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# Dietary patterns in relation to testosterone levels and severity of impaired kidney function among middle-aged and elderly men in Taiwan: a cross-sectional study

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## Abstract

**Background:** Chronic Kidney Disease (CKD), characterized by an impaired kidney function, is associated with low testosterone levels. This study investigated the association between dietary patterns, testosterone levels, and severity of impaired kidney function among middle-aged and elderly men.

**Methods:** This cross-sectional study used the database from a private health-screening institute in Taiwan between 2008 and 2010. Men aged 40 years old and older ( $n = 21,376$ ) with estimated glomerular filtration rate (eGFR)  $< 90$  mL/min/1.73 m<sup>2</sup> and proteinuria were selected. Among 21,376 men, 256 men had available measurements of testosterone levels. Dietary assessment was conducted using a food frequency questionnaire and three dietary patterns (fried-processed, vege-seafood, and dairy-grain dietary patterns) were identified using principal component analysis.

**Results:** Men in the lower tertiles (T1 and T2) of eGFR had significantly decreased testosterone levels by 0.8 (95% CI:  $-1.40, -0.20$ ) and 0.9 nmol/L (95% CI:  $-1.43, -0.33$ ). Furthermore, serum triglycerides (TG) levels were inversely associated with testosterone levels ( $\beta = -0.51$ , 95% CI:  $-0.77, -0.24$ ). Men in the higher tertile of fried-processed dietary pattern scores were associated with decreased testosterone levels by 0.8 nmol/L (95% CI:  $-1.40, -0.16$ ), reduced testosterone-to-TG (T/TG) ratio by 1.8 units (95% CI:  $-2.99, -0.53$ ), and increased risk of moderate/severe impaired kidney function (eGFR  $< 60$  mL/min/1.73 m<sup>2</sup>) and proteinuria severity by 1.35 (95% CI: 1.15, 1.58) and 1.18 (95% CI: 1.02, 1.37) times respectively. In contrast, the vege-seafood dietary pattern was negatively associated with severity of impaired kidney function and proteinuria after multivariable adjustment, but had no association with testosterone levels and T/TG ratio.

**Conclusions:** The fried-processed dietary pattern is negatively associated with testosterone levels but positively associated with the severity of impaired kidney function. However, the vege-seafood and dairy-grain dietary patterns appear to have beneficial effects.

**Keywords:** Dietary pattern, Testosterone, Kidney function, Principal component analysis

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## Background

Chronic kidney disease (CKD), characterized by an impaired kidney function, is recognized as a global health problem with major adverse outcomes that include cardiovascular disease (CVD) and end-stage renal disease (ESRD). In Taiwan, a large prospective 13-year cohort study showed that the national prevalence of CKD in 2007 was around 12% and less than 4% of the participants were aware of this disease [1]. The prevalence of CKD started to increase at age 40–44 years old (7.4%) and doubled between the ages of 55–59 years (13.1%) and 60–64 years (26.1%) [1]. Moreover, according to the US Renal Data System in 2014, Taiwan had the highest prevalence and incidence rate of ESRD worldwide, especially in the middle-aged and elderly populations [2]. Patients with ESRD spent approximately 7% of the total annual budget from the Taiwan national health insurance program, indicating that CKD not only threatens population health but also becomes a financial burden to the nation [3].

Decreased testosterone levels are the most common gonadal problem among men with impaired kidney function [4]. A cross-sectional study by Yilmaz et al. showed that serum total testosterone levels were decreased by 10% in CKD stage 2 and 42% in CKD stage 5, compared to those in stage 1 and the prevalence of hypogonadism was increased from 75% in men with CKD stage 1 to 92% in CKD stage 5 [5]. Additionally, testosterone deficiency has been associated with reduced kidney function, CKD progression, and all-cause mortality, especially by CVD. A possible explanation is that testosterone has vasodilatory effects on blood vessels, which can prevent the progression of atherosclerosis, ischemia, and endothelial dysfunction [6, 7].

Dietary patterns have been linked to many risks for developing chronic diseases [8]. In previous studies, high intake of protein and Western diet (red and processed meats, animal/saturated fats, and sugar/sweets) were associated with increased risk of CKD, while high intake of vegetables, fruits, and whole grains reduced the risk of decreased kidney function [9–11]. However, previous studies examining the effects of dietary patterns in subjects with impaired kidney function were mostly conducted in Western countries and among women [9–11]. In fact, Taiwanese men have a higher rate of kidney disease than women [1] and there is no published study investigating the correlation of dietary patterns with the severity of impaired kidney function among Taiwanese men. Although a recent study among type 2 diabetes patients in Taiwan showed that diets rich in fish and vegetables were associated with better kidney function [12], further study is still needed to investigate the association between dietary patterns and the onset of impaired kidney function in

a larger sample size. Additionally, to our knowledge, few studies have described the relationship between diets and testosterone levels [13–15]. Notably, a previous study found a significant increase in serum testosterone levels in overweight or obese men on a low-fat diet for 12 weeks and weight maintenance for another 40 weeks [13]. In contrast, two small cross-sectional studies did not find an association between vegetarian diets and testosterone levels in men [14, 15]. In addition, no study has yet examined the relationship between dietary patterns and testosterone levels among men with impaired kidney function. Therefore, this study aims to investigate the relationship between dietary patterns, testosterone, and severity of impaired kidney function among middle-aged and elderly men in Taiwan.

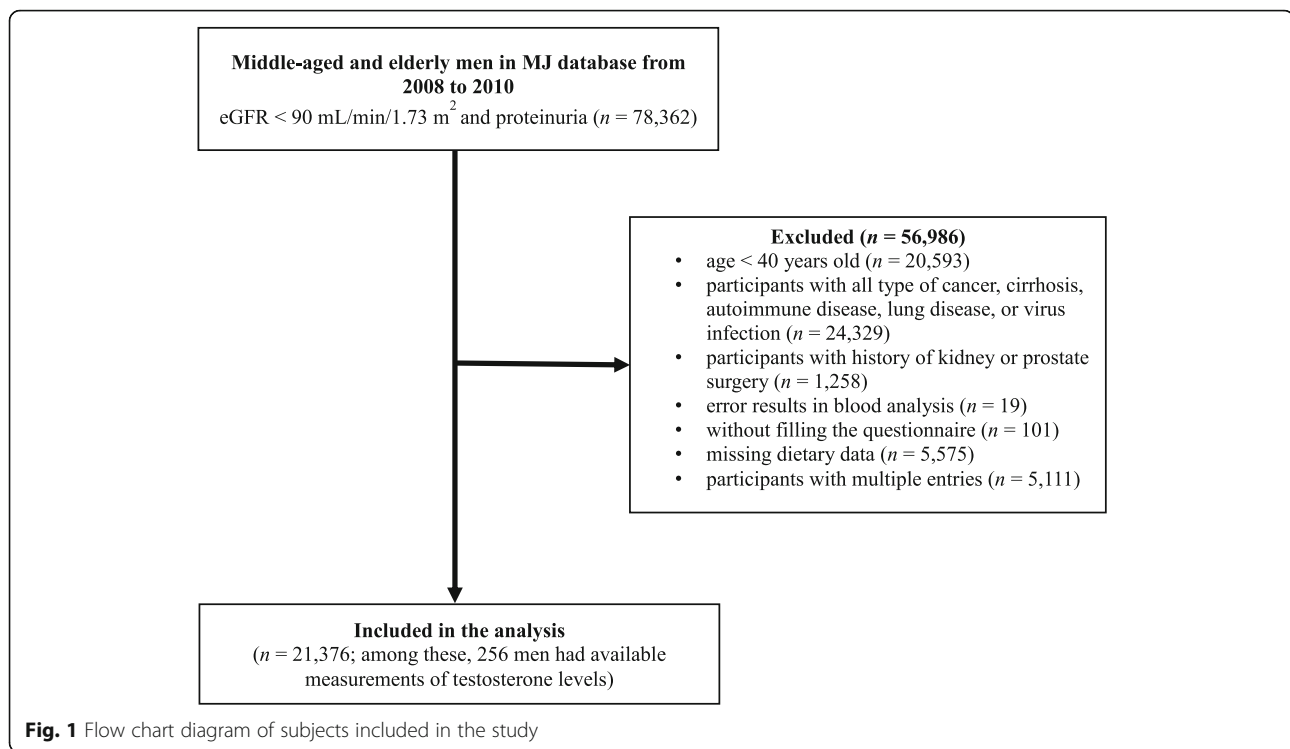
## Methods

### Data source and study subjects

This cross-sectional study was conducted with samples collected from Mei Jau (MJ) Health Management Institution database, Taiwan. The MJ Health Management Institution is a membership-oriented private institute with four health check-up centers (Taipei, Taoyuan, Taichung, and Kaohsiung) in Taiwan, which provides periodic health examination (mostly one examination per year per person) to its members. Subjects completed a self-reported questionnaire to collect information on sociodemographic data, lifestyle, medical history, and dietary habit before blood and urine tests, anthropometric measurement, and a physical examination. All individuals had signed a consent form authorized by the MJ Health Screening Center to process the data generated from medical screening. The data were treated as highly confidential, without personal identification, and for research purposes only. The data consisted of 78,362 middle-aged and elderly men with estimated glomerular filtration rates (eGFR) < 90 mL/min/1.73 m<sup>2</sup> and proteinuria were retrieved from the MJ database between 2008 and 2010. After excluding with some criteria in Fig. 1, a total of 21,376 men were recruited for analysis. The Institutional Ethical Review of Taipei Medical University (TMU-JIRB N201802006) approved this study.

### Clinical and biochemical data

Body weight and height were measured by an auto-anthropometer (Nakamura KN-5000A, Nakamura, Tokyo, Japan). Waist and hip circumferences were measured by a flexible tape. After resting for 5 min, blood pressure was measured twice at 10 min intervals on the right arm in a sitting position by a standardized sphygmomanometer. Hypertension was defined as systolic blood pressure ≥ 140 mmHg or diastolic blood pressure ≥ 90 mmHg, use of an anti-hypertensive drug, or a self-report of hypertension



diagnosis by a physician [16]. Body mass index (BMI) was calculated as weight in kilograms divided by the square of height in meters.

Fasting blood glucose (FBG), triglycerides (TG), total cholesterol (TC), high density lipoprotein-cholesterol (HDL-C), low density lipoprotein-cholesterol (LDL-C), C-reactive protein (CRP), and blood urea nitrogen (BUN) were analyzed (Toshiba C8000 auto-analyzers, Tokyo, Japan) in each subject after fasting for at least 8 h. Diabetes mellitus was defined as FBG  $\geq$  7.0 mmol/L (126 mg/dL), use of a hypoglycemic agent, or a self-report of diabetes diagnosis by a physician [17]. Serum testosterone levels were measured by chemiluminescent microparticle immunoassay (CMIA, Architect System, Abbott, IL, USA). The architect testosterone assay measures total testosterone. However, among 21,376 men, only 256 men had serum testosterone data. Serum creatinine levels were analyzed by uncompensated Jaffe method with alkaline picrate kinetic test. The Roche Miditron M semi-automated computer-assisted urinalysis system (Combur-10 test M dipstick, Basel, Switzerland) was used to measure urinary protein. The result was reported as one plus (equivalent to micro-albuminuria) or more pluses (equivalent to macro-albuminuria). All specimens were analyzed at MJ Health Screening Central Laboratory. The internal and external quality control techniques were performed by the laboratory and the calibrations were supplied by the manufacturer [18]. The laboratory followed Westgard multi-rules for sample quality control and the coefficient of variation ranged from 1 to

3%. Furthermore, we used the equation from Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) to estimate eGFR [19]:

If creatinine  $\leq$  79.56  $\mu$ mol/L (0.9 mg/dL) in male:

$$\text{eGFR (mL/ min/1.73 m}^2\text{)} = 141 \times (\text{creatinine}/0.9)^{-0.411} \times (0.993)^{\text{age}}$$

If creatinine > 79.56  $\mu$ mol/L (0.9 mg/dL) in male:

$$\text{eGFR (mL/ min/1.73 m}^2\text{)} = 141 \times (\text{creatinine}/0.9)^{-1.209} \times (0.993)^{\text{age}}$$

Based on eGFR levels, we categorized men with impaired kidney function into mild ( $n = 19,519$ ) and moderate/severe ( $n = 1,857$ ) groups which were defined as eGFR 60–89 mL/min/1.73 m<sup>2</sup> with positive urinary protein and eGFR < 60 mL/min/1.73 m<sup>2</sup>, respectively [20].

#### Dietary assessment and other covariates

A standardized and validated self-administered semi-quantitative food frequency questionnaire was used to assess dietary habit [21, 22]. The frequency and servings of dietary intake were assessed according to the consumption of twenty-two food groups daily or weekly in the past month before data collection. The frequency intake for each food group was categorized into five responses from the lowest to the highest

using the following units of measurement: a bowl for vegetables and oil added vegetables or salad, a glass for milk and sugar-sweetened beverages, and a serving for the remaining food groups. The definition of the portion size (serving) was described previously [22].

The demographic and lifestyle data such as age, education levels, marital status, family income, smoking, drinking alcohol (no, < 1 time/week; yes,  $\geq$  1–2 times/week), sleeping, physical activity, and medical history (having a history and using drug for CVD, hypertension, and diabetes mellitus) were also collected. The questions regarding physical activity were the frequency and time for exercise during the past 2 weeks. The frequency of exercise had five response options that ranged from none or rarely to 2–3 times per day, while the exercise time had four response options that ranged from less than 30 min to more than 2 h. We defined physical activity as ‘yes’ if the subject engaged in exercise for more than 1 h in a week and ‘no’ if otherwise [22]. A structured questionnaire used to collect demographic and lifestyle data in this study was validated prior to its use and has been documented elsewhere [1, 18]. Moreover, CVD was defined as having a history of CVD or use of a cardiovascular drug.

### Statistical analysis

Statistical analysis was performed by using IBM® SPSS Statistics version 20.0 (International Business Machines Corp., Armonk, NY, USA). Categorical and continuous variables were presented as number (percentage) of the subjects and mean  $\pm$  standard deviation (SD), respectively. A general linear model and a chi-square test for continuous and categorical data respectively, were used for analyzing the characteristics across tertiles of eGFR. Due to the limited number of men who had testosterone levels, we decided to use tertiles of eGFR for the characteristics of the study instead of impaired kidney function categories (only few men who had testosterone levels in the moderate/severe impaired group). Thus, using the severity of kidney function categories as characteristics of the study subjects may not represent the complete results.

Dietary patterns were identified by using principal component analysis (PCA). Three dietary patterns were identified after an orthogonal varimax rotation based on eigenvalue > 1.4 and with the cut-off value of factor loading  $\geq$  0.30. Factor loadings are equivalent to a simple correlation between the food items and the extracted factor or pattern. Higher factor loadings indicate that the food shares more variance with that pattern. Factor scores were calculated for each pattern by summing up the intake of food items or groups weighed by their factor loadings, and thus each

subject received a score for each identified dietary pattern [23]. For further analysis, dietary pattern scores were divided into tertiles. A multivariable linear regression analysis was performed to examine the independent variables associated with testosterone levels and the relationship across tertiles of dietary pattern scores with testosterone levels and kidney function biomarkers. The data were presented as regression coefficient ( $\beta$ ) and 95% confidence interval (95% CI). Meanwhile, a multivariable logistic regression analysis was used to identify the association of dietary pattern scores across tertiles with impaired kidney function and proteinuria severity. The data were presented as odds ratio (OR) with 95% CI. Three adjustment models were used: model 1 adjusted for age, model 2 adjusted for age and BMI, and model 3 adjusted for model 2 and further adjusted for education level, marital status, family income, smoking, drinking alcohol, sleeping, physical activity, cardiovascular disease, hypertension, diabetes, and CRP levels. A *P*-value < 0.05 was considered statistically significant.

## Results

### Characteristics of the subjects

Among 21,376 men aged  $\geq$ 40 years old (mean  $51.9 \pm 10.0$  years), men with at least mild impaired kidney function ( $n = 19,519$ ) had a mean of eGFR  $77.7 \pm 8.6$  mL/min/1.73 m<sup>2</sup> (data not shown). Men in the lowest tertile (T1) of eGFR tended to be older, heavier, and had higher prevalence of cardiovascular disease, hypertension, and diabetes mellitus (57.5, 46.7, and 44.6% respectively,  $P < 0.001$ ) than those in the higher tertiles (T2 and T3) of eGFR (Table 1). In contrast, men in the lowest tertile of eGFR had lower serum testosterone levels compared to those in the higher tertiles of eGFR ( $4.6 \pm 1.3$  vs.  $4.8 \pm 1.4$  and  $5.4 \pm 1.9$  nmol/L in T1 vs. T2 and T3,  $P = 0.001$ ).

### Dietary patterns

Table 2 shows three dietary patterns identified by PCA. The cumulative percentage of variance explained by three factors was 34.1% (13.8, 11.6, and 8.7% for each factor, respectively). The three dietary patterns were named based on the interpretation and components of the pattern structure. The first factor was named as the fried-processed dietary pattern and characterized by frequent consumption of ten food groups: deep-fried foods, preserved or processed foods, sauce, organ meats, sugar-sweetened beverages, meats, jam or honey, fried rice or flour products, instant noodles, and eggs. The second factor was named as the vegetable dietary pattern and consisted of six food groups: dark-colored vegetables, light-colored vegetables, oil added vegetables or salad, seafood, legumes or

**Table 1** Characteristics of the subjects across tertiles of eGFR,  $n = 21,376^a$ 

	Total	Tertile of eGFR <sup>b</sup>			<i>P</i> <sup>c</sup>
		T1 ( $n = 7,128$ )	T2 ( $n = 7,149$ )	T3 ( $n = 7,099$ )	
Demographic data, lifestyle, and medical history					
Age, years	51.9 ± 10.0	58.0 ± 10.9	50.7 ± 8.5	47.1 ± 6.7	< 0.001
Education level, <i>n</i> (%)					< 0.001
< High school	3583 (17.0)	1898 (53.0)	1012 (28.2)	673 (18.8)	
High school	9354 (44.4)	2854 (30.5)	3169 (33.9)	3331 (35.6)	
> High school	8150 (38.6)	2254 (27.7)	2885 (35.4)	3011 (36.9)	
Marital status, <i>n</i> (%)					< 0.001
Never married	767 (3.8)	148 (19.3)	247 (32.2)	372 (48.5)	
Married	18412 (91.0)	6134 (33.3)	6207 (33.7)	6071 (33.0)	
Widowed/divorced	1048 (5.2)	455 (43.4)	320 (30.5)	273 (26.1)	
Family income, <i>n</i> (%)					< 0.001
< 800,000 NTD	7395 (37.7)	2975 (40.2)	2295 (31.0)	2125 (28.8)	
810,000–1.6 million NTD	8197 (41.8)	2338 (28.5)	2873 (35.1)	2986 (36.4)	
> 1.61 million NTD	4036 (20.6)	1103 (27.3)	1471 (36.5)	1462 (36.2)	
Smoking, <i>n</i> (%)					< 0.001
Never	11469 (55.4)	4017 (35.0)	3816 (33.3)	3636 (31.7)	
Past	3140 (15.2)	1120 (35.7)	1078 (34.3)	942 (30.0)	
Current	6086 (29.4)	1728 (28.4)	2046 (33.6)	2312 (38.0)	
Drinking alcohol, <i>n</i> (%)					0.001
No	14648 (73.0)	4950 (33.8)	4856 (33.1)	4842 (33.1)	
Yes	5405 (27.0)	1678 (31.1)	1855 (34.3)	1872 (34.6)	
Sleeping, <i>n</i> (%)					< 0.001
< 6 h	4828 (23.1)	1747 (36.2)	1572 (32.6)	1509 (31.2)	
6–7 h	10776 (51.5)	3407 (31.6)	3681 (34.2)	3688 (34.2)	
> 7 h	5324 (25.4)	1798 (33.8)	1753 (32.9)	1774 (33.3)	
Physical activity, <i>n</i> (%)					0.07
No	12554 (66.9)	4066 (32.4)	4213 (33.6)	4275 (34.0)	
Yes	6199 (33.1)	2073 (33.4)	2118 (34.2)	2008 (32.4)	
Cardiovascular disease, <i>n</i> (%)	1298 (6.5)	746 (57.5)	329 (25.3)	223 (17.2)	< 0.001
Hypertension, <i>n</i> (%)	6533 (44.0)	3051 (46.7)	1941 (29.7)	1541 (23.6)	< 0.001
Diabetes mellitus, <i>n</i> (%)	2356 (11.0)	1052 (44.6)	666 (28.3)	638 (27.1)	< 0.001
Anthropometric measurements					
BMI (kg/m <sup>2</sup> )	24.7 ± 3.1	24.9 ± 3.0	24.6 ± 3.0	24.5 ± 3.2	< 0.001
Systolic pressure (mmHg)	124.7 ± 16.7	128.1 ± 17.9	124.0 ± 16.2	122.0 ± 15.4	< 0.001
Diastolic pressure (mmHg)	76.9 ± 11.2	77.9 ± 11.5	76.8 ± 11.1	75.9 ± 11.0	< 0.001
Biochemical data					
FBG (mmol/L)	6.0 ± 1.3	6.1 ± 1.5	6.0 ± 1.2	5.9 ± 1.3	< 0.001
TG (mmol/L)	1.7 ± 1.2	1.7 ± 1.1	1.6 ± 1.1	1.7 ± 1.4	0.33
TC (mmol/L)	5.3 ± 0.9	5.3 ± 0.9	5.3 ± 0.9	5.2 ± 0.9	0.001
HDL-C (mmol/L)	1.4 ± 0.3	1.3 ± 0.3	1.4 ± 0.3	1.4 ± 0.3	0.001
LDL-C (mmol/L)	3.2 ± 0.8	3.2 ± 0.8	3.2 ± 0.8	3.1 ± 0.8	< 0.001
TC/HDL-C ratio	4.0 ± 0.9	4.0 ± 0.9	4.0 ± 0.9	3.9 ± 0.9	< 0.001
Testosterone (nmol/L) <sup>d</sup>	5.0 ± 1.6	4.6 ± 1.3	4.8 ± 1.4	5.4 ± 1.9	0.001
CRP (nmol/L)	23.2 ± 47.5	25.7 ± 48.9	22.3 ± 54.3	21.3 ± 37.2	< 0.001
Kidney function biomarkers					
BUN (mmol/L)	5.4 ± 1.4	6.0 ± 1.8	5.2 ± 1.1	4.9 ± 1.0	< 0.001
Creatinine (μmol/L)	100.8 ± 19.6	113.5 ± 29.0	98.4 ± 5.3	90.4 ± 4.1	< 0.001
eGFR (mL/min/1.73 m <sup>2</sup> )	75.4 ± 11.3	63.0 ± 7.7	74.5 ± 3.0	83.2 ± 3.4	< 0.001
Urinary protein, <i>n</i> (%)					< 0.001

**Table 1** Characteristics of the subjects across tertiles of eGFR,  $n = 21,376^a$  (Continued)

	Total	Tertile of eGFR <sup>b</sup>			<i>P</i> <sup>c</sup>
		T1 ( $n = 7,128$ )	T2 ( $n = 7,149$ )	T3 ( $n = 7,099$ )	
+ 1	20260 (94.8)	6481 (32.0)	6909 (34.1)	6870 (33.9)	
+ 2	641 (3.0)	311 (48.5)	163 (25.4)	167 (26.1)	
+ 3	303 (1.4)	188 (62.0)	63 (20.8)	52 (17.2)	
+ 4	172 (0.8)	148 (86.1)	14 (8.1)	10 (5.8)	

NTD new taiwan dollar, BMI body mass index, FBG fasting blood glucose, TG triglycerides, TC total cholesterol, HDL-C high density lipoprotein-cholesterol, LDL-C low density lipoprotein-cholesterol, TC/HDL-C total cholesterol-to-high density lipoprotein-cholesterol, CRP C-reactive protein, BUN blood urea nitrogen, eGFR estimated glomerular filtration rate

<sup>a</sup> Continuous data are presented as mean  $\pm$  SD, and categorical data are presented as number (percentage)

<sup>b</sup> Tertiles of eGFR, T1:  $\leq 71.7$  mL/min/1.73 m<sup>2</sup>; T2: 71.8–81.2 mL/min/1.73 m<sup>2</sup>; T3:  $\geq 81.3$  mL/min/1.73 m<sup>2</sup>

<sup>c</sup> *P* was analyzed using general linear model for continuous variables and chi-square test for categorical variables

<sup>d</sup> Testosterone: all,  $n = 256$ ; T1,  $n = 78$ ; T2,  $n = 91$ ; T3,  $n = 87$

beans, and rice or flour products. The last factor was named as the dairy-grain dietary pattern and consisted of six food groups: dairy products, milk, bread, root crops, fruits, and whole grains. Fruits, meats, and root crops had a factor-loading  $\geq 0.30$  in the fried-processed and vege-seafood dietary patterns; while jam or honey was appeared in the vege-seafood and dairy-grain dietary patterns. However, these food groups could only belong to one factor (the one with a greater factor loading value).

### Testosterone and kidney function

The association between serum testosterone levels and kidney function is demonstrated in Table 3. After multivariable adjustment, serum testosterone levels among men in the lower tertiles (T1 and T2) of eGFR were significantly decreased by approximately 0.8 and 0.9 nmol/L ( $\beta = -0.80$ , 95% CI:  $-1.40, -0.20$  and  $\beta = -0.88$ , 95% CI:  $-1.43, -0.33$ ,  $P < 0.05$ ) compared to those in the highest tertile (T3) of eGFR. Furthermore, BMI was negatively correlated with testosterone levels ( $\beta = -0.11$ , 95% CI:  $-0.19, -0.04$ ,

**Table 2** Factor loadings of three dietary patterns identified by principal component analysis<sup>a</sup>

	Fried-processed dietary pattern	Vege-seafood dietary pattern	Dairy-grain dietary pattern
Dark-colored vegetables	–	0.756	–
Light-colored vegetables	–	0.742	–
Fruits	–	–	0.403
Root crops	–	–	0.47
Legumes/beans	–	0.431	–
Fried rice/flour products	0.47	–	–
Preserved/processed foods	0.599	–	–
Organ meats	0.501	–	–
Sauce	0.567	–	–
Instant noodles	0.424	–	–
Deep-fried foods	0.663	–	–
Meats	0.485	–	–
Eggs	0.39	–	–
Seafood	–	0.444	–
Dairy products	–	–	0.657
Milk	–	–	0.586
Bread	–	–	0.499
Whole grains	–	–	0.392
Sugar-sweetened beverages	0.501	–	–
Jam or honey	0.478	–	–
Rice or flour products	–	0.398	–
Oil added vegetables/salad	–	0.567	–

<sup>a</sup> Factor loading below 0.30 was not shown in the table for simplicity. The food group with smaller factor loading value was eliminated as appearing in more than one dietary factor



$P < 0.01$ ), particularly obese men had significantly decreased testosterone levels ( $\beta = -0.88$ , 95% CI:  $-1.52, -0.25$ ,  $P < 0.01$ ) compared to men with normal BMI (data not shown). TG and TC/HDL-C ratio were also negatively associated with testosterone levels ( $\beta = -0.51$ , 95% CI:  $-0.77, -0.24$  and  $\beta = -0.42$ , 95% CI:  $-0.75, -0.08$ ,  $P < 0.05$ ). Correspondingly, age-adjusted or age and BMI-adjusted linear regression showed that serum testosterone levels were positively associated with eGFR levels ( $\beta = 0.83$ , 95% CI 0.17, 1.49 and  $\beta = 0.68$ , 95% CI: 0.01, 1.36,  $P < 0.05$ ) (Additional file 1: Table S1). Moreover, after multivariable adjustment, age and overweight or obese BMI were negatively correlated with eGFR ( $\beta = -0.57$ , 95% CI:  $-0.59, -0.55$ ,  $\beta = -1.17$ , 95% CI:  $-1.63, -0.71$ , and  $\beta = -1.18$ , 95% CI:  $-1.55, -0.80$ ,  $P < 0.001$  for all). All blood lipid components were negatively associated with eGFR levels ( $P < 0.001$ ), but HDL-C levels were positively associated with eGFR levels ( $\beta = 1.82$ , 95% CI: 1.26, 2.39,  $P < 0.001$ ) (Additional file 1: Table S1).

#### Dietary patterns, testosterone levels, and severity of impaired kidney function

Table 4 shows the association between dietary pattern scores across tertiles and serum testosterone levels. After multivariable adjustment, the fried-processed dietary pattern was significantly associated with decreased testosterone levels by 0.8 nmol/L ( $\beta = -0.78$ , 95% CI:  $-1.40, -0.16$ ,  $P < 0.05$ ) and reduced testosterone-to-triglycerides (T/TG) ratio by 1.8 units

( $\beta = -1.76$ , 95% CI:  $-2.99, -0.53$ ,  $P < 0.01$ ) among men in the higher tertile (T2) of dietary pattern scores compared to those in the lowest tertile (T1) of dietary pattern scores. However, the vege-seafood or dairy-grain dietary pattern was not significantly correlated with testosterone levels or T/TG ratio. Moreover, the fried-processed dietary pattern was positively associated with BUN and blood creatinine levels ( $\beta = 0.09$ , 95% CI: 0.04, 0.15 and  $\beta = 0.94$ , 95% CI: 0.08, 1.80,  $P < 0.05$ ), but negatively associated with eGFR ( $\beta = -0.64$ , 95% CI:  $-1.05, -0.24$ ,  $P < 0.01$ ) among men in the higher tertile (T2) of dietary pattern scores. In contrast, compared with the lowest tertile (T1), the higher tertiles (T2 and T3) of vege-seafood dietary pattern scores were negatively correlated with creatinine levels, but positively correlated with eGFR levels.

Furthermore, a multivariable logistic regression analysis revealed that men in the highest tertile (T3) of fried-processed dietary pattern scores were significantly associated with 35 and 18% increased risks of moderate/severe impaired kidney function and proteinuria severity respectively, compared to those in the lowest tertile (T1) (Fig. 2). In contrast, the highest tertile (T3) of vege-seafood dietary pattern scores had inverse association with moderate/severe impaired kidney function (OR = 0.82, 95% CI: 0.69, 0.97,  $P < 0.05$ ) and proteinuria severity (OR = 0.85, 95% CI: 0.73, 0.99,  $P < 0.05$ ). However, after

**Table 3** Multivariable linear regression analysis of independent variables affecting serum testosterone levels (nmol/L),  $n = 256$

	Model 1 <sup>a</sup>		Model 2 <sup>b</sup>		Model 3 <sup>c</sup>	
	$\beta$ (95% CI)	<i>P</i>	$\beta$ (95% CI)	<i>P</i>	$\beta$ (95% CI)	<i>P</i>
Age (years)	0.01 (-0.02, 0.03)	0.63	0.00 (-0.02, 0.03)	0.78	0.01 (-0.03, 0.04)	0.78
BMI (kg/m <sup>2</sup> )	-0.11 (-0.18, -0.05)	< 0.001	-0.11 (-0.18, -0.05)	< 0.001	-0.11 (-0.19, -0.04)	0.004
FBG (mmol/L)	-0.15 (-0.32, 0.01)	0.07	-0.09 (-0.26, 0.08)	0.30	-0.21 (-0.53, 0.11)	0.20
TG (mmol/L)	-0.49 (-0.68, -0.30)	< 0.001	-0.42 (-0.61, -0.22)	< 0.001	-0.51 (-0.77, -0.24)	< 0.001
TC (mmol/L)	-0.12 (-0.34, 0.09)	0.25	-0.07 (-0.28, 0.14)	0.50	-0.14 (-0.42, 0.13)	0.31
HDL-C (mmol/L)	0.69 (0.00, 1.37)	0.05	0.36 (-0.34, 1.06)	0.31	0.53 (-0.29, 1.35)	0.20
LDL-C (mmol/L)	0.04 (-0.21, 0.28)	0.77	0.09 (-0.15, 0.32)	0.48	-0.02 (-0.34, 0.31)	0.93
TC/HDL-C ratio	-0.36 (-0.59, -0.13)	0.002	-0.23 (-0.48, 0.01)	0.06	-0.42 (-0.75, -0.08)	0.014
BUN (mmol/L)	-0.08 (-0.25, 0.10)	0.38	-0.07 (-0.24, 0.10)	0.40	0.01 (-0.19, 0.21)	0.92
Creatinine ( $\mu$ mol/L)	-0.03 (-0.04, -0.01)	0.015	-0.02 (-0.04, 0.00)	0.049	-0.02 (-0.04, 0.01)	0.13
eGFR (mL/min/1.73 m <sup>2</sup> )						
T3	Ref		Ref		Ref	
T2	-0.72 (-1.18, -0.26)	0.002	-0.62 (-1.08, -0.17)	0.007	-0.88 (-1.43, -0.33)	0.002
T1	-0.94 (-1.44, -0.44)	0.001	-0.81 (-1.31, -0.31)	0.002	-0.80 (-1.40, -0.20)	0.010

BMI body mass index, FBG fasting blood glucose, TG triglycerides, TC total cholesterol, HDL-C high density lipoprotein-cholesterol, LDL-C low density lipoprotein-cholesterol, TC/HDL-C total cholesterol-to-high density lipoprotein-cholesterol, BUN blood urea nitrogen, eGFR estimated glomerular filtration rate

<sup>a</sup> Model 1: adjusted for age

<sup>b</sup> Model 2: adjusted for age and BMI

<sup>c</sup> Model 3: adjusted for model 2 and for education level, marital status, family income, smoking, drinking alcohol, sleeping, physical activity, cardiovascular disease, hypertension, diabetes, and CRP levels

**Table 4** Multivariable linear regression analysis of across tertiles of dietary pattern scores associated with serum testosterone and kidney function biomarkers <sup>a</sup>

	Fried-processed dietary pattern				Vege-seafood dietary pattern			
	T2	P	T3	P	T2	P	T3	P
	β (95% CI)		β (95% CI)		β (95% CI)		β (95% CI)	
Testosterone (nmol/L)	-0.78 (-1.40, -0.16)	0.014	-0.19 (-0.86, 0.48)	0.58	-0.05 (-0.82, 0.71)	0.89	0.04 (-0.73, 0.81)	0.92
T/TG ratio	-1.76 (-2.99, -0.53)	0.005	-0.81 (-2.14, 0.53)	0.24	0.54 (-0.99, 2.07)	0.48	0.94 (-0.60, 2.49)	0.23
BUN (mmol/L)	0.09 (0.04, 0.15)	0.001	0.06 (-0.00, 0.12)	0.06	-0.00 (-0.06, 0.05)	0.89	-0.03 (-0.09, 0.03)	0.36
Creatinine (μmol/L)	0.94 (0.08, 1.80)	0.031	0.21 (-0.67, 1.09)	0.64	-1.03 (-1.89, -0.17)	0.018	-1.47 (-2.32, -0.61)	0.001
eGFR (mL/min/1.73 m <sup>2</sup> )	-0.64 (-1.05, -0.24)	0.002	-0.39 (-0.81, 0.03)	0.07	0.46 (0.05, 0.86)	0.027	1.01 (0.60, 1.41)	< 0.001

T/TG testosterone-to-triglycerides, BUN blood urea nitrogen, eGFR estimated glomerular filtration rate

<sup>a</sup> Data were analyzed by model 3 adjusted for age, BMI, education level, marital status, family income, smoking, drinking alcohol, sleeping, physical activity, cardiovascular disease, hypertension, diabetes, and CRP levels using T1 as the reference group ( $n = 21,376$ , except for testosterone levels and T/TG ratio,  $n = 256$ ). Dairy-grain dietary pattern is not listed in the table because there was no association between the dairy-grain dietary pattern and all the variables above

adjusting for model 3, the vege-seafood dietary pattern was not significantly correlated with proteinuria severity (OR = 0.93, 95% CI: 0.77, 1.12,  $P = 0.46$ , data not shown). Similarly, the association between proteinuria severity and the fried-processed dietary pattern was altered after further adjustment in model 2 (OR = 1.06, 95% CI: 0.91, 1.23,  $P = 0.46$ , data not shown). Moreover, the highest tertile (T3) of dairy-grain dietary pattern scores were also significantly associated with reduced risk of proteinuria severity (OR = 0.78, 95% CI: 0.63, 0.95,  $P < 0.05$ ), but not significantly correlated with the severity of impaired kidney function.

## Discussion

### Testosterone and kidney function

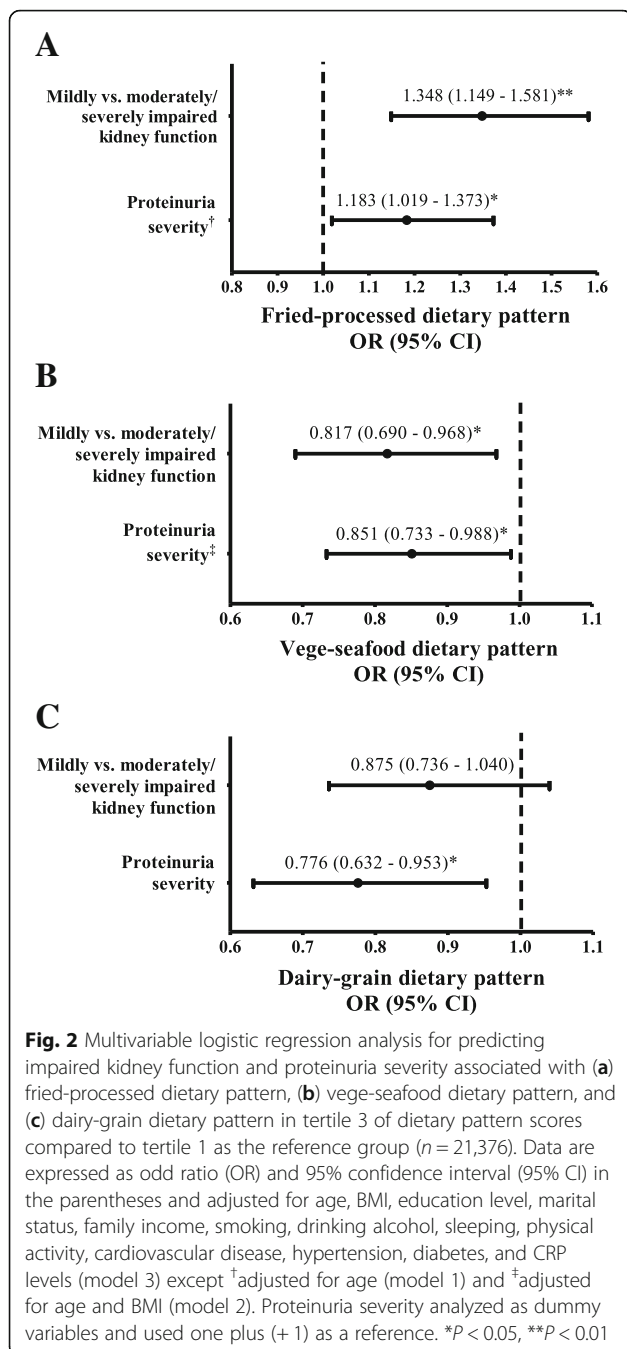
Our study found that men aged  $\geq 40$  years with lower tertiles of eGFR were associated with a decrease in serum testosterone levels (T2 and T1  $\beta = -0.80$ , 95% CI -1.40, -0.20 and  $\beta = -0.88$ , 95% CI -1.43, -0.33,  $P < 0.05$ , respectively). The association between kidney function and testosterone levels in men has been reported in previous studies [7, 24–27]. Japanese adult men with low salivary testosterone levels (22.0 pg/mL, the 5th percentile) were associated with reduced eGFR ( $-3.43$  mL/min/1.73 m<sup>2</sup>, 95% CI:  $-6.02$ ,  $-0.84$  mL/min/1.73 m<sup>2</sup>) compared to those in the 90th percentile of salivary testosterone levels (156.9 pg/mL) [7]. In agreement, our study also found that 1 nmol/L increment of serum testosterone levels were correlated with increased eGFR levels by 0.7 mL/min/1.73 m<sup>2</sup> after adjustment by age and BMI. However, there are other confounders that potentially influenced these results (Additional file 1: Table S1). Furthermore, our study did not find any association between serum testosterone levels and the risk of impaired kidney function or proteinuria severity (Additional file 1: Table S2) likely due to the small number of men who had serum testosterone data.

The possible mechanisms for the association between impaired kidney function and low testosterone levels are multi-factorial including uremia [28] and increased prolactin levels [29]. Comorbid conditions such as hypertension and diabetes may also contribute to reduce the levels of testosterone [30]. Uremia, commonly seen in patients with impaired kidney function, could also impair the hypothalamic-pituitary-gonadal axis and reduce testosterone production [31]. In addition, a previous study showed that prolactin levels were inversely associated with kidney function (eGFR) due to the reduction of prolactin clearance by the kidney [29]. Both uremia and prolactin negatively affect gonadal function by inhibiting the secretion of gonadotropin-releasing hormone and luteinizing hormone for testosterone production in the Leydig cells [32]. Our study also found that TG and TC/HDL-C ratio were independently associated with decreased testosterone levels. Consistent with our findings, previous studies have shown a negative correlation between testosterone levels and TG levels in men with impaired kidney function [25, 33]. Decreased testosterone levels by 41%, accompanied by 58% increase in TG levels were reported in men with advanced CKD stage compared to those in early CKD stage [25]. Additionally, lower testosterone levels and higher TG levels contribute to a rapid loss of kidney function [34] and all-cause mortality [5, 6, 24, 27]. Therefore, maintaining normal testosterone levels might have a beneficial effect to prevent a rapid decline in kidney function and CKD related outcomes.

### Dietary patterns and testosterone levels

To the best of our knowledge, our study is the first study to investigate the association between dietary patterns and testosterone levels in middle-aged and elderly men with impaired kidney function. In this study, we demonstrated that the fried-processed dietary pattern was independently associated with decreased testosterone levels ( $\beta = -0.78$ , 95% CI:  $-1.40$ ,  $-0.16$ ,  $p < 0.05$ ) and T/TG





ratio ( $\beta = -1.76$ , 95% CI:  $-2.99, -0.53$ ,  $P < 0.01$ ). In our study, the fried-processed dietary pattern consisted of food groups that are high in calories, saturated fat, artificial sweets, and sodium. Previous prospective studies have described the association between low caloric diets and testosterone levels [35–37]. Plasma total testosterone levels have been reported to be significantly increased after an 8-week low caloric diet (850–900 kcal/d) in obese nondiabetic men [35]. In a clinical trial of overweight or obese men on low fat and energy-

restricted diet (~ 1600 kcal/d) for 52 weeks, significant improvements in serum testosterone levels were found [13]. Limited published articles have investigated specific dietary patterns with male testosterone levels. A recent cross-sectional study in 125 Taiwanese men reported that higher adherence to a typical high-calorie Western diet was associated with decreased testosterone level by 0.87 ng/mL [38]. However, another study found that men consuming vegetarian diets had no association with serum testosterone levels [14, 15]. Consistently, we did not find an association between vege-seafood dietary pattern and testosterone levels. Elevated testosterone levels seem to occur in overweight or obese men following a low-fat or low-calorie diet, presumably due to weight loss and less aromatization of testosterone to estradiol [39]. However, the potential links between dietary patterns and testosterone levels are unclear, but might be due to Western diet-related insulin resistance [38]. Insulin stimulates testosterone production and insulin resistance may impair this process in the Leydig cells [40]. Whereas, previous studies were conducted in smaller samples, hence further larger prospective studies are needed to assess the effect of dietary patterns on male testosterone levels.

#### Dietary patterns and kidney function

Our study also found that the fried-processed dietary pattern was positively associated with increased risk of moderate/severe impaired kidney function and proteinuria severity. However, the association between the fried-processed dietary pattern with proteinuria severity was altered after further adjustment in model 2 (OR = 1.06, 95% CI: 0.91, 1.23,  $P = 0.46$ ) and model 3 (OR = 1.02, 95% CI: 0.83, 1.25,  $P = 0.88$ ). Previous studies have reported that subjects with higher adherence to typical Western dietary patterns were associated with increased creatinine levels and decreased eGFR levels [9–11, 41, 42]. The Nurses' Health Study has found that high adherence to a Western dietary pattern, characterized by high intake of meat, processed meat, saturated fat, and sweets, was positively correlated with rapid eGFR decline and albuminuria [9–11]. Two recent studies among Singaporean Chinese and US adults also addressed that red meat and processed meat were associated with the onset of CKD and kidney failure [41, 42]. In addition, a Southern dietary pattern, consisted of primarily fried-processed foods, organ meats, and sweetened beverages, was correlated with increased mortality in CKD subjects [43]. In general, animal-based protein with high phosphorus and biological value could produce acid metabolites and increase dietary acid load, which may provoke the kidney to excrete excess acid. A dietary acid load was found to be positively

associated with the progression of advanced CKD stage and albuminuria [44, 45]. In addition, our study also showed that there was a negative association between dairy-grain dietary pattern and proteinuria severity. Differences in the direction between protein-rich dietary patterns and kidney dysfunction could possibly be explained by the composition of amino acids and fatty acids between non-dairy and dairy foods. Dairy foods contain large amounts of branched-chain amino acids, which have a lesser effect on kidney function [46].

In our study, the vege-seafood dietary pattern had an inverse association with the severity of impaired kidney function and proteinuria. Consistent with our findings, a cross-sectional study in Taiwan demonstrated that a dietary pattern with high intake of fish and vegetables had a positive association with kidney function [12]. Diets rich in whole grains, fruits, and low-fat dairy products were also correlated with lower risks of albuminuria [46] and mortality [47]. Moreover, a large prospective cohort study reported that plant sources of protein; like legumes/nuts, dairy protein, fish, and seafood reduced the risk of CKD [42]. Plant-based protein is predominantly alkaline-forming. Diets high in plant-based protein, fruits, and vegetables have been reported to be associated with higher blood bicarbonate levels, lower blood pressure, reduced renal acid load, decreased uremic toxin and inflammatory markers, and reduced serum creatinine and proteinuria [46, 48–51]. Seafood that is rich in polyunsaturated fatty acids has anti-inflammatory properties, which may also improve kidney function [52].

### Strengths and limitations

The strength of this study is a relatively large study representative that was used to determine the association between dietary patterns and severity of impaired kidney function in men. Additionally, we used the CKD-EPI equation to estimate eGFR levels due to its higher accuracy than other equations [19]. However, present study had certain limitations. First, due to the cross-sectional study design, our study cannot establish a causal relationship between dietary patterns and severity of impaired kidney function. A temporal relationship may occur as we observed in the present study, and could be affected by the dietary advice given to all participants due to their disease conditions. Furthermore, subjects who had received health screenings may have been health-conscious and changed their dietary habits. Second, data availability of other male sex hormones such as sex hormone binding globulin, free and albumin-bound testosterone are limited. Measurement of other testosterone forms

may be more sensitive to determine male sex hormone status. Moreover, we used the data from the full sample to derive dietary patterns, whereas the testosterone analysis was conducted on a very small subset (1.2% of the full sample). Third, the self-reported food frequency questionnaire may have self-reporting bias and measurement errors. Although a food frequency questionnaire is the most common method used to evaluate dietary patterns, given its feasibility and affordability in nutritional epidemiological studies, we cannot neglect that self-reported dietary assessment methods are known for both omission (e.g. unreported or underreporting food that was eaten) and intrusion errors (e.g. lack of memories when reporting). However, in the present study, we used a validated and standardized semi-quantitative food frequency questionnaire that has been published elsewhere [21, 22]. Nevertheless, future studies using rigorous scientific methods such as randomized trials are needed to investigate the role of dietary patterns in the development of chronic diseases. Fourth, due to the lack of quantitative data concerning urine protein, we used urinary protein dipstick as a marker of proteinuria instead of measuring albuminuria, which is more standardized measurement of kidney damage. Moreover, clinically diagnosed CKD was defined as a patient with eGFR < 60 ml/min/1.73 m<sup>2</sup> for ≥3 months, with or without persistent albuminuria [20]. Although MJ Health Institute provides periodic health screenings to its members, not all subjects had an annual examination. Hence, we cannot identify clinically diagnosed CKD based on one single measurement only. Therefore, our results may not necessarily reflect the clinically diagnosed CKD subjects, and not generalize to those with impaired kidney function and without clinical proteinuria. Finally, some potential confounders such as total energy and protein intake might affect the study findings.

### Conclusion

In conclusion, our findings suggest that frequent consumption of fried-processed dietary pattern is associated with decreased testosterone levels and T/TG ratio as well as increased risk of impaired kidney function and proteinuria severity. In contrast, the vege-seafood dietary pattern is negatively associated with moderately/severely impaired kidney function and proteinuria severity, but has no association with either testosterone levels or T/TG ratio. Further studies with prospective measurements are needed to elucidate the association between dietary patterns and testosterone levels in men with impaired kidney function.

## Additional file

**Additional file 1: Table S1.** Multivariable linear regression analysis of independent variables affecting eGFR (mL/min/1.73 m<sup>2</sup>), *n* = 21,376. **Table S2.** Multivariable logistic regression analysis for predicting moderate/severe impaired kidney function and proteinuria severity associated with serum testosterone and lipid profiles <sup>a</sup>. (DOCX 27 kb)

### Abbreviations

BMI: Body mass index; BUN: Blood urea nitrogen; CKD: Chronic kidney disease; CRP: C-reactive protein; CVD: Cardiovascular disease; DBP: Diastolic blood pressure; eGFR: Estimated glomerular filtration rate; FBG: Fasting blood glucose; HDL-C: High density lipoprotein-cholesterol; LDL-C: Low density lipoprotein-cholesterol; PCA: Principal component analysis; SBP: Systolic blood pressure; TC: Total cholesterol; TC/HDL-C: Total cholesterol-to-HDL-C ratio; TG: Triglycerides

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### Authors' contributions

ALK and JJC conceived and designed the study; CYH and HHR managed the dataset and retrieved the data; LYL revised the language of the entire manuscript; ALK analyzed and interpreted the data; ALK and JJC wrote the manuscript. All authors have read and approved the final manuscript.

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### Availability of data and materials

The data that support the findings of this study are available from Mei Jau (MJ) Health Institute, but restricted for research use only. The data are not publicly available. Data are available from the authors upon reasonable request and with permission of MJ Health Institute.

### Ethics approval and consent to participate

The study was approved by the Taipei Medical University-Joint Institutional Review Board (TMU-JIRB) no. 201802006. All the participants signed a written informed consent authorized by Mei Jau (MJ) Health Institute.

### Consent for publication

The data provided by Mei Jau (MJ) Health Institute to the researchers did not include any personal information, and all participants were adults. Not applicable.

### Competing interests

The authors declare that they have no competing interests.

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