Hematuria is a risk factor towards end-stage renal disease - A propensity score analysis

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Abstract

Background: It remains unclear whether microscopic hematuria accelerates the progression of chronic kidney disease (CKD). Since the risk of microscopic hematuria was expected very weak, we employed a propensity score analysis.

Methods: A retrospective CKD cohort of 803 participants in a single institution was analyzed. The degree of microscopic hematuria was scaled as 0, 0.5, 1, 2, 3 for negative, 1+, 2+, 3+ in the urinary qualitative analysis by dipstick, respectively. Time-averaged microscopic hematuria (TA-MH) was calculated by trapezoidal rule. The propensity score was estimated using 23 baseline covariates by multivariate binary logistic regression for the threshold of TA-MH at ≥ 0.5 or ≥ 1.0. Kaplan-Meier analysis after propensity score matching was also examined.

Results: The incidence rate of end-stage renal disease (ESRD) was 33.9 per 1,000 person-years over median follow-up of 4.3 years. A Cox regression analysis stratified by quintile on the propensity scores showed that TA-MH ≥ 0.5 was a risk for ESRD (HR 1.72, 95% CI 1.08-2.75; p = 0.023) but not TA-MH ≥ 1.0 (HR 1.34, 95% CI 0.76-2.40; p = 0.315). Kaplan-Meier analysis after propensity score matching reproduced the similar results (TA-MH ≥ 0.5, HR 1.80, 95% CI 1.03-3.12; p = 0.046; TA-MH ≥ 1.0, HR 1.73, 95% CI 0.84-3.54; p = 0.145). Kaplan-Meier analysis after propensity score matching was also examined.

Conclusions: Mild degree of microscopic hematuria is a risk factor for CKD progression to ESRD. The present result may help clinicians to recognize the CKD patients at higher risk due to the presence of microscopic hematuria.

Introduction

Early recognition and intervention against risk factors responsible for the progression of chronic kidney disease (CKD) are expected to improve renal outcomes of patients at risk [1]. Major risk factors of subsequent incidence of end-stage renal disease (ESRD) are hypertension, proteinuria and anemia in addition to preceding kidney dysfunction [2,3]. However, the second-line risk predictors remains to be clarified; candidates are hyperuricemia, hyperphosphatemia, metabolic acidosis, dyslipidemia, hypoalbuminemia, etc. Although hematuria has long been appreciated as a clinical presentation of the kidney diseases, it remains to be elucidated whether hematuria is a risk factor for progression of CKD. Recently, it has been suggested that young persons with persistent isolated microscopic hematuria may indeed have an increased risk for ESRD as compared with individuals without this condition [4]. On the other hand, there are several contradictory results, ruling out the impact of microscopic hematuria on the outcome of IgA nephropathy [5,6]. Therefore, microscopic hematuria has not reached consensus as a risk factor of ESRD [7].

The most probable reason for the conflicting results may lie on the small effect of microscopic hematuria compared with other major confounders such as proteinuria and hypertension. Thus, a propensity score analysis was employed, which is increasingly being used to estimate causal effects in the observational studies because one can replicate the prospective randomized controlled trial by minimizing baseline confounding [8]. Another important property long overlooked is the time-varying nature of microscopic hematuria in the clinical course of the CKD. In this study time-averaged microscopic hematuria (TA-MH) in the follow-up was calculated and used to investigate whether microscopic hematuria may be a risk factor of CKD progression.

Patients and methods

Study protocol and ethical statement

The retrospective CKD cohort already reported (n = 803) was used [9-12], and in the current study the propensity score analysis was used in such a way as used in the previously published articles [9,12]. Inclusion criteria consisted of CKD stage 3 to 4, age 20 to 84 years and follow-up period ≥ 1 year. On the other hand, patients with nephrotic syndrome, malignancy, obstructive nephropathy, acute kidney injury and gout were excluded. All the patients were followed up to 6 years until censoring or reaching the initiation of dialysis. The present study was approved by the institutional review board (IRB) in the Teikyo University Review Board #14-115 and was executed in accordance with the principle of the Helsinki Declaration. Written informed consent was waived after approval of IRB and the patient records and information were anonymized and de-identified prior to analysis.

Parameters analyzed

The demographic characteristics included sex, age, body mass index (BMI), original kidney diseases (diabetic nephropathy, chronic

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Key words: microscopic hematuria, time-averaged microscopic hematuria, chronic kidney disease, end-stage renal disease, propensity score analysis

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glomerulonephritis, nephrosclerosis and other diseases) and systolic blood pressure (SBP). Diabetic nephropathy was defined as treated with antidiabetic agents or as having a long history of the disease. Chronic glomerulonephritis was defined as having biopsy-proven diagnosis or as absence of diabetes mellitus or hypertension. Nephrosclerosis was defined as treated with antihypertensive agents or as having a long history of hypertension. Other disease included polycystic kidney disease, tubulointerstitial nephritis and undetermined causes of CKD.

Blood parameters included hemoglobin (Hb), white blood cell (WBC), platelet (Plt), albumin (Alb), uric acid (UA), sodium (Na), potassium (K), chloride (Cl), Na-Cl (as a surrogate of HCO₃⁻), albumin-corrected calcium (cCa), inorganic phosphorus (P), low-density lipoprotein cholesterol (LDL-C) and C-reactive protein (CRP). Urine parameters included spot urine proteinuria (expressed as gram per gram creatinine excretion) and spot urine hematuria by dipstick (coded as four grades of 0 to 3 according to 0, 1+, 2+, and 3+ and as 0.5 ±). Due to the distribution, C-reactive protein and proteinuria were log-transformed for analyses.

Blood was tested using hematology autoanalyzer (Sysmex XE-5000, Kobe, Japan) and blood chemistry parameters were measured by routine measurements using autoanalyzer (LABOSPECT 008, Hitachi High-Technologies Corporation, Tokyo, Japan). Creatinine concentration in serum and urine was measured by an enzymatic method and urinary protein concentration measured by a pyrocatechol violet-metal complex assay method. Hematuria measured at every visit was recorded until censoring or reaching estimated GFR 5 ml/min/1.73 m² and calculated as TA-MH by trapezoidal rule [9,12]. Estimated GFR was evaluated using the Modification of Diet in Renal Disease (MDRD) study equation for Japanese population [13]. The grade of CKD was classified based on the Kidney Disease Outcomes Quality Initiative (KDQI) practice guidelines [1].

Use of angiotensin converting enzyme inhibitor or angiotensin II receptor blocker (combined as RASi) and use of antplatelet drugs were recorded as yes (coded as 1) or no (coded as 0). The baseline covariates used for the propensity score estimate modeling became 23 in total.

Endpoints of renal outcomes

A primary endpoint was defined as the incidence of ESRD (initiation of hemodialysis or peritoneal dialysis). Death was treated as censoring because the present study focused on the effect of hematuria on the subsequent ESRD rather than the risk of mortality [11]. Competing risk method was not employed due to a small portion of death censoring (n = 10).

A propensity score analysis

The threshold of TA-MH in the follow-up was tested at 0.5 and 1.0. The probability to reach above or equal to the threshold was determined by a multivariate binary logistic regression using the aforementioned 23 baseline covariates. Since the distribution of propensity score of two groups differed widely, patients whose propensity scores not overlapped between two groups were trimmed, then the remaining subsample was re-stratified on the quintiles of the propensity scores [14,15]. A stratified Cox proportional hazards model was conducted in the substrata on the quintiles of the propensity scores [8]. Then, a pooled hazard ratio (HR) of the higher group of TA-MH was obtained as a crude HR.

Participants separated by the threshold of TA-MH (0.5 and 1.0) were matched using a greedy method with a 1:1 pair. The caliper size was first set at 0.25 times standard deviation of the propensity scores and gradually narrowed down until the balance of baseline 23 covariates were achieved between 2 groups [8]. The model of assignment was estimated by c-statistics and the balance between two groups was checked by paired comparison tests [16]. A time-to-event survival analysis was examined by a Kaplan-Meier analysis with stratified log-rank test [8,17].

Statistical analyses

Values for categorical variables are given as number (percentage) and values for continuous variables are given as mean ± standard deviation or median [interquartile range] depending on the distribution. C-statistics for the accuracy of the propensity score models were obtained by the area under the receiver operating characteristic (AUROC) curve for the threshold [18]. For a Cox proportional hazards model, any covariate was tested for its proportional hazards assumption using both a time-dependent Cox regression and a Schoenfeld residual plot. Stratified log-rank test following Kaplan-Meier analysis after propensity score matching was performed by stratification on quintile of the matched subsample. Statistical analyses were performed using SPSS version 22 (IBM, Tokyo) and STATA version 14 (StataCorp LP, College Station, TX, USA). A p value less than 0.05 was considered statistically significant.

Results

Clinical characteristics of the cohort

During the follow-up period of median 4.3 [interquartile range 2.6-5.7] years, 110 out of 803 patients progressed to ESRD. The incidence rate was 33.9 per 1,000 person-years. The patients (n = 236) having mild degree of hematuria (TA-MH ≥ 0.5) in the follow-up showed the following baseline characteristics as shown Table 1. Estimated GFR was lower and chronic glomerulonephritis was more prevalent. Hemoglobin and serum albumin were lower and serum uric acid and serum phosphorus were higher. In addition, proteinuria and microscopic hematuria were greater in degree. The patients (n = 108) having greater degree of hematuria (TA-MH ≥ 1.0) in the follow-up showed the following baseline characteristics as shown Table 2. Chronic glomerulonephritis was more prevalent but diabetic nephropathy less prevalent. Hemoglobin and serum albumin were lower while serum uric acid and serum phosphorus were higher. Proteinuria and microscopic hematuria were also greater.

Taken altogether, the patients presenting greater degree of microscopic hematuria, either 0.5 or 1.0 in the follow-up, were confounded with established risk factors such as anemia, hypoalbuminuria, hyperuricemia, hyperphosphatemia and proteinuria. It is no way questionable that hematuria did not show an independent risk factor by standard Cox regression analysis [9] (supplementary information).

A propensity score analysis

Stratified Cox proportional hazards model: A propensity score analysis could overcome the aforementioned problem. The threshold of TA-MH was examined at 0.5 and 1.0. A subsample after trimming non-overlapped patients was re-stratified on the quintiles of the propensity scores and then subjected to survival analysis by a stratified Cox proportional hazards model. Proportional hazards assumption was not violated and multicollinearity was not seen. The result disclosed that the significantly high HR was found at 0.5 of the threshold of TA-MH (HR 1.72, 95% CI 1.08-2.75, p = 0.023). On the other hand, there was no
Propensity score matching: Following propensity score matching, all the baseline covariates were well balanced by paired analysis when TA-MH 0.5 and 1.0 was used, respectively (Table 1 and 2). C-statistics estimated by the area under the ROC curve were all greater than 0.8 (Table 4), suggesting the high discrimination accuracy [16]. Then, two groups divided by the threshold of TA-MH were subjected to a Kaplan-Meier analysis (Table 4) that were plotted in Figure 1a and 1b. The patients showing TA-MH ≥ 0.5 in the follow-up showed significantly higher HR for ESRD (HR 1.80, 95%CI 1.03-3.12, p = 0.046), whereas the threshold of TA-MH ≥ 1.0 did not reach statistical significance (Table 4).

Discussion

In the present study we could show the significant impact of microscopic hematuria in the follow-up on the subsequent incidence of ESRD by applying the propensity score analysis. A pooled crude HR of TA-MH greater than or equal to 0.5 versus the lower group was significantly as high as 1.72. To our astonishment, the target value in the follow-up should be below ± in dipstick test. The present study is the first report elucidating that even the mild hematuria in the follow-up underscores a risk factor of CKD progression, which was elucidated by applying the propensity score analysis.

Strength of the present study is two folds; calculation of TA-MH in the follow-up and the propensity score analysis. In most previous reports, hematuria was tested only once at baseline. Microscopic hematuria is likely to change with a clinical course of any types of CKD, regarded as one of the time-varying parameters. Thus, TA-MH is better for assessing the impact of microscopic hematuria on CKD progression. In practice, arithmeticly calculated time-averaged values have been utilized in time-to-event survival analyses [19-22]. Meanwhile, a propensity score analysis has come into rapid use in the clinical medicine because it can approximate randomized controlled trials using retrospective observational cohorts. The method also enables one to investigate the causal effect which cannot be otherwise executed due to serious issues such as ethical problems and inability of intervention. Propensity score analysis by use of time-averaged values of test parameters has a great advantage to freely scrutinize the threshold of target values as demonstrated here and previously [9].

Although microscopic hematuria has been considered as a diagnostic marker of any kidney disease including glomerular diseases, its impact on CKD progression has not been investigated in many large

Table 1. Balance of baseline covariates before and after propensity score matching for TA-MH 0.5

| Characteristics | Before matching (n = 803) | After matching (n = 270) |
|-----------------|--------------------------|-------------------------|
| **Blood**       |                          |                         |
| Hb (g/dL)       | 13.0 ± 1.9               | 12.6 ± 2.0              | 12.7 ± 1.8 | 12.6 ± 2.0 | 0.786 |
| WBC (×10^9/L)   | 64.7 ± 21.3              | 67.6 ± 21.7             | 0.078      | 65.7 ± 23.4 | 66.2 ± 21.3 | 0.879 |
| Plt (×10^12/L)  | 21.9 ± 6.4               | 22.2 ± 7.7              | 0.515      | 21.6 ± 6.2 | 21.8 ± 6.6 | 0.792 |
| Alb (g/dL)      | 4.0 ± 0.5                | 3.8 ± 0.5               | < 0.001    | 3.8 ± 0.5 | 3.9 ± 0.5 | 0.176 |
| UA (mg/dL)      | 6.4 ± 1.4                | 6.7 ± 1.4               | 0.020      | 6.5 ± 1.5 | 6.6 ± 1.5 | 0.602 |
| Na (mEq/L)      | 140.8 ± 2.7              | 140.7 ± 2.6             | 0.682      | 140.7 ± 2.4 | 140.7 ± 2.8 | 0.818 |
| K (mEq/L)       | 4.5 ± 0.5                | 4.5 ± 0.6               | 0.759      | 4.5 ± 0.5 | 4.5 ± 0.6 | 0.875 |
| Ca (mg/dL)      | 35.5 ± 2.4               | 35.1 ± 2.7              | 0.053      | 35.2 ± 2.1 | 35.2 ± 2.7 | 0.961 |
| P (mg/dL)       | 8.8 ± 0.5                | 8.9 ± 0.5               | 0.009      | 9.0 ± 0.5 | 8.9 ± 0.5 | 0.375 |
| CRP (mg/dL)     | 0.08[0.04-0.18]          | 0.10[0.05-0.24]         | 0.045      | 0.08[0.05-0.17] | 0.10[0.05-0.24] | 0.581 |
| LDLC (mg/dL)    | 112.2 ± 31.0             | 107.6 ± 29.0            | 0.050      | 107.8 ± 30.7 | 106.8 ± 29.4 | 0.798 |
| **Spot urine**  |                          |                         |
| TPU/CrU (g/g Cr)| 0.36[0.16-0.99]          | 0.73[0.25-1.75]         | < 0.001    | 0.49[0.22-1.55] | 0.48[0.19-1.36] | 0.496 |
| **Drug use**    |                          |                         |
| RASi, Yes (%)   | 319 (56.3)               | 118 (50.0)              | 0.120      | 80 (59.3) | 73 (54.1) | 0.406 |
| Antiplatelet drugs, Yes (%) | 60 (10.6) | 25 (10.6) | 1.000 | 11 (8.1) | 16 (11.9) | 0.297 |

Note: Values for categorical variables are given as number (percentage); values for continuous variables are given as mean ± standard deviation or median [interquartile range]. For statistical analyses, CRP and TPU/CrU were log-transformed. Conversion factors for units: creatinine in mg/dL to μmol/L, x 88.4; uric acid in mg/dL to μmol/L, x 59.48.

*Unpaired t test or McNemar test

Abbreviations: TA-MH, time-averaged microscopic hematuria; eGFR, estimated glomerular filtration rate; DMN, diabetic nephropathy; CGN, chronic glomerulonephritis; BMI, Body Mass Index; SBP, systolic blood pressure; Hb, hemoglobin; WBC, white blood cells; Pth, Platelet; Alb, albumin; UA, uric acid; Na, sodium; K, potassium; Cl, chloride; cCa, albumin-corrected calcium; P, phosphorus; LDL-C, low-density lipoprotein cholesterol; TPU/CrU, urine total protein divided by urine creatinine; UB_score, urine blood score; RASI, RAS inhibitor.
Shima T (2017) Hematuria is a risk factor towards end-stage renal disease - A propensity score analysis

**Table 2. Balance of baseline covariates before and after propensity score matching by TA-MH 1.0.**

| Characteristics | Before matching (n = 803) | After matching (n = 150) |
|-----------------|---------------------------|--------------------------|
| TA-MH < 1       | TA-MH ≥ 1                 | p value†                |
| n = 695         | n = 108                   |                          |
| Age (y)         | 62.4 ± 12.7               | 0.126                    |
| eGFR (mL/min/1.73m²) | 41.1 ± 13.1              | 0.889                    |
| Male (%)        | 442 (63.6)                | 0.087                    |
| DMN (%)         | 171 (24.0)                | 0.014                    |
| CGN (%)         | 139 (20.0)                | 0.002                    |
| BMI (kg/m²)     | 24.5 ± 4.3                | 0.035                    |
| SBP (mmHg)      | 137.8 ± 20.9              | 0.075                    |
| TA-MH < 1       | TA-UA ≥ 1                | p value†                |
| n = 75          | n = 75                   |                          |
| Age (y)         | 62.0 ± 14.1               | 0.126                    |
| eGFR (mL/min/1.73m²) | 38.9 ± 13.0              | 0.889                    |
| Male (%)        | 38 (50.7)                 | 0.087                    |
| DMN (%)         | 16 (21.3)                 | 0.014                    |
| CGN (%)         | 18 (24.0)                 | 0.002                    |
| BMI (kg/m²)     | 23.5 ± 4.4                | 0.035                    |
| SBP (mmHg)      | 137.4 ± 20.0              | 0.075                    |

**Blood Parameters**

|                         | Before matching (n = 803) | After matching (n = 150) |
|-------------------------|---------------------------|--------------------------|
| HB (g/dL)               | 12.9 ± 1.9                | 12.4 ± 2.1               |
| WBC (<10⁹/μL)           | 65.3 ± 21.1               | 65.7 ± 22.0              |
| PΙ (<10⁹/μL)            | 22.0 ± 6.6                | 22.0 ± 8.0               |
| Alb (g/dL)              | 4.0 ± 0.5                 | 3.8 ± 0.4                |
| UA (mg/dL)              | 6.4 ± 1.4                 | 6.8 ± 1.5                |
| Na (mEq/L)              | 140.7 ± 2.7               | 140.6 ± 2.9              |
| K (mEq/L)               | 4.5 ± 0.5                 | 4.7 ± 0.6                |
| Na-CI (mEq/L)           | 35.4 ± 2.4                | 35.0 ± 2.9               |
| cCa (mg/dL)             | 8.8 ± 0.5                 | 8.9 ± 0.5                |
| P (mg/dL)               | 3.4 ± 0.5                 | 3.5 ± 0.6                |
| CRP (mg/dL)             | 0.08[0.05-0.20]           | 0.07[0.05-0.17]          |
| LDL-C (mg/dL)           | 111.5 ± 31.1              | 109.4 ± 31.2             |
| Spot urine              |                           |                          |
| TPU/Cr (g/g Cr)         | 0.40[0.17-1.09]           | 0.70[0.27-1.64]          |
| UB_score                | 0.34 ± 0.61               | 1.39 ± 1.06              |
| Antiplatelet drugs, Yes (%) | 74 (10.6)             | 7 (9.3)                  |

**Drug use**

RASi, Yes (%) 74 (10.6) 7 (9.3) 0.749

**Notes:** Values for categorical variables are given as number (percentage); values for continuous variables are given as mean ± standard deviation or median [interquartile range]. For statistical analyses, CRP and TPU/Cr were log-transformed. Conversion factors for units: creatinine in mg/dL to μmol/L, x 88.4; uric acid in mg/dL to μmol/L, x 59.48.

*Unpaired t test or McNemar test

†Paired t test or McNemar test

**Abbreviations:** TA-MH, time-averaged microscopic hematuria; eGFR, estimated glomerular filtration rate; DMN, diabetic nephropathy; CGN, chronic glomerulonephritis; BMI, Body Mass Index; SBP, systolic blood pressure; HB, hemoglobin; WBC, white blood cells; PΙ, Platelet; Alb, albumin; UA, uric acid; Na, sodium; K, potassium; Cl, chloride; cCa, albumin-corrected calcium; P, phosphorus; LDLC, low-density lipoprotein cholesterol; TPU/Cr, urine total protein divided by urine creatinine; UB score, urine blood score; RASi, RAS inhibitor.

**Table 3. Stratified Cox regression for ESRD by thresholds of TA-MH.**

| Threshold of TA-MH | Patients after trimming (Total/Incident) | HR | 95% CI | p value† |
|-------------------|------------------------------------------|----|--------|----------|
| ≥ 0.5             | 765/108                                  | 1.72 | 1.08 - 2.75 | 0.023 |
| ≥ 1.0             | 717/96                                   | 1.34 | 0.76 - 2.40 | 0.315 |

**Abbreviations:** ESRD, end-stage renal disease; TA-MH, time-averaged microscopic hematuria; HR, hazard ration; CI, confidence interval

**Table 4. Kaplan-Meier analysis after propensity score matching.**

| Threshold of TA-MH | C-statistics | Before matching | After matching | p value (stratified log-rank) |
|--------------------|--------------|----------------|----------------|-------------------------------|
| 0.5                | 0.86         | 567/55         | 135/21         | 1.80 | 1.03 - 3.12 | 0.046 |
| 1                  | 0.90         | 695/88         | 72/12          | 0.75 | 0.84 - 3.54 | 0.145 |

**Abbreviations:** TA-MH, time-averaged microscopic hematuria; HR, hazard ratio; CI, confidence interval

epidemiological studies [23,24]. This fact may be in part due to the difficulty in assessing microscopic hematuria quantitatively [25,26]. Recently, Vivante et al. reported that persistent asymptomatic isolated microscopic hematuria was significantly associated with increased risk of ESRD over 22 years of follow-up in 1.2 million persons aged 16 through 25 years [4]. This study suggests that hematuria in the absence of a specific diagnosis can be a risk factor of ESRD and the classical term "benign hematuria" or "benign familial hematuria" may be a misnomer, especially in most cases where neither a histopathologic nor a genetic diagnosis is made [4]. Goto et al. also showed that mild hematuria was associated with increased risk of ESRD over 10 years of follow-up in 790 IgA nephropathy patients. Most recently, Lin et al. showed that microscopic hematuria was associated with increased risk of ESRD in 1,799 patients with stage 3 to 5 nondiabetic CKD. Whereas, there are several reports denying microscopic hematuria as a risk factor of CKD progression in patients with IgA nephropathy [5,6].
Hematuria could induce renal tubular epithelial cell damage by the cytotoxic, oxidant and inflammatory effects of hemoglobin-related molecules such as heme and iron [28]. Erythrocyte passage across the glomerular filtration barrier induces mechanical stress to erythrocyte, leading to leakage of hemoglobin from erythrocyte. Tubular epithelial cells can uptake hemoglobin in the urinary space via magalin-cubulin receptor system. Hemoglobin is then transformed into heme and globin [29]. Free heme is extremely toxic, promoting lipid oxidation, inducing apoptosis and releasing inflammatory chemokines [30]. This heme-induced oxidation and inflammation in tubular cells could be related to CKD progression in the patients presenting microscopic hematuria [28]. Future studies are needed to elucidate the mechanism how hematuria exacerbates kidney damage in CKD milieu. In any event the present finding may open up a new horizon for the impact of hematuria on the progression of CKD.

Some limitations of the present study should be mentioned. The first limitation is that we assessed microscopic hematuria only by dipstick test but not by urinary sediment. So there is a possibility that false positive seen at myoglobinuria and false negative seen at use of ascorbic acid may exist. Next, in female patients there is the possibility that menstrual blood may come to be mixed in urine. Not with standing, the mean age of the cohort greater than 60 may negate this possibility. Thirdly, there is the potential presence of unmeasured confounders which cannot be avoided in any observational study. Lastly, there is a possibility of misspecification of the propensity score model which cannot be asserted by any means, either. Despite these limitations, a propensity score analysis clearly captures its unbounded potential to examine many test conditions such as risk factors and target thresholds. Particularly intervention is not easy to perform such as microscopic hematuria.

**Conclusion**

The propensity score analysis unveiled that mild hematuria in the follow-up can predict the risk of ESRD. Target range of hematuria in the follow-up is less than ± in dipstick test to inhibit the progression of CKD to ESRD. The results may facilitate vigorous surveillance of microscopic hematuria which has been prone to be ignored in CKD practice. The mechanism whereby mild hematuria renders kidney injury in the setting of CKD should be elucidated in animal and human studies.

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Hematuria is a risk factor towards end-stage renal disease - A propensity score analysis

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