients due to excessive missing information may create a
311 selection bias. In recent years, the use of new drugs has greatly improved the prognosis of DN,
312 thus further comparisons of the prognosis of the three groups after treatment are needed.
313

314 **Conclusions**

315 In conclusion, this respective single-center cohort research based on a Chinese population
316 demonstrated that the risk of endpoint events differed among DN, NDKD, and DN+NDKD
317 groups. Patients with DN were susceptible to getting the kidney endpoints. In the T2DM patients
318 with CKD, it has been found that, low serum ALB, Hb, higher glycosylated hemoglobin and
319 increased SBP, were independent risk parameters for the occurrence of endpoint events.
320 Therefore, it is crucial to focus on the DN group and implement early preventive or therapeutic

321 measures in order to delay the occurrence of these endpoints.

322

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325

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Figure 1

Flow chart of participant selection in this study.

From the 315 patients in our single center, we screened patients who met the inclusion criteria. A total of 268 participants were included and divided into three groups based on kidney puncture results.

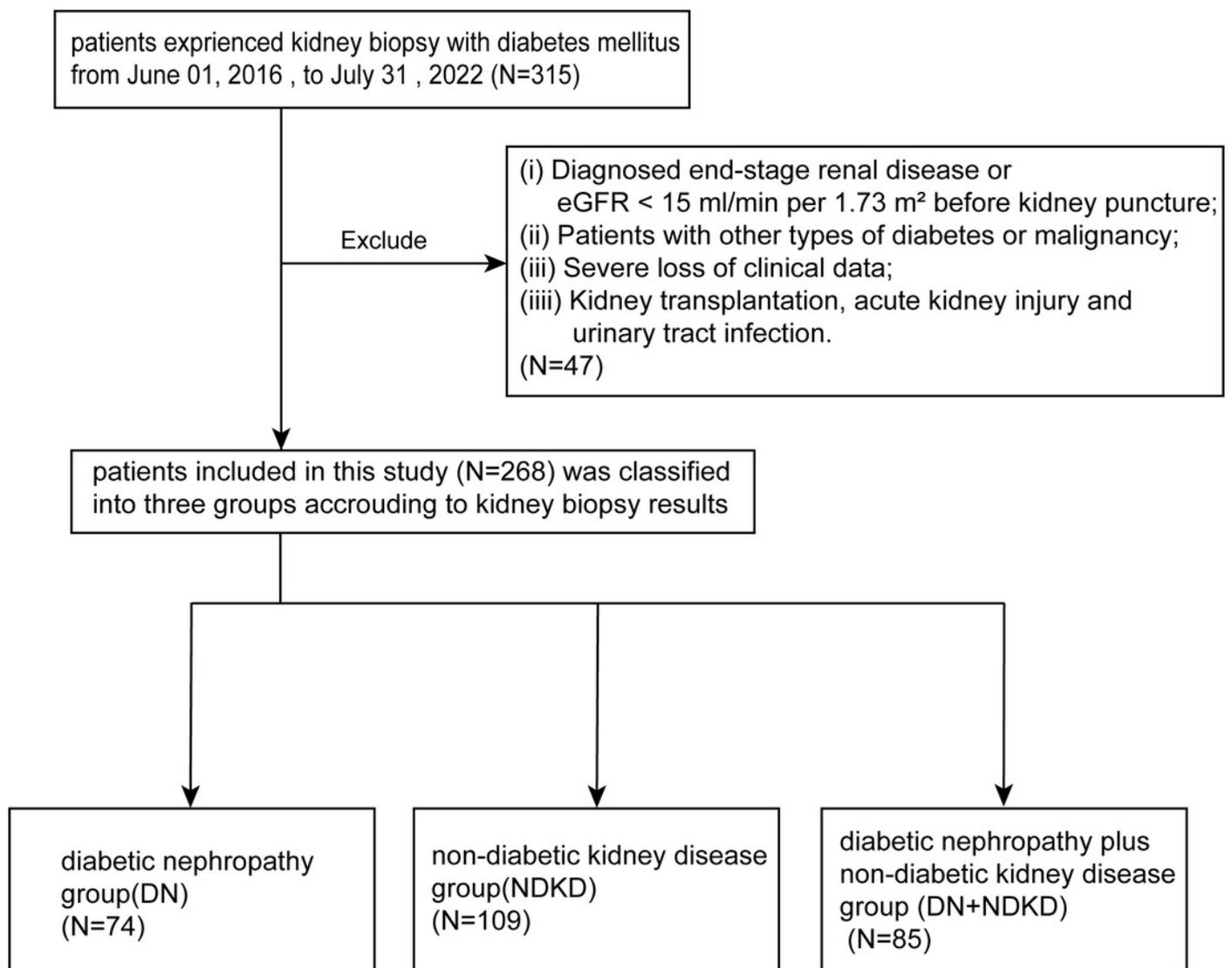


Table 1 (on next page)

Baseline characteristics and drug treatment of patients in the DN group, NDKD group and DN+NDKD group.

1

Characteristic	Overall(n=268)	DN(n=74)	NDKD(n=109)	DN+NDKD (n=85)	p-value
Age(median [IQR])	52.50(15.00)	50.00(18.00)	54.00(15.00)	53.00(12.00)	0.133
Gender = male (%)	180.00 (67.20)	55.00(74.30)	59.00(54.10)	66.00(77.60)	0.001**
Cigarette (%)	60.00 (22.40)	21.00(28.40)	18.00(16.50)	21.00(24.70)	0.138
family history of diabetes(%)	23.00(8.58)	12(16.22)	7(6.42)	4(4.71)	0.021*
RASi (%)	184 (68.70)	53.00(71.60)	80.00(73.40)	51.00(60.00)	0.101
Immunosuppressant(%)	23.00 (8.60)	3.00 (4.10)	16.00 (14.70)	4.00 (4.70)	0.013*
Glucocorticoid (%)	51 (19.00)	1.00 (1.40)	40.00(36.70)	10.00 (11.80)	0.000**
Insulin (%)	122(45.50)	46.00 (62.20)	37.00(33.90)	39.00(45.90)	0.000**
Follow-up time(mean (SD))	16.44(13.74)	15.32(10.00) ^a	16.44 (19.00) ^b	15.00(11.00) ^a	0.009**
The duration of DM (median [IQR])	36.00(82.25)	60.00(111.00) ^a	24.00(57.00) ^b	42.00(79.50) ^a	0.007**
DR(%)	43.00(16.00)	27.00(36.50)	4.00(3.70)	12.00(14.10)	0.000**
Sbp(mean (SD))	138.43 (22.81)	141.70(22.74)	136.97(21.30)	137.46(24.66)	0.348
Dbp(mean (SD))	86.93(13.68)	87.99(12.73)	86.97(13.76)	85.94(14.46)	0.643
Hb (mean (SD))	125.82(23.73)	122.01(5.75) ^a	129.69(20.61) ^b	125.04(25.05)	0.048*
RBC (median [IQR])	4.29(1.00)	4.16(1.13) ^a	4.34(1.01) ^b	4.30(1.13)	0.043*
24h proteinuria(median [IQR])	2484.76(4431.73)	3438.53(5331.20)	2148.00(4583.10)	1925.80(3860.20)	0.055
24h urine protein >3.5g(%)	87.00(32.46)	30.00(40.54)	34.00(31.19)	23.00(27.06)	0.126
Urinary	30.15(49.84)	25.15(28.15)	41.30(87.95)	14.20(63.62)	0.000

sediment)) ^a) ^b) ^a	4 ^{**}
RBC(median[IQR])					0 ^{**}
BUN (median[IQR])	7.29(3.97)	8.21(4.53) ^a	6.60(3.00) ^b	7.97(4.08) ^a	0.00
Scr (median[IQR])	103.50(77.75)	115.00(84.38) ^a	89.00(48.50) ^b	122.00(100.50) ^a	0.00
EGFR(median[IQR])	64.95(48.35)	59.10(50.48) ^a	78.00(41.95) ^b	55.50(44.25) ^a	0.00
ALB(median[IQR])	37.80(13.48)	34.95(11.55)	38.80(15.65)	39.70(10.90)	0.13
Blood glucose(median[IQR])	8.32(4.47)	8.98(5.49)	8.26(3.80)	8.23(4.41)	0.16
Glycosylated hemoglobin(median [IQR])	6.85(1.50)	7.60(1.80) ^a	6.60(1.10) ^b	6.80(1.00) ^b	0.00
TC(median[IQR])	4.77(1.99)	4.90(1.74)	4.80(2.20)	4.58(1.80)	0.23
TG(median[IQR])	2.51(2.48)	2.72(3.01)	2.61(2.47)	2.39(2.23)	0.26
HDL-C(median[IQR])	0.98(0.38)	0.96(0.34)	1.04(0.38) ^a	0.93(0.39) ^b	0.04
LDL-C(median[IQR])	2.65(1.27)	2.60(1.10)	2.74(1.32)	2.45(1.42)	0.44
IgG (mean (SD))	9.87(3.74)	9.70(4.40)	9.60(5.55)	10.49(3.94)	0.05
IgA (median [IQR])	2.54(1.43)	2.45(1.49)	2.64(1.61)	2.53(1.19)	0.73
IgM (median[IQR])	0.97(0.60)	0.98(0.61)	1.05(0.68)	0.89(0.37)	0.10
C3 (median[IQR])	0.96(0.25)	0.92(0.23) ^a	1.03(0.26) ^b	0.94(0.18) ^a	0.00
C4 (median[IQR])	0.26(0.09)	0.27(0.09)	0.26(0.10)	0.25(0.09)	0.60

2 Notes: Data are presented as medians with ranges, or counts and percentages. ^a and ^b represent
3 instances where there are significant differences between ^a and ^b. P* $<$ 0.05, P** $<$ 0.01, P*** $<$ 0.001
4 Abbreviations: DR, diabetic retinopathy; DM, diabetes mellitus; UACR, urinary albumin/creatinine
5 ratio; Sbp, systolic blood pressure; Dbp, diastolic blood pressure; Hb; BUN, Blood Urea Nitrogen
6 ; Scr, serum creatinine; EGFR, estimated glomerular filtration rate; ALB, albumin; TC, total
7 cholesterol; TG, triglyceride; HDL-C, high density lipid-cholesterol; LDL-C, low density lipid-

8 cholesterol.

9

Figure 2

Pathologic manifestations of DN.

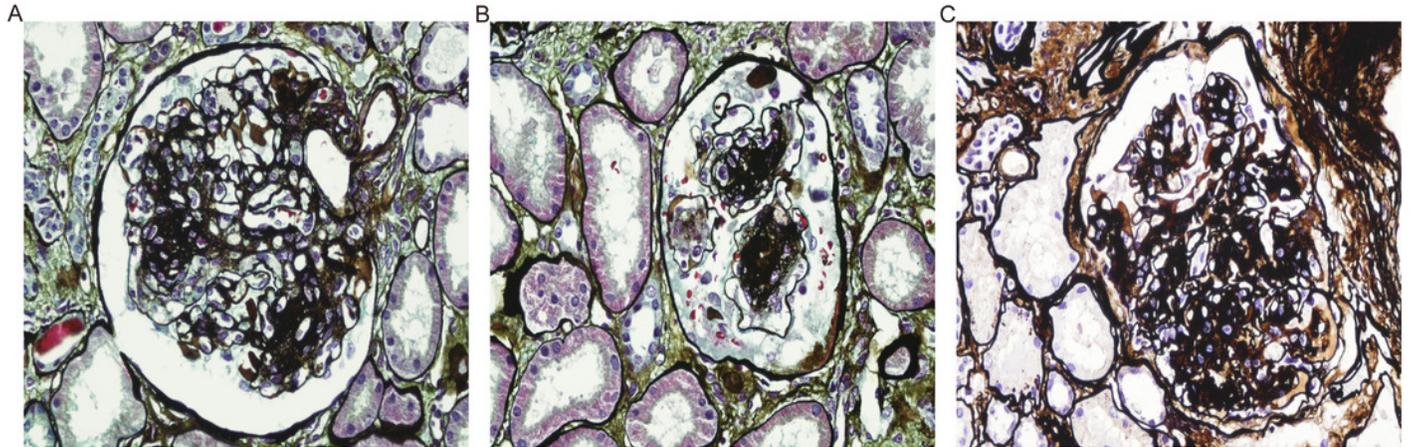


Table 2 (on next page)

Comparison of pathological characteristics between NDKD and DN+NDKD groups.

1

Pathological characteristic	NDKD (109)	DN+NDKD (85)
IgA nephropathy	26	15
Membranous nephropathy	45	12
Hypertensive nephropathy	10	34
Henoch-Schoenlein purpura nephritis	5	0
Obesity-related nephropathy	4	0
Focal segmental glomerulosclerosis	3	3
light chain deposition disease	1	0
Kidney amyloidosis	1	0
Tubulointerstitial nephritis	2	8
Thrombotic Microangiopathy	2	1
Hepatitis B virus-related nephropathy	1	1
Sclerosing glomerulonephritis	3	2
Minimal change disease	2	1
Proliferative glomerular lesions	4	1
Acute tubular necrosis	0	2
HCV associated glomerulonephritis	0	1
Post-infectious glomerulonephritis	0	1
Crescentic glomerulonephritis	0	3

2

Figure 3

Comparison of renal survival rate in the DN group, NDKD group and DN+NDKD group.

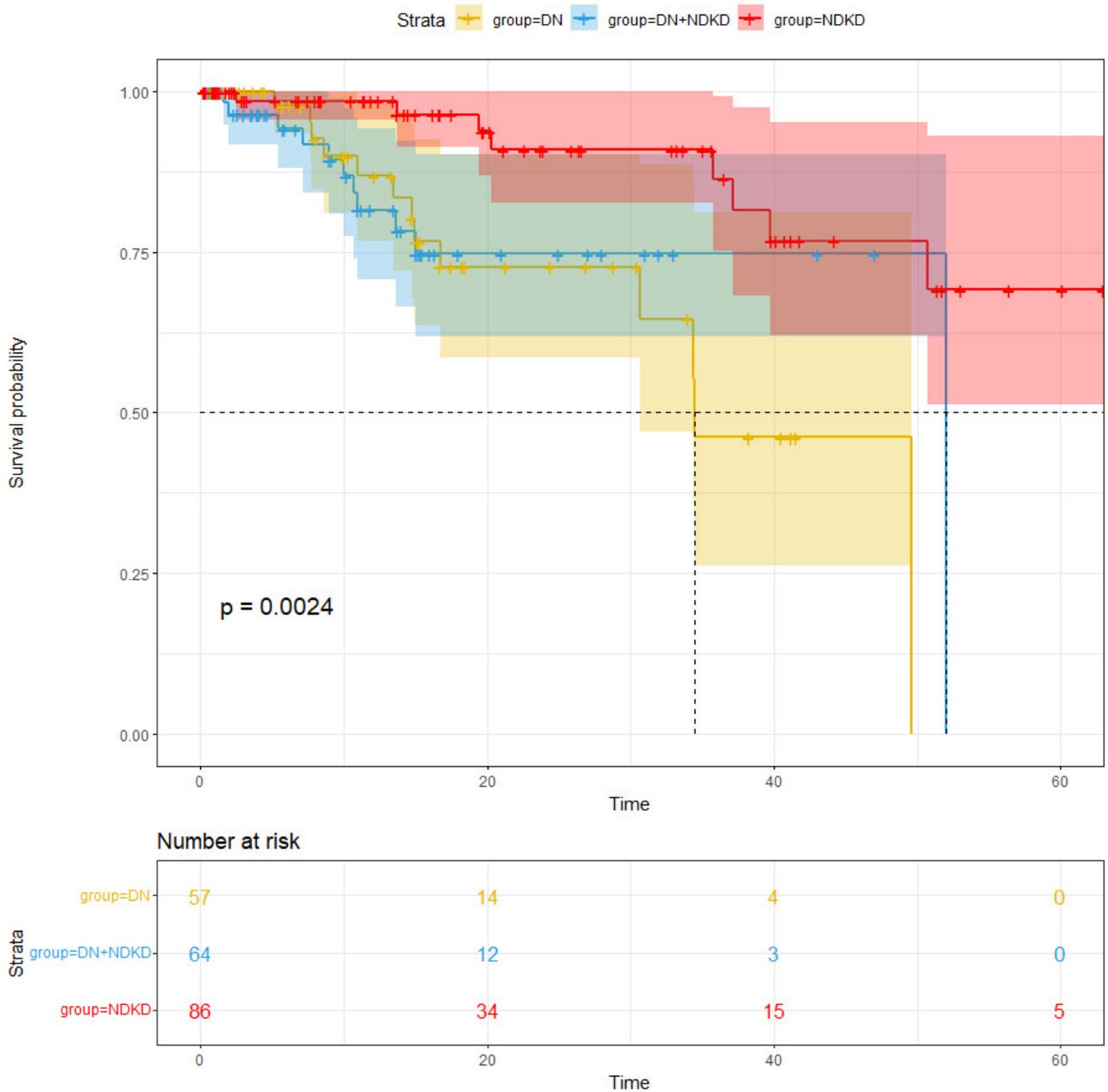


Table 3 (on next page)

Univariate Cox regression analyses for endpoints.

1 (A) The univariate Cox analysis results of the DN and NDKD group

Characteristic	DN		NDKD	
	HR(95%CI)	p-value	HR(95%CI)	p-value
Age	1.008(0.957-1.062)	0.759	1.026(0.958-1.100)	0.461
Gender(male)	1.052(0.283-3.904)	0.940	0.686(0.166-2.830)	0.602
SBP	1.047(1.019-1.076)	0.001**	0.99(0.957-1.025)	0.580
Hb	0.935(0.899-0.973)	0.001**	0.976(0.939-1.015)	0.221
Urinary sediment RBC	1.027(1.008-1.048)	0.007**	0.993(0.975-1.011)	0.452
Serum Alb	0.792(0.704-0.893)	0.000***	0.93(0.859-1.006)	0.071
Scr	1.013(1.005-1.022)	0.002**	0.997 (0.98-1.015)	0.777
24h urine protein	1.00(1.00-1.00)	0.047*	1.00(1.00-1.00)	0.050*
TC	1.093(0.721-1.657)	0.676	1.004(0.652-1.545)	0.986
TG	0.813(0.65-1.017)	0.07	1.01(0.818-1.246)	0.928
HDL	1.606(0.44-5.866)	0.473	2.389(0.325-17.535)	0.392
LDL	1.163(0.642-2.107)	0.618	0.761(0.382-1.513)	0.436
IgG	0.757(0.588-0.974)	0.03*	0.759(0.581-0.991)	0.043*
IgA	1.428(0.818-2.492)	0.21	0.586(0.263-1.306)	0.191
IgM	0.523 (0.16-1.709)	0.283	1.568(0.33-7.462)	0.572
C3	0.173 (0.005-6.185)	0.336	0.020(0.000-0.937)	0.046*

2

3 (B) The univariate Cox analysis results for the DN+NDKD group and whole cohort

Characteristic	DN+NDKD		All patients	
	HR(95%CI)	p-value	HR(95%CI)	p-value
Age	1.010(0.942-1.083)	0.783	1.006(0.973-1.041)	0.725
Gender(male)	0.398 (0.105-	0.177	0.845(0.408-	0.65

	1.516)		1.749)	
SBP	1.032(1.003-1.062)	0.028*	1.024(1.009-1.04)	0.002**
Hb	0.976(0.953-0.999)	0.042*	0.963(0.947-0.979)	0.000***
Urinary sediment RBC	1.00(1.00-1.00)	0.769	1.00(1.00-1.00)	0.728
Serum Alb	0.931(0.872-0.993)	0.029*	0.93(0.895-0.967)	0.000***
Scr	1.004(1.002-1.007)	0.003**	1.003(1.001-1.005)	0.001**
24h urine protein	1.00(1.00-1.00)	0.463	1.00(1.00-1.00)	0.024*
TC	1.056(0.729-1.528)	0.774	0.952(0.765-1.185)	0.662
TG	0.947(0.728-1.232)	0.684	0.933(0.818-1.065)	0.304
HDL	1.734(0.277-10.838)	0.556	1.377(0.511-3.708)	0.527
LDL	1.214(0.771-1.912)	0.402	0.966(0.699-1.334)	0.833
IgG	0.949(0.789-1.141)	0.579	0.897(0.804-1.002)	0.054
IgA	1.07(0.583-1.964)	0.828	0.876(0.64-1.199)	0.41
IgM	4.049(1.537-10.668)	0.005**	1.992(1.119-3.549)	0.019*
C3	0.142(0.002-11.771)	0.386	0.062(0.008-0.49)	0.008**

Table 4 (on next page)

Multivariate Cox regression analyses for endpoints.

1 (A)DN

Characteristic	HR(95%CI)	p-value	the Chi-square values	p-value
ALB	0.685(0.559-0.839)	<0.001***	36.846	<0.001***
24h proteinuria	0.999(0.999-1.000)	0.006*		
SBP	1.047(1.006-1.089)	0.024*		
age	0.890(0.802-0.988)	0.028*		

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4 (B)NDKD

Characteristic	HR(95%CI)	p-value	the Chi-square values	p-value
Urinary sediment RBC	0.975(0.948-1.003)	0.082	9.896	0.042*
24h proteinuria	1.000(1.000-1.001)	0.019*		
IgG	0.803(0.584-1.105)	0.178		
C3	0.001(0.000-0.356)	0.021*		

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7 (C) DN+NDKD

Characteristic	HR(95%CI)	p-value	the Chi-square values	p-value
ALB	0.828(0.724-0.947)	0.006**	53.626	<0.001***
Scr	1.011(1.005-1.018)	0.001**		
IgM	13.708(3.611-52.034)	<0.001***		
SBP	1.050(1.007-1.094)	0.021*		

age	0.851(0.756-0.958)	0.008**
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10 (D)Total patients

Characteristic	HR(95%CI)	p-value	the Chi-square values	p-value
SBP	1.018(1.002-1.035)	0.025*	50.029	<0.001***
Hb	0.979(0.961-0.997)	0.023*		
C3	0.133(0.014-1.228)	0.075		
ALB	0.952(0.910-0.996)	0.032*		
Glycosylated hemoglobin	1.338(1.080-1.658)	0.008**		

11 p* < 0.05; P** < 0.01; P***<0.001

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