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Serum klotho is inversely associated with metabolic syndrome in chronic kidney disease: results from the KNOW-CKD study

Hyo Jin Kim¹, Joongyub Lee^{2,3}, Dong-Wan Chae⁴, Kyu-Beck Lee⁵, Su Ah Sung⁶, Tae-Hyun Yoo⁷, Seung Hyeok Han⁷, Curie Ahn⁸ and Kook-Hwan Oh^{8*} 

Abstract

Background: Metabolic syndrome (MS) is prevalent in chronic kidney disease (CKD). Klotho, a protein linked to aging, is closely associated with CKD. Each component of MS and klotho has an association. However, little is known about the association between klotho and MS per se. We investigated the association between serum klotho levels and MS using baseline cross-sectional data obtained from a large Korean CKD cohort.

Methods: Of the 2238 subjects recruited in the KoreaN Cohort Study for Outcome in Patients With Chronic Kidney Disease (KNOW-CKD) between 2011 and 2016, 484 patients with missing data on serum klotho and extreme klotho values (values lower than the detectable range or > 6000 pg/mL) or with autosomal dominant polycystic kidney disease patients were excluded. The data of the remaining 1754 subjects were included in the present study. MS was defined using the revised National Cholesterol Education Program Adult Treatment Panel (NCEP-ATP) III criteria. Serum klotho levels were measured using an enzyme-linked immunosorbent assay.

Results: Mean patient age was 54.9 ± 12.1 years and 1110 (63.3%) were male. The prevalence of MS among all study subjects was 63.7% ($n = 1118$). The median serum klotho level was 527 pg/mL (interquartile range [IQR]: 418–656 pg/mL). Serum klotho level was significantly lower in MS patients than patients without MS (Median [IQR]; 521 pg/mL [413, 651] vs. 541 pg/mL [427, 676], respectively; $P = 0.012$). After adjusting for age, sex, estimated glomerular filtration rate, and overt proteinuria, serum klotho was independently associated with MS (adjusted odds ratio [OR], 0.44; 95% confidence interval, 0.23–0.82; $P = 0.010$). Furthermore, the adjusted OR for MS was found to be significantly increased at serum klotho levels of < 518 pg/mL (receiver operating characteristic curve cut-off value).

Conclusions: Serum klotho was inversely associated with the presence of MS in patients with CKD.

Trial registration: This trial was registered on ClinicalTrials.gov on 26 June 2012 (<https://clinicaltrials.gov/NCT01630486>).

Keywords: Chronic kidney disease, Klotho, Metabolic syndrome

Background

Metabolic syndrome (MS) is a cluster of risk factors for diabetes and cardiovascular (CV) disease and is characterized by hypertension, hyperglycemia, hypertriglyceridemia, decreased high-density lipoprotein (HDL) cholesterol, and abdominal obesity [1, 2]. Furthermore, MS is common in chronic kidney

disease (CKD) patients [3]. Previous studies showed that MS was independently associated with an increased risk for CKD and vice versa [4, 5].

Klotho, a protein linked to aging, is closely associated with CKD. In a previous study, klotho knock-out mice exhibited similarities with CKD patients, such as, hyperphosphatemia, ectopic soft tissue calcification, and arteriosclerosis [6], which suggested CKD might result from a state of klotho deficiency. Thus, in addition to serving as a biomarker for CKD, klotho deficiency is also viewed as a pathogenetic indicator of renal and extra-renal complications in CKD [7].

* Correspondence: khoh@snu.ac.kr

⁸Department of Internal Medicine, Seoul National University Hospital, Seoul National University College of Medicine, 101 Daehak-Ro, Jongno-Gu, Seoul 03080, Korea

Full list of author information is available at the end of the article



Recent reports indicated *klotho* is associated with each component of MS. Arking et al. [8] showed that a functional variant of the *Klotho* gene was associated with increased systolic blood pressure and decreased HDL-cholesterol levels in 525 Jewish subjects. In experimental studies, *klotho* has been shown to ameliorate vascular endothelial dysfunction, to increase nitric oxide production, and to reduce elevated blood pressure in an animal models of MS [9, 10]. *Klotho* polymorphism was significantly associated with glucose metabolism in apparently healthy Korean females [11]. In addition, genetic variants of *Klotho* have been associated with insulin resistance and hypertriglyceridemia [12]. Accordingly, it was suggested in all of these studies that *klotho* be considered a candidate molecule in MS. However, little is known about the association between *klotho* and MS per se in CKD patients. In the present study, we investigated the metabolic profile characteristics of Korean CKD patients and evaluated the association between serum *klotho* and MS using baseline cross-sectional data obtained from a large Korean CKD cohort.

Methods

Study population

We conducted a cross-sectional analysis of clinical profiles obtained at enrollment for the KoreaN Cohort Study for Outcome in Patients With Chronic Kidney Disease (KNOW-CKD) study. KNOW-CKD is a nationwide prospective cohort study, including predialysis subjects with CKD from stage 1 to 5, aged between 20 and 75 years and recruited from nine clinical centers of major university-affiliated Korean hospitals. Details of the rationale and study design of the KNOW-CKD have been described elsewhere [13]. Of 2238 CKD patients recruited between June 2011 and January 2016, 484 were excluded for the following reasons: 125 for missing serum *klotho* data; 9 for a serum *klotho* level lower than the detectable range; 3 for an extremely higher *klotho* level (> 6000 pg/mL); and 347 for autosomal dominant polycystic kidney disease patients because large cystic masses in their kidneys and liver can cause an overestimation of abdominal obesity. Extreme *klotho* values were not influence value by Cook's distance analysis. Finally, 1754 CKD patients were included in the cross-sectional analysis. *Klotho* expression is known to decline in cases of acute kidney injury, regardless of whether it is caused by ischemia, infection, a toxin, or ureteral obstruction [14–16]. However, in the present study, we did not enroll patients with acute illness at the time of initial enrollment. Therefore, it is unlikely that initial *klotho* levels were influenced by acute illness. The study protocol was approved by the ethical committees of participating clinical centers, that is, by the Institutional Review Boards of Seoul National University Hospital (1104–089-359),

Seoul National University Bundang Hospital (B-1106/129–008), Yonsei University Severance Hospital (4–2011-0163), Kangbuk Samsung Medical Center (2011–01-076), Seoul St. Mary's Hospital (KC11OIMI0441), Gil Hospital (GIRBA2553), Eulji General Hospital (201105–01), Chonnam National University Hospital (CNUH-2011-092), and Pusan Paik Hospital (11–091) in 2011. All study subjects provided written informed consent. This study was performed in accordance with the principles of the Declaration of Helsinki.

Clinical data collection and laboratory analyses

Laboratory values and demographic characteristics at enrollment were extracted from an electronic case-reporting form (<http://www.phactax.org>) developed by the assistance of the Division of Data Management at Seoul National University Medical Research Collaborating Center. Waist circumferences were measured midway between the lower part of the lowest rib and the upper iliac crest using a standardized protocol at clinics, as recommended by the World Health Organization (WHO) [17]. Brachial systolic blood pressure (SBP) and diastolic blood pressure (DBP) were obtained by averaging two measurements (separated by a minimum rest of 5 min in the sitting position) obtained using a calibrated oscillometric device (BP-203RV III; Omron Co., Kyoto, Japan). Serum creatinine levels were measured by an isotope dilution mass spectrometry (IDMS)-traceable method [18] at a central laboratory. Estimated glomerular filtration rate (eGFR) was calculated using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) creatinine equation [19]. Fasting blood glucose and triglyceride levels were measured in the hospital laboratories of participating center according to a standardized protocol. Overt proteinuria was defined as a 24-h urine protein result of > 500 mg/day. Serum α -*klotho* level was measured using an enzyme linked immunosorbent assay (ELISA) kit (Immuno-Biological Laboratories Co., Gunma, Japan) according to the manufacturer's instructions [20]. The intra- and inter-assay coefficients of variation of this kit were 2.7–3.5% (*klotho* levels 186.64–2968.78 pg/mL) and 2.9–11.4% (*klotho* levels 165.47–2903.01 pg/mL), respectively. The data for intra- and inter-assay coefficients of variations were validated in our central laboratory by measurements of a serum control in 20 repeats on each ELISA plate. The intra- and inter-assay coefficients of variations were 0.57–1.78% and 3.01–6.12% (*klotho* levels 120.30–4468.50 pg/mL), respectively. The standard curve was shown to be linear up to 4000 pg/mL. C-terminal FGF23 was measured using a second generation human FGF23 ELISA kit (Immutopics, San Clemente, California, USA) according to the manufacturer's instructions.

Table 1 The clinical characteristics of study subjects at enrollment with respect to the presence of metabolic syndrome

	Total (N = 1754)	Subjects without MS (n = 636)	Subjects with MS (n = 1118)	P-value
Age (years)	54.9 ± 12.1	52.3 ± 12.8	56.3 ± 11.4	< 0.001
Sex, male, n (%)	1110 (63.3)	387 (60.8)	723 (64.7)	0.111
SBP (mmHg)	128.5 ± 17.0	124.9 ± 16.4	130.6 ± 17.0	< 0.001
DBP (mmHg)	76.0 ± 11.2	75.5 ± 10.9	76.5 ± 11.4	0.059
BMI (kg/m ²)	24.7 ± 3.4	22.9 ± 2.9	25.8 ± 3.2	< 0.001
Waist circumference (cm)	88.1 ± 9.8	81.4 ± 8.4	91.7 ± 8.5	< 0.001
DM, n (%)	700 (39.9)	110 (17.3)	590 (52.8)	< 0.001
Hypertension, n (%)	1719 (98.0)	606 (95.3)	1113 (99.6)	< 0.001
Preexisting CV disease, n (%)	298 (17.0)	72 (11.3)	226 (20.2)	< 0.001
CAD, n (%)	110 (6.3)	21 (3.3)	89 (8.0)	< 0.001
PVD, n (%)	76 (4.3)	22 (3.5)	54 (4.8)	0.175
Cerebrovascular disease, n (%)	110 (6.3)	24 (3.8)	86 (7.7)	0.001
HF, n (%)	28 (1.6)	9 (1.4)	19 (1.7)	0.648
Arrhythmia, n (%)	52 (3.0)	19 (3.0)	33 (3.0)	0.966
Cause of CKD, n (%)				< 0.001
GN, n (%)	746 (42.5)	387 (60.8)	359 (32.1)	
Diabetic nephropathy, n (%)	491 (28.0)	72 (11.3)	419 (37.5)	
Hypertension, n (%)	389 (22.2)	142 (22.3)	247 (22.1)	
Others, n (%)	128 (7.3)	35 (5.5)	93 (8.3)	
Laboratory findings				
eGFR (mL/min/1.73m ²)	49.1 ± 28.6	53.1 ± 29.6	46.8 ± 27.9	< 0.001
Hemoglobin (g/dL)	12.7 ± 2.1	12.8 ± 2.0	12.7 ± 2.1	0.798
Albumin (g/dL)	4.1 ± 0.4	4.1 ± 0.4	4.1 ± 0.5	0.357
Uric acid (mg/dL)	7.2 ± 1.9	7.1 ± 1.9	7.3 ± 1.9	0.025
Creatinine (mg/dL)	1.9 ± 1.2	1.8 ± 1.1	2.0 ± 1.2	0.002
hsCRP, median, (IQR) (mg/dL)	0.07 (0.03, 0.17)	0.04 (0.02, 0.11)	0.08 (0.03, 0.21)	0.005
Fasting blood sugar (mg/dL)	113.9 ± 42.7	100.4 ± 28.5	121.5 ± 47.2	< 0.001
Total cholesterol (mg/dL)	173.4 ± 40.4	173.5 ± 36.2	173.3 ± 42.5	0.917
Triglyceride (mg/dL)	164.1 ± 103.1	106.3 ± 52.0	196.0 ± 110.2	< 0.001
LDL cholesterol (mg/dL)	95.0 ± 32.7	96.5 ± 30.1	95.7 ± 34.1	0.609
HDL cholesterol (mg/dL)	48.2 ± 15.5	56.5 ± 15.0	43.7 ± 13.8	< 0.001
Overt proteinuria ^a , n (%)	942 (61.0)	319 (58.3)	623 (62.5)	0.108
Medication, n (%)				
ACEi or ARB, n (%)	1521 (86.9)	526 (83.0)	995 (89.1)	< 0.001
Diuretics, n (%)	443 (25.3)	99 (15.6)	344 (25.3)	< 0.001
Statin, n (%)	998 (56.9)	323 (50.8)	675 (60.4)	< 0.001
Ca-based P binders, n (%)	161 (9.2)	67 (10.5)	94 (8.4)	0.179

^a24-h urine protein > 500 mg/day

MS metabolic syndrome, SBP systolic blood pressure, DBP diastolic blood pressure, BMI body mass index, DM diabetes mellitus, CV cardiovascular, CAD coronary artery disease, HF heart failure, PVD peripheral vascular disease, CKD chronic kidney disease, GN glomerulonephritis, eGFR estimated glomerular filtration rate by CKD-EPI creatinine equation, hsCRP high sensitivity C-reactive protein, IQR interquartile range, LDL low-density lipoprotein, HDL high-density lipoprotein, ACEi angiotensin converting enzyme inhibitor, ARB angiotensin II receptor blocker, Ca calcium, P phosphorus

The intra- and inter-assay coefficients of variation as informed by the manufacturer were 1.4–2.4% (FGF23 levels 33.7–302 RU/mL) and 2.4–4.7% (FGF23 levels 33.6–293 RU/mL), respectively.

Definition of metabolic syndrome

MS was defined using the modified National Cholesterol Education Program Adult Treatment Panel (NCEP-ATP) III criteria according to American Heart Association/National Heart, Lung, and Blood Institute Scientific Statement, which adopted the Asian-Pacific waist circumference threshold by WHO. Subjects with three or more of the following 5 metabolic components were defined as having MS [2, 21]: 1) history of hypertension, elevated blood pressure (systolic blood pressure \geq 130 mmHg, or diastolic blood pressure \geq 85 mmHg), or taking anti-hypertensive medication; 2) history of diabetes mellitus, elevated fasting plasma glucose (\geq 100 mg/dL), or the use of anti-diabetes medication; 3) elevated triglycerides (\geq 150 mg/dL), or on drug treatment for elevated triglycerides; 4) reduced HDL cholesterol ($<$ 40 mg/dL for men and $<$ 50 mg/dL for women), or on drug treatment for reduced HDL cholesterol; and 5) elevated waist circumference (\geq 90 cm for men and \geq 80 cm for women) according to the Asia-Pacific criteria [21, 22].

Statistical analysis

Categorical variables were evaluated using the Chi-square test or Fisher’s exact test and presented as frequencies and percentages. Continuous variables were analyzed with independent sample *t*-tests or the Mann-Whitney U test. Results are presented as the mean \pm standard deviation (SD) for normally distributed variables and the median (interquartile range [IQR]) for variables with skewed distributions. Log transformation was used to normalize klotho variability. We performed binominal logistic regression model analysis with adjustments (the Enter method), including variables that were significant in a univariate analysis or other clinically relevant variables to investigate the independent risk factors related to MS. To explore the association between MS and eGFR, or between MS and serum klotho considering the possible non-linearity, we adopted 3-knots restricted cubic splines with 95% confidence interval (CI) which described odds ratios (ORs) of MS according to the eGFR or klotho level. Knots were chosen using SAS LGTPHCURV9 Macro by Li et al. at the eGFR of 13.4, 42.7, and 106 ml/min/1.73m²; 257, 526, and 946 pg/mL for klotho, respectively. An eGFR of 60 ml/min/1.73m², as a reference value of decreased renal function, and a serum klotho of 518 pg/mL (ROC [receiver operating characteristic] curve cut-off value) were taken as the reference point (OR = 1.00). The final model was adjusted for confounders. In subgroup analyses using an

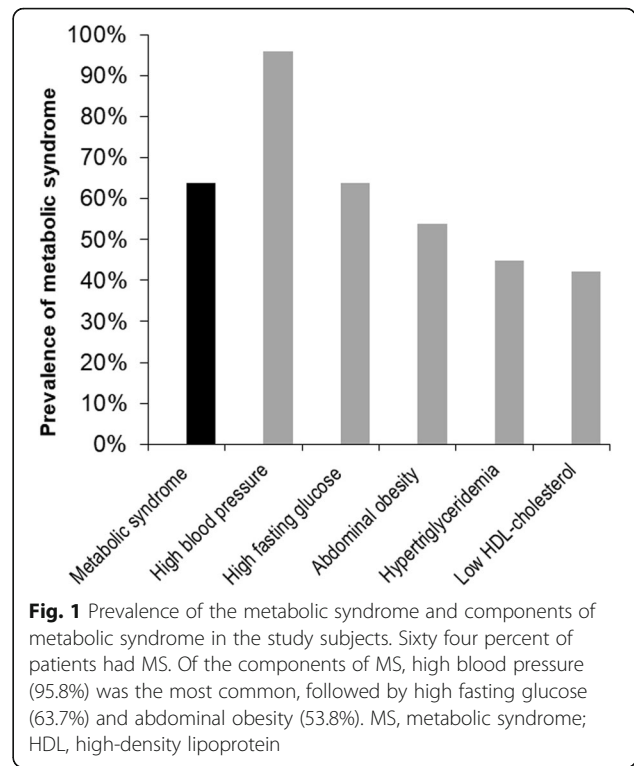


Fig. 1 Prevalence of the metabolic syndrome and components of metabolic syndrome in the study subjects. Sixty four percent of patients had MS. Of the components of MS, high blood pressure (95.8%) was the most common, followed by high fasting glucose (63.7%) and abdominal obesity (53.8%). MS, metabolic syndrome; HDL, high-density lipoprotein

adjusted binominal logistic regression model (adjusted for age, sex, eGFR, overt proteinuria), we categorized patients according to age (aged $<$ 50 vs. \geq 50 years), sex, and CKD stages. *P*-values of $<$ 0.05 were considered statistically significant. The SPSS statistical software (SPSS version 18.0, Chicago, IL, USA) and SAS

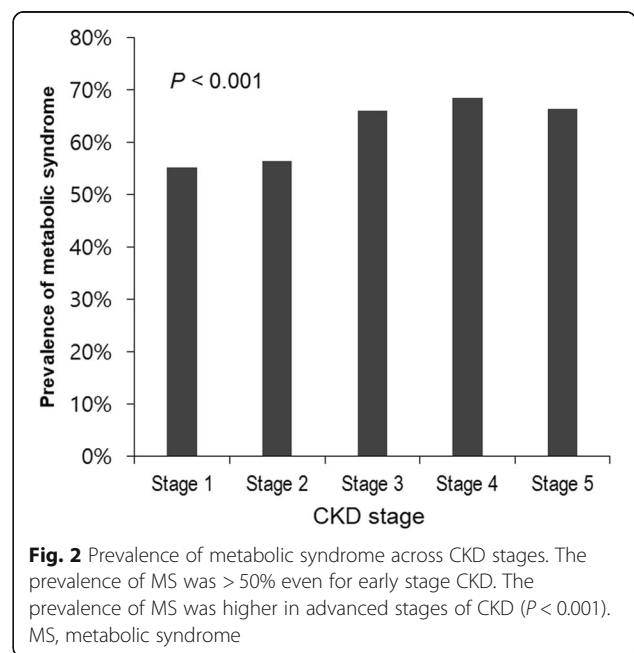


Fig. 2 Prevalence of metabolic syndrome across CKD stages. The prevalence of MS was $>$ 50% even for early stage CKD. The prevalence of MS was higher in advanced stages of CKD ($P <$ 0.001). MS, metabolic syndrome

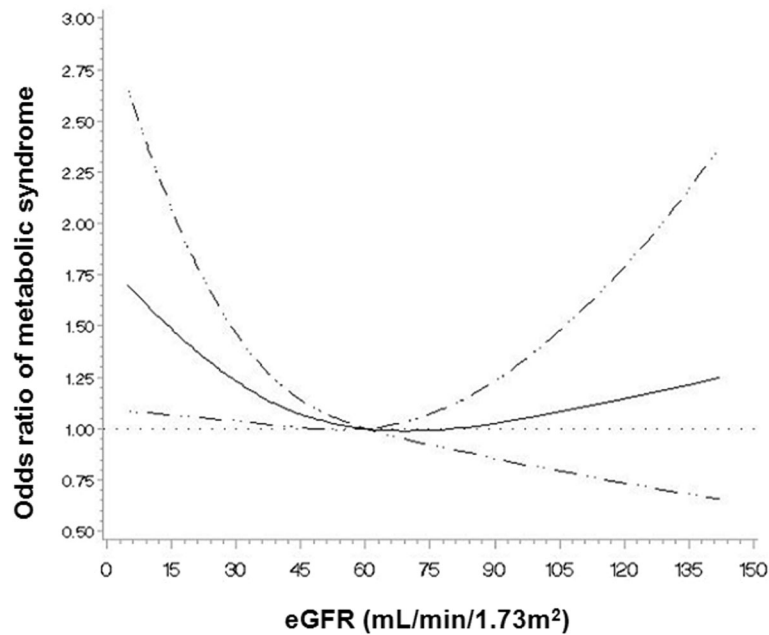


Fig. 3 Multivariable-adjusted odds ratios of metabolic syndrome according to levels of estimated glomerular filtration rate. An eGFR of 60 ml/min/1.73m², as a reference value of decreased renal function, was taken as the reference point (OR = 1.00). The adjusted OR of MS was significantly increased at eGFR levels of < 60 ml/min/1.73m². The model was adjusted for age, sex, eGFR, and overt proteinuria. The solid line represents the multivariable-adjusted ORs of MS according to levels of eGFR. The dashed lines indicate 95% confidence intervals. eGFR, estimated glomerular filtration rate; OR, odds ratio; MS, metabolic syndrome

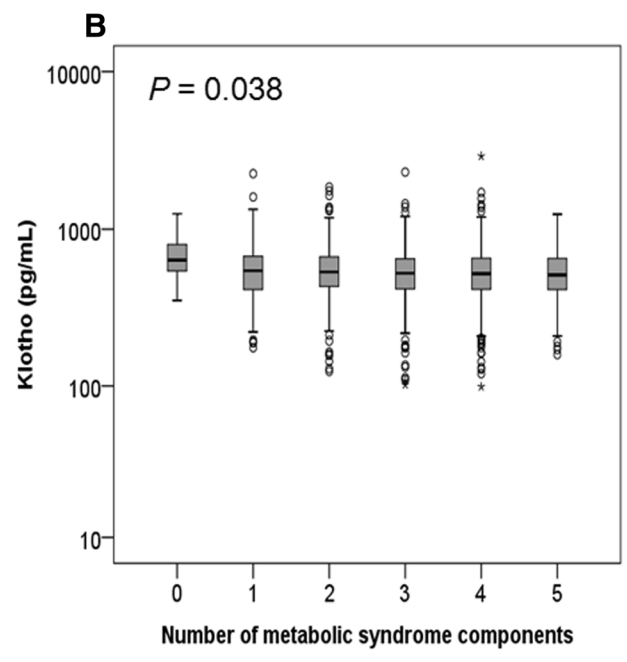
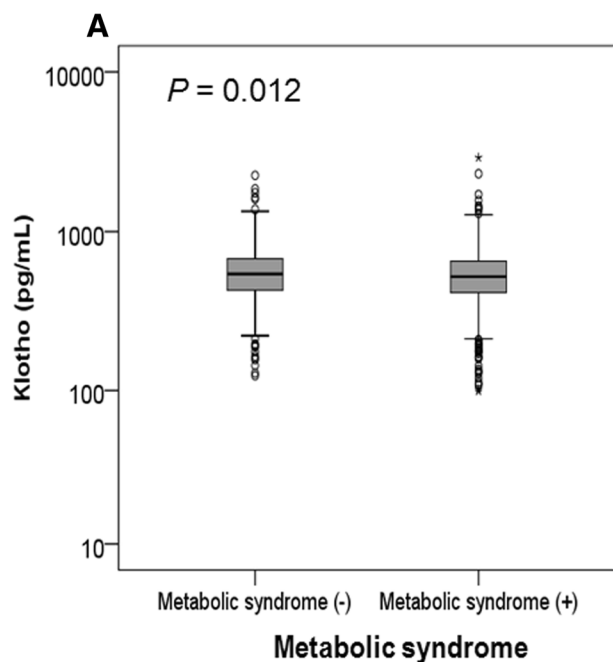


Fig. 4 Serum klotho level according to metabolic syndrome and numbers of metabolic syndrome components. **a** Serum klotho was significantly lower in MS patients compared with patients without MS (median [interquartile range]; 521 pg/mL [413, 651] vs. 541 pg/mL [427, 676], respectively; $P = 0.012$). **b** Klotho levels tended to decrease as numbers of MS components increased ($P = 0.038$). MS, metabolic syndrome

(Version 9.4, SAS Institute Inc., Cary, NC, USA) were used for all descriptive and outcome analyses.

Results

Baseline clinical characteristics of subjects according to the presence of metabolic syndrome

The clinical characteristics of patients at enrollment are shown in Table 1. Mean age of the 1754 study subjects was 54.9 ± 12.1 years, and 1110 (63.3%) were male. Median serum klotho level was 527 pg/mL (IQR: 418–656 pg/mL). Among them, 1118 (63.7%) patients exhibited MS. The frequencies of the five MS component in the study subjects are shown in Fig. 1. High blood pressure (95.8%) was the most prevalent component, followed by high fasting glucose (63.7%) and abdominal obesity (53.8%). Figure 2 presents the prevalence of MS across CKD stages. The prevalence of MS was > 50% even for early stage CKD. The prevalence of MS was higher in advanced stages of CKD ($P < 0.001$, P for linear trend < 0.001). As show in Fig. 3, the adjusted OR (adjusted for age, sex, eGFR, and overt proteinuria) of MS was significantly increased at eGFR levels of < 60 ml/min/1.73m². Subjects were divided into two groups according to the presence of MS. Patients with MS were older ($P < 0.001$, Table 1) and had higher blood pressure ($P < 0.001$), body mass index ($P < 0.001$), and waist circumference ($P < 0.001$). At enrollment, diabetes mellitus (DM) ($P < 0.001$), hypertension ($P < 0.001$), and preexisting CV disease ($P < 0.001$) were more prevalent among subjects with MS. Mean eGFR ($P < 0.001$) was lower and uric acid ($P = 0.025$) level was higher in MS patients. Serum klotho was significantly lower in MS patients compared with patients without MS (Median [IQR]; 521 pg/mL [413, 651] vs. 541 pg/mL [427, 676], respectively; $P = 0.012$; Fig. 4a). In addition, klotho levels tended to decrease as numbers of MS components increased (Fig. 4b; $P = 0.038$).

Clinical characteristics of subjects stratified by serum klotho level

The clinical characteristics of patients dichotomized by serum klotho level (lower than median vs. equal to or higher than median) are shown in Table 2. Age, sex, blood pressure, and underlying comorbidities were similar in these two groups. However, patients with a lower klotho level had a higher mean uric acid level ($P < 0.001$) and a higher C-reactive protein (CRP) level ($P = 0.005$). Mean eGFR ($P < 0.001$) and hemoglobin ($P < 0.001$) were lower in the low serum klotho level group. Serum klotho levels across CKD stages are shown in Fig. 5. Advanced CKD stages were associated with lower serum klotho levels ($P < 0.001$).

Table 2 Baseline characteristics of subjects stratified by serum klotho level

	Klotho groups		P-value
	Lower than median (n = 877) (99–526 pg/mL)	Equal to or higher than median (n = 877) (527–2909 pg/mL)	
Age (years)	55.3 ± 11.9	54.5 ± 12.3	0.152
Sex, male, n (%)	556 (63.4)	554 (63.2)	0.921
SBP (mmHg)	128.9 ± 16.7	128.2 ± 17.2	0.387
DBP (mmHg)	76.3 ± 11.5	76.0 ± 10.9	0.568
BMI (kg/m ²)	24.9 ± 3.4	24.6 ± 3.4	0.060
Waist circumference (cm)	88.4 ± 9.8	87.8 ± 9.8	0.203
DM, n (%)	348 (39.7)	352(40.1)	0.845
Hypertension, n (%)	866 (98.7)	853 (97.3)	0.026
Preexisting CV disease, n (%)	144 (16.4)	154 (17.6)	0.525
CAD, n (%)	46 (5.2)	64 (7.3)	0.076
PVD, n (%)	40 (4.6)	36 (4.1)	0.639
Cerebrovascular disease, n (%)	50 (5.7)	60 (6.8)	0.325
HF, n (%)	11 (1.3)	17 (1.9)	0.253
Arrhythmia, n (%)	23 (2.6)	29 (3.3)	0.398
Cause of CKD, n (%)			0.012
GN, n (%)	358 (40.8)	388 (44.2)	
Diabetic nephropathy, n (%)	253 (28.8)	238 (27.1)	
Hypertension, n (%)	214 (24.4)	175 (20.0)	
Others, n (%)	52 (5.9)	73 (8.7)	
Laboratory findings			
eGFR (mL/min/1.73m ²)	44.9 ± 26.0	53.6 ± 30.6	< 0.001
Hemoglobin (g/dL)	12.5 ± 2.0	13.0 ± 2.1	< 0.001
Albumin (g/dL)	4.1 ± 0.4	4.1 ± 0.5	0.953
Uric acid (mg/dL)	7.5 ± 1.9	7.0 ± 1.9	< 0.001
Creatinine (mg/dL)	2.0 ± 1.2	1.8 ± 1.1	< 0.001
hsCRP, median, (IQR) (mg/dL)	0.07 (0.03, 0.18)	0.06 (0.02, 0.15)	0.005
Fasting blood sugar (mg/dL)	110.9 ± 34.9	117.0 ± 49.1	0.003
Total cholesterol (mg/dL)	172.8 ± 39.7	174.0 ± 41.0	0.531
Triglyceride (mg/dL)	168.6 ± 107.0	159.7 ± 98.9	0.072
LDL cholesterol (mg/dL)	95.0 ± 32.3	97.0 ± 33.1	0.205
HDL cholesterol (mg/dL)	47.9 ± 15.9	48.6 ± 15.1	0.356
Overt proteinuria ^a , n (%)	475 (61.3)	467 (60.7)	0.821
Klotho, median, (IQR) (pg/mL)	418 (337, 475)	656 (583, 774)	< 0.001
Medication, n (%)			
ACEi or ARB, n (%)	773 (88.2)	748 (85.5)	0.088
Diuretics, n (%)	342 (39.0)	295 (33.6)	0.058
Statin, n (%)	514 (58.6)	484 (55.2)	0.309
Ca-based P binders, n (%)	88 (10.0)	73 (8.3)	0.396
Metabolic syndrome, n (%)	574 (65.5)	544 (62.0)	0.136

^a24-hour urine protein > 500 mg/day
MS metabolic syndrome, SBP systolic blood pressure, DBP diastolic blood pressure, BMI body mass index, DM diabetes mellitus, CV cardiovascular, CAD coronary artery disease, HF heart failure, PVD peripheral vascular disease, CKD chronic kidney disease, GN glomerulonephritis, eGFR estimated glomerular filtration rate by CKD-EPI creatinine equation, hsCRP high sensitivity C-reactive protein, IQR interquartile range, LDL low-density lipoprotein, HDL high-density lipoprotein, ACEi angiotensin converting enzyme inhibitor, ARB angiotensin II receptor blocker, Ca calcium, P phosphorus

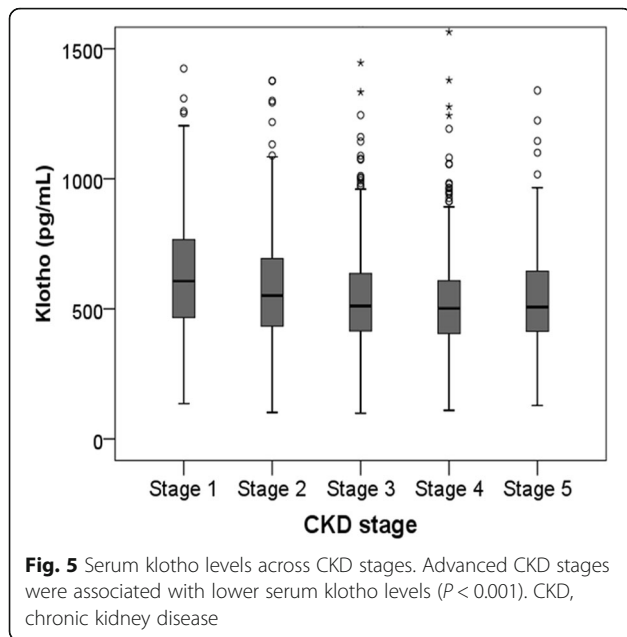


Fig. 5 Serum klotho levels across CKD stages. Advanced CKD stages were associated with lower serum klotho levels ($P < 0.001$). CKD, chronic kidney disease

Associations of metabolic syndrome and each metabolic syndrome component with serum klotho

In the univariate analysis, serum log klotho (OR, 0.40; 95% confidence interval [CI], 0.23–0.72; $P = 0.002$) and eGFR (OR, 0.99; 95% CI, 0.98–0.99; $P < 0.001$) were inversely associated with the presence of MS. On the other hand, age (OR, 1.03; 95% CI, 1.02–1.04; $P < 0.001$) was positively associated with MS. Binominal logistic regression analysis, adjusted for age, sex, eGFR, and overt proteinuria, showed that log klotho was independently associated with the presence of MS (adjusted OR, 0.44; 95% CI, 0.23–0.82; $P = 0.010$; model B in Table 3). As shown in Fig. 6, the adjusted OR of MS was significantly increased at serum klotho levels of < 518 pg/mL (the ROC cut-off value). Furthermore, adjusted for age, sex, eGFR, overt proteinuria, and drug information (ACEi or ARB, diuretics, statin), log klotho was still independently

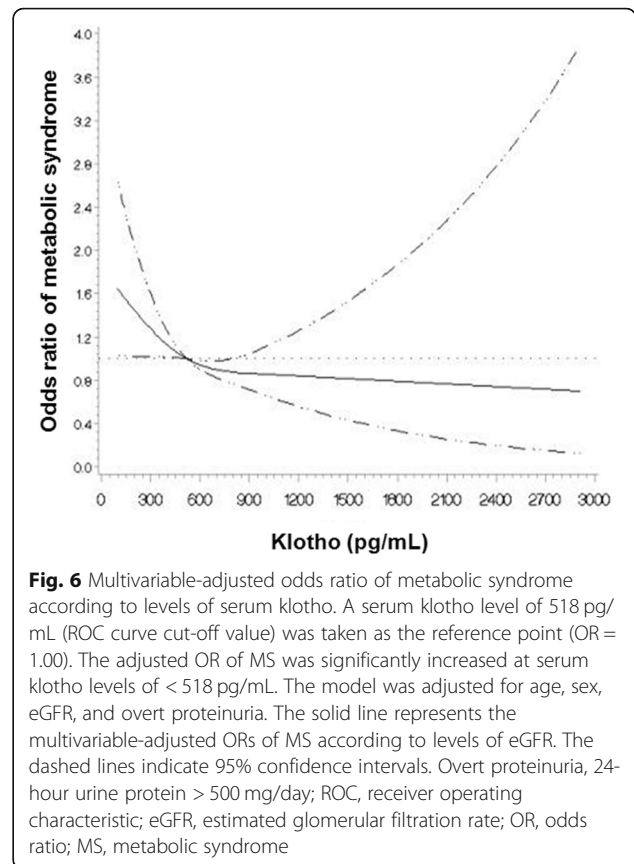


Fig. 6 Multivariable-adjusted odds ratio of metabolic syndrome according to levels of serum klotho. A serum klotho level of 518 pg/mL (ROC curve cut-off value) was taken as the reference point (OR = 1.00). The adjusted OR of MS was significantly increased at serum klotho levels of < 518 pg/mL. The model was adjusted for age, sex, eGFR, and overt proteinuria. The solid line represents the multivariable-adjusted ORs of MS according to levels of eGFR. The dashed lines indicate 95% confidence intervals. Overt proteinuria, 24-hour urine protein > 500 mg/day; ROC, receiver operating characteristic; eGFR, estimated glomerular filtration rate; OR, odds ratio; MS, metabolic syndrome

associated with the presence of MS (adjusted OR, 0.51; 95% CI, 0.27–0.98; $P = 0.045$).

ORs of relations between each MS component and serum klotho are shown in Table 4. Only high blood pressure (adjusted OR, 0.05; 95% CI, 0.01–0.29; $P = 0.001$) and hypertriglyceridemia (adjusted OR, 0.48; 95% CI, 0.27–0.87; $P = 0.016$) were found to be independently associated with log klotho. High fasting glucose, abdominal obesity, and low HDL cholesterol were not significantly associated with serum klotho.

Table 3 Multivariable logistic regression analysis presenting associations between log klotho and metabolic syndrome

Variable	Model A ^a	Model B ^b
	Adjusted OR (95% CI)	Adjusted OR (95% CI)
Log klotho	0.44 (0.25–0.80) ^d	0.44 (0.23–0.82) ^d
Age (per year)	1.03 (1.02–1.04) ^d	1.03 (1.02–1.04) ^d
Sex (male vs. female)	1.15 (0.94–1.41)	1.13 (0.91–1.40)
eGFR (per mL/min/1.73m ²)		1.00 (0.99–1.00)
Overt proteinuria ^c		1.26 (1.01–1.58) ^d

OR Odds ratio, CI confidence interval, eGFR estimated glomerular filtration rate by CKD-EPI creatinine equation

^aModel A adjusted for log klotho, age, and sex

^bModel B adjusted for covariates in model A plus eGFR, and overt proteinuria

^c24-hour urine protein > 500 mg/day

^dsignificant association with MS ($P < 0.05$). $P < 0.05$ was considered significant

Subgroup analysis

We further analyzed the relationship between klotho and MS in several groups. Significant interaction was found in age group (aged < 50 vs. ≥ 50 years), suggesting that the association between klotho and MS was particularly evident in subjects aged < 50 years (adjusted OR, 0.18; 95% CI, 0.06–0.56; $P = 0.003$ vs. adjusted OR, 0.72; 95% CI, 0.33–1.57; $P = 0.412$ in aged ≥ 50 years). Klotho levels tended to decrease as age increased (Pearson’s correlation coefficient $r = -0.083$, $P < 0.001$), but the value was not large. Klotho levels were not significantly different in aged < 50 and ≥ 50 years (Median [IQR]; 535 pg/mL [418, 668] vs. 525 pg/mL [418, 655], respectively; $P = 0.465$). There were no statistical interactions with sex group (interaction $P =$

Table 4 Multivariable logistic regression analysis presenting associations between log klotho and each component of metabolic syndrome

Variable	Adjusted OR (95% CI)				
	High blood pressure	High fasting glucose	Abdominal obesity	Hypertriglyceridemia	Low HDL-cholesterol
Log klotho	0.05 (0.01–0.29)	1.32 (0.70–2.51)	0.68 (0.36–1.28)	0.48 (0.27–0.87)	0.71 (0.38–1.31)
P-value	0.001	0.389	0.235	0.016	0.271

Adjusted for covariates log klotho, age, sex, eGFR, and overt proteinuria

OR Odds ratio, CI confidence interval, HDL high-density lipoprotein, eGFR estimated glomerular filtration rate by CKD-EPI creatinine equation

0.605) and CKD stages (interaction $P = 0.425$) for the association between klotho and MS.

Discussion

MS, a common clinical phenotype demonstrating as a combination of metabolic abnormalities including, hypertension, hyperglycemia, hypertriglyceridemia, decreased high-density lipoprotein cholesterol, and central obesity, is common in CKD [23]. In the present study, 63.7% of the 1754 CKD patients enrolled exhibited MS. The prevalence of MS exceeded 50% even in patients with early stage CKD and its prevalence was higher in advanced CKD stages. In the univariate analysis, serum klotho was significantly lower in patients with MS. After adjusting for various factors, such as, age, sex, eGFR, and overt proteinuria, serum klotho level remained an independent factor associated with MS in CKD.

In previous studies, genetic variants of *Klotho* increased the risk of MS which could be related to their susceptibility to high fasting glucose, high blood pressure, hypertriglyceridemia, and decreased HDL cholesterol [8, 12]. *Klotho* has a role in the regulation of vascular tone through homeostatic interplay between the renin-angiotensin system and nitric oxide [24], which might explain the association between klotho and blood pressure. Furthermore, the *Klotho* has been shown to be a target gene for PPAR γ , a key transcription factor controlling lipid metabolism and insulin sensitivity [25]. Thus, we speculated these factors might play major roles in the association of klotho with glucose and lipid metabolism.

MS is a cluster of risk factors of diabetes and CV disease. Subjects with CKD present higher incidence of CV disease and higher CV mortality than the general population [26, 27]. Novel risk factors such as persistent inflammation, oxidative stress, and endothelial dysfunction can be linked to CV disease in CKD patients [28]. Fundamental manifestations of MS include insulin resistance and adipose tissue expansion, and the latter leads to oxidative stress and chronic inflammation, which aggravate insulin resistance [29–31]. Furthermore, these pro-inflammatory manifestations of MS might play roles in the pathogenesis of endothelial dysfunction and atherosclerosis [32]. *Klotho* is an anti-aging gene that extends life span when overexpressed and accelerates aging when disrupted [6]. Aging which reflects cellular

senescence is closely associated with inflammatory reactions, oxidative damage, and endothelial dysfunction [33–35]. *Klotho* has also been shown to be an anti-inflammatory modulator [36, 37], to be closely associated with oxidative stress [38, 39]. *Klotho* plays a role in the protection against endothelial dysfunction [10]. The kidney is the principal organ responsible for the production of klotho, and CKD is known to be a *klotho* deficient state. Accordingly, the effects of klotho mentioned above suggest an association between klotho and MS beyond individual components of MS.

The present study is the first study, to our knowledge, to report the independent association between serum klotho level and MS per se especially in CKD patients. In previous studies, there were not many human studies, and most of studies were related to the association between genetic variants of *Klotho* and individual components of MS. In addition, few studies have been performed in patients with CKD. Furthermore, it was the strength of our study that we included a large-scale CKD cohort patients and obtained serum klotho level from a large number of our CKD patients. However, the study has several limitations that warrant consideration. First, because it was based on cross-sectional analysis, it could not demonstrate causality between klotho and MS. It could not provide detailed mechanistic links between klotho and MS. We just described the association between klotho and MS, not the causality. Second, serum klotho exhibits circadian variations [40]. They showed that midnight serum klotho level was an approximately 40% reduction with gradual return to near baseline values in the early morning hours. Serum klotho level decreased less than 20% compared with morning klotho level until 6 p.m. In the present study, blood samples were not obtained at any fixed time of the day. However, circadian variation of serum klotho may have been less in our study because blood sampling was done before 6 p.m.

Conclusions

Serum klotho levels were found to be independently and inversely associated with the presence of MS. Further studies are warranted to elucidate the nature of the mechanistic link between klotho and MS in CKD patients.

Abbreviations

ACEi: Angiotensin converting enzyme inhibitor; ARB: Angiotensin II receptor blocker; Ca: Calcium; CI: Confidence interval; CKD: Chronic kidney disease; CKD-EPI: Chronic kidney Disease Epidemiology Collaboration; CRP: C-reactive protein; CV: Cardiovascular; DBP: Diastolic blood pressure; DM: Diabetes mellitus; eGFR: Estimated glomerular filtration rate; ELISA: Enzyme-linked immunosorbent assay; HDL: High-density lipoprotein; IDMS: Isotope dilution mass spectrometry; IQR: Interquartile range; KNOW-CKD: KoreaN Cohort Study for Outcome in Patients With Chronic Kidney Disease; MS: Metabolic syndrome; NCEP-ATP: National Cholesterol Education Program Adult Treatment Panel; OR: Odds ratio; P: Phosphorus; ROC: Receiver operating characteristic; SBP: Systolic blood pressure; SD: Standard deviation; WHO: World Health Organization

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Availability of data and materials

The data set that support the findings is not publicly available. Limited de-identified data sets may be available from the authors upon reasonable request and with permission of the study sponsor.

Authors' contributions

HJK and K-HO conceived and designed the study, analyzed and interpreted the data, wrote the manuscript, contributed to discussions, and reviewed and edited the manuscript. JL made substantial contributions to the analysis of data and drafted the article. D-WC, K-BL, SAS, T-HY, SHH, and CA contributed to the study conception and design, contributed to data interpretation and discussion, and reviewed and edited the manuscript. All authors approved the final manuscript. K-HO was the principal investigator and is responsible for the integrity of the work as a whole. All authors revised the manuscript content and approved the final version.

Ethics approval and consent to participate

The study protocol was approved by the ethical committee of each participating clinical center, including the Institutional Review Boards of Seoul National University Hospital, Severance Hospital, Kangbuk Samsung Medical Center, Seoul St. Mary's Hospital, Gil Hospital, Eulji General Hospital, Chonnam National University Hospital, and Busan Paik Hospital. All participating patients provided written informed consent. The study protocol was in accordance with the principles of the Declaration of Helsinki.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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Author details

¹Department of Internal Medicine, Dongguk University College of Medicine, Gyeongju-si, Gyeongsangbuk-do, Korea. ²School of Medicine, Inha University, Incheon, Korea. ³Department of Prevention and Management, Inha University Hospital, Incheon, Korea. ⁴Department of Internal Medicine, Seoul National University Bundang Hospital, Seongnam-si, Gyeonggi-do, Korea. ⁵Department of Internal Medicine, Kangbuk Samsung Hospital, Sungkyunkwan University School of Medicine, Seoul, Korea. ⁶Department of Internal Medicine, Nowon Eulji Medical Center, Eulji University, Seoul, Korea. ⁷Department of Internal Medicine, Yonsei University College of Medicine, Seoul, Korea. ⁸Department of Internal Medicine, Seoul National University Hospital, Seoul National University College of Medicine, 101 Daehak-Ro, Jongno-Gu, Seoul 03080, Korea.

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References

- Laaksonen DE, Lakka HM, Niskanen LK, Kaplan GA, Salonen JT, Lakka TA. Metabolic syndrome and development of diabetes mellitus: application and validation of recently suggested definitions of the metabolic syndrome in a prospective cohort study. *Am J Epidemiol*. 2002;156:1070–7.
- Grundy SM, Cleeman JI, Daniels SR, Donato KA, Eckel RH, Franklin BA, et al. Diagnosis and management of the metabolic syndrome: an American Heart Association/National Heart, Lung, and Blood Institute Scientific Statement. *Circulation*. 2005;112:2735–52.
- Thomas G, Sehgal AR, Kashyap SR, Srinivas TR, Kirwan JP, Navaneethan SD. Metabolic syndrome and kidney disease: a systematic review and meta-analysis. *Clin J Am Soc Nephrol*. 2011;6:2364–73.
- Kurella M, Lo JC, Chertow GM. Metabolic syndrome and the risk for chronic kidney disease among nondiabetic adults. *J Am Soc Nephrol*. 2005;16:2134–40.
- Nashar K, Egan BM. Relationship between chronic kidney disease and metabolic syndrome: current perspectives. *Diabetes Metab Syndr Obes*. 2014;7:421–35.
- Kuro-o M, Matsumura Y, Aizawa H, Kawaguchi H, Suga T, Utsugi T, et al. Mutation of the mouse *klotho* gene leads to a syndrome resembling ageing. *Nature*. 1997;390:45–51.
- Hu MC, Kuro-o M, Moe OW. *Klotho* and chronic kidney disease. *Contrib Nephrol*. 2013;180:47–63.
- Arking DE, Atzmon G, Arking A, Barzilai N, Dietz HC. Association between a functional variant of the *KLOTHO* gene and high-density lipoprotein cholesterol, blood pressure, stroke, and longevity. *Circ Res*. 2005;96:412–8.
- Yamagishi T, Saito Y, Nakamura T, Takeda S, Kanai H, Sumino H, et al. Troglitazone improves endothelial function and augments renal *klotho* mRNA expression in Otsuka Long-Evans Tokushima Fatty (OLETF) rats with multiple atherogenic risk factors. *Hypertens Res*. 2001;24:705–9.
- Saito Y, Nakamura T, Ohyama Y, Suzuki T, Iida A, Shiraki-Iida T, et al. In vivo *klotho* gene delivery protects against endothelial dysfunction in multiple risk factor syndrome. *Biochem Biophys Res Commun*. 2000;276:767–72.
- Rhee EJ, Oh KW, Yun EJ, Jung CH, Lee WY, Kim SW, et al. Relationship between polymorphisms G395A in promoter and C1818T in exon 4 of the *KLOTHO* gene with glucose metabolism and cardiovascular risk factors in Korean women. *J Endocrinol Investig*. 2006;29:613–8.
- Majumdar V, Christopher R. Association of exonic variants of *Klotho* with metabolic syndrome in Asian Indians. *Clin Chim Acta*. 2011;412:1116–21.
- Oh KH, Park SK, Park HC, Chin HJ, Chae DW, Choi KH, et al. KNOW-CKD (KoreaN cohort study for outcome in patients with chronic kidney disease): design and methods. *BMC Nephrol*. 2014;15:80.
- Kuro-o M. *Klotho* in health and disease. *Curr Opin Nephrol Hypertens*. 2012; 21:362–8.
- Hu MC, Moe OW. *Klotho* as a potential biomarker and therapy for acute kidney injury. *Nat Rev Nephrol*. 2012;8:423–9.
- Seo MY, Yang J, Lee JY, Kim K, Kim SC, Chang H, et al. Renal *Klotho* expression in patients with acute kidney injury is associated with the severity of the injury. *Korean J Intern Med*. 2015;30:489–95.
- WHO expert committee. Physical status: the use and interpretation of anthropometry. Report of a WHO Expert Committee. *World Health Organ Tech Rep Ser*. 1995;854:1–452.
- Siekman L. Determination of creatinine in human serum by isotope dilution-mass spectrometry. Definitive methods in clinical chemistry, IV. *J Clin Chem Clin Biochem*. 1985;23:137–44.
- Levey AS, Stevens LA, Schmid CH, Zhang YL, Castro AF 3rd, Feldman HI, et al. A new equation to estimate glomerular filtration rate. *Ann Intern Med*. 2009;150:604–12.
- Yamazaki Y, Imura A, Urakawa I, Shimada T, Murakami J, Aono Y, et al. Establishment of sandwich ELISA for soluble alpha-*Klotho* measurement: age-dependent change of soluble alpha-*Klotho* levels in healthy subjects. *Biochem Biophys Res Commun*. 2010;398:513–8.
- Alberti KG, Eckel RH, Grundy SM, Zimmet PZ, Cleeman JI, Donato KA, et al. Harmonizing the metabolic syndrome: a joint interim statement of the international diabetes federation task force on epidemiology and prevention; national heart, lung, and blood institute; American heart association; world heart federation; international atherosclerosis society; and international association for the study of obesity. *Circulation*. 2009;120:1640–5.

22. Hara K, Matsushita Y, Horikoshi M, Yoshiike N, Yokoyama T, Tanaka H, et al. A proposal for the cutoff point of waist circumference for the diagnosis of metabolic syndrome in the Japanese population. *Diabetes Care*. 2006;29:1123–4.
23. Chen J, Muntner P, Hamm LL, Jones DW, Batuman V, Fonseca V, et al. The metabolic syndrome and chronic kidney disease in U.S. adults. *Ann Intern Med*. 2004;140:167–74.
24. Yang J, Matsukawa N, Rakugi H, Imai M, Kida I, Nagai M, et al. Upregulation of cAMP is a new functional signal pathway of Klotho in endothelial cells. *Biochem Biophys Res Commun*. 2003;301:424–9.
25. Zhang H, Li Y, Fan Y, Wu J, Zhao B, Guan Y, et al. Klotho is a target gene of PPAR-gamma. *Kidney Int*. 2008;74:732–9.
26. Causes of death. United States renal data system. *Am J Kidney Dis*. 1998;32:S81–8.
27. Foley RN, Parfrey PS, Sarnak MJ. Clinical epidemiology of cardiovascular disease in chronic renal disease. *Am J Kidney Dis*. 1998;32:S112–9.
28. Stenvinkel P, Carrero JJ, Axelsson J, Lindholm B, Heimbürger O, Massy Z. Emerging biomarkers for evaluating cardiovascular risk in the chronic kidney disease patient: how do new pieces fit into the uremic puzzle? *Clin J Am Soc Nephrol*. 2008;3:505–21.
29. Zhang X, Lerman LO. The metabolic syndrome and chronic kidney disease. *Transl Res*. 2017;183:14–25.
30. Wahba IM, Mak RH. Obesity and obesity-initiated metabolic syndrome: mechanistic links to chronic kidney disease. *Clin J Am Soc Nephrol*. 2007;2:550–62.
31. Spahis S, Borys JM, Levy E. Metabolic syndrome as a multifaceted risk factor for oxidative stress. *Antioxid Redox Signal*. 2016.
32. Tziomalos K, Athyros VG, Karagiannis A, Mikhailidis DP. Endothelial dysfunction in metabolic syndrome: prevalence, pathogenesis and management. *Nutr Metab Cardiovasc Dis*. 2010;20:140–6.
33. Rangel-Zuniga OA, Corina A, Lucena-Porras B, Cruz-Teno C, Gomez-Delgado F, Jimenez-Lucena R, et al. TNFA gene variants related to the inflammatory status and its association with cellular aging: from the CORDIOPREV study. *Exp Gerontol*. 2016;83:56–62.
34. Hamilton ML, Van Remmen H, Drake JA, Yang H, Guo ZM, Kewitt K, et al. Does oxidative damage to DNA increase with age? *Proc Natl Acad Sci U S A*. 2001;98:10469–74.
35. Salmon AB. Beyond diabetes: does obesity-induced oxidative stress drive the aging process? *Antioxidants (Basel)*. 2016;5:24.
36. Izquierdo MC, Perez-Gomez MV, Sanchez-Nino MD, Sanz AB, Ruiz-Andres O, Poveda J, et al. Klotho, phosphate and inflammation/ageing in chronic kidney disease. *Nephrol Dial Transplant*. 2012;27(Suppl 4):iv6–10.
37. Zhao Y, Banerjee S, Dey N, LeJeune WS, Sarkar PS, Brobey R, et al. Klotho depletion contributes to increased inflammation in kidney of the db/db mouse model of diabetes via RelA (serine)536 phosphorylation. *Diabetes*. 2011;60:1907–16.
38. Oh HJ, Nam BY, Lee MJ, Kim CH, Koo HM, Doh FM, et al. Decreased circulating klotho levels in patients undergoing dialysis and relationship to oxidative stress and inflammation. *Perit Dial Int*. 2015;35:43–51.
39. Yamamoto M, Clark JD, Pastor JV, Gurnani P, Nandi A, Kurosu H, et al. Regulation of oxidative stress by the anti-aging hormone klotho. *J Biol Chem*. 2005;280:38029–34.
40. Carpenter TO, Insogna KL, Zhang JH, Ellis B, Nieman S, Simpson C, et al. Circulating levels of soluble klotho and FGF23 in X-linked hypophosphatemia: circadian variance, effects of treatment, and relationship to parathyroid status. *J Clin Endocrinol Metab*. 2010;95:E352–7.

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