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METABOLIC SYNDROME - AN EMERGING RISK FACTOR FOR CHRONIC KIDNEY DISEASE

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ABSTRACT

Objective: Metabolic syndrome (MetS) is a common health problem worldwide. According to third national health and nutrition examination survey criteria, about 47 million people have MetS. It is defined as having three or more of the following five risk factors including abdominal obesity, increased triglyceride levels, low-density lipoprotein cholesterol level, elevated blood pressure, and elevated fasting glucose levels. These components of MetS are major risk factors for the development of chronic kidney disease (CKD) also. CKD is a major public problem and it is a major risk factor for the development of cardiovascular disease. Hence, the aim of the current study was to evaluate the association between MetS and CKD.

Methods: A total of 50 patients reporting to Saveetha Dental College and Hospitals were enrolled into the study which includes 25 patients with MetS and 25 healthy individuals. 5 mL of venous blood was collected and centrifuged. Then, it is analyzed for fasting blood sugar (FBS), serum triglycerides, serum urea, and creatinine using the standard kit method. The data obtained were subjected to statistical analysis using the SPSS software.

Results: The mean body mass index, FBS, serum creatinine, and triglyceride levels were higher in MetS patients in comparison to healthy individuals. The mean body mass index (BMI), FBS, serum urea, serum creatinine, and triglyceride levels in the control group and MetS group were 27.75±3.67, 84.8±12.5, 17.52±5.2, 0.91±0.17, and 96.5±60.13 and 35.14±4.25, 108.8±34.69, 21.4±5.9, 1.0±0.14, and 239.76±51.21, respectively. There was a significant difference in the mean BMI, FBS, urea, creatinine, and triglyceride levels of the above group.

Conclusion: Serum urea and creatinine levels were significantly higher in MetS individuals. Hence, MetS could be a one of the risk factors for the development of CKD.

Keywords: Metabolic Syndrome, Chronic Kidney Disease, Urea, Creatinine.

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INTRODUCTION

Metabolic syndrome (MetS) previously called syndrome X is defined by a constellation of interconnected physiological, biochemical, clinical, and metabolic factors that directly increases the risk of cardiovascular disease, type 2 diabetes mellitus, and all cause of mortality [1,2]. MetS is a state of chronic low-grade inflammation as a consequence of complex interplay between genetic and environmental factors. Insulin resistance, visceral adiposity, atherogenic dyslipidemia, endothelial dysfunction, genetic susceptibility, elevated blood pressure, hypercoagulable state, and chronic stress are the several factors which constitute the MetS [3-5].

There are currently two major definitions for MetS provided by the International Diabetes Federation (IDF) and the revised National Cholesterol Education Program, respectively.

IDF [6]

Central adiposity (waist circumference ≥ 90 cm for men, ≥ 80 cm for women) and two or more of the following:

- Fasting triglycerides (TGL) >150 mg/dL or on specific medication
- HDL cholesterol <40 mg/dL and <50 mg/dL for men and women, respectively, or on specific medication
- Blood pressure >130 mm systolic or >85 mm diastolic or previous diagnosis or on specific medication
- Fasting plasma glucose ≥100 mg/dL or previously diagnosed type 2 diabetes.

National Cholesterol Education Program and Adult Treatment Panel III (NCEP: ATP III)

Three or more of the following

- Central obesity waist circumference >102 cm (M), >88 cm (F)
- Hypertriglyceridemia TGL ≥ 150 mg/dL or specific medication

- Low HDL cholesterol <40 mg/dL and <50 mg/dL for men and women, respectively, or on specific medication
- Hypertension blood pressure ≥130 mm systolic or ≥85 mm diastolic or on specific medication
- Fasting plasma glucose ≥100 mg/dL or on specific medication previously diagnosed type 2 diabetes.

Despite MetS is associated with many disorders, it has been clearly associated with chronic kidney disease (CKD) also. Each component of MetS has been associated with both CKD incidence and progression [7]. Major component of MetS is insulin resistance. Insulin is an anti-inflammatory hormone. Hence, insulin resistance leads to inflammation, oxidative stress, and renal insufficiency [8,9]. Each and every component of MetS increases the oxidative stress and reactive oxygen species which leads to renal insufficiency and CKD [9,10].

CKD is recognized as a major public health problem. It is defined as a glomerular filtration rate <60 mL/min/1.73 m² and/or markers of kidney damage for at least 3 months. It is assessed by estimating serum creatinine concentration and the urine albumin concentration [11-13]. The major causes for the development of CKD are hypertension, diabetes, obesity, and other causes are anemia, hyperphosphatemia, high plasma C-reactive protein and fibrinogen, accumulation of endogenous inhibitors of nitric oxide synthase, and high sympathetic activity [14,15].

The major causes for the development of CKD are the major components of the development of MetS also. Hence, the aim of this study was evaluation of the association between MetS and CKD.

Table 1: BMI, FBS, TGL, urea, and creatinine levels in two groups

Parameters	Mean±SD		p value
	Control	Metabolic Syndrome (MetS)	
BMI	27.75±3.67	35.14±4.25	0.0001
FBS	84.8±12.5	108.8±34.69	0.002
TGL	96.6±60.13	239.76±51.21	0.0001
Urea	17.52±5.2	21.4±5.9	0.007
Creatinine	0.91 ± 0.17	1.0 ± 0.14	0.047

BMI: Body mass index, FBS: Fasting blood sugar, TGL: Triglycerides

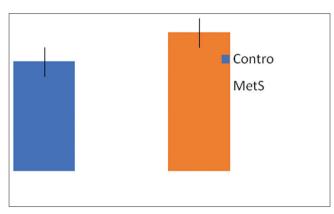


Fig. 1: Mean body mass index level in two groups

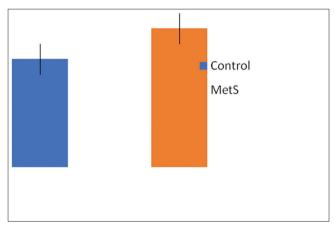


Fig. 2: Mean fasting blood sugar levels in two groups

METHODS

Patients were selected from those attending the Outpatient Department of Saveetha Dental College and Hospitals and divided into two groups as follows:

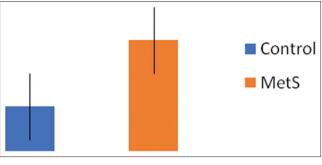
- Group I normal healthy individuals 25 individuals
- Group II patients with MetS 25 individuals.

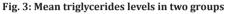
Inclusion criteria

- Individuals with the age group of 35–55 years.
- Individuals with type 2 diabetes mellitus (fasting blood sugar [FBS] ≥100 mg/dL).
- Individual with normal body mass index (BMI) (18.9–24.9) and obese BMI (≥30).
- Serum TGL levels more than 150 mg/dL.

Exclusion criteria

• Individuals with other systemic illness such as cardiovascular disease, renal failure, stroke, and endocrine illness.





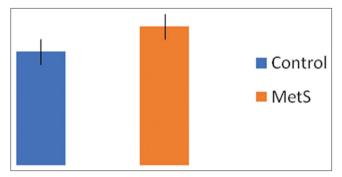


Fig. 4: Mean urea levels in two groups

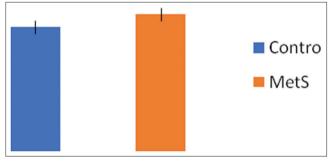


Fig. 5: Mean creatinine levels in two groups

- Individuals with acute illness like fever.
- · Immunocompromised individuals.

Sample collection

Informed consent was obtained from the patient before sample collection. 5 mL of fasting venous blood was collected and distributed in plain collection tubes and centrifuged in 3000 rpm for serum. Then, serum was separated, and then, it is analyzed for FBS by GOD-POD method, serum TGL by colorimetric enzymatic method, serum urea by urease method, and creatinine by Jaffe's method using ERBA CHEM 5 plus analyzer.

RESULTS

All the data were analyzed using SPSS package. Student *t*-test analysis was done to find out significant differences between the two groups. All the tests were considered statistically significant at p<0.05 level.

DISCUSSION

The results of this study revealed that there is a significant difference in mean BMI, FBS, TGL, urea, and creatinine levels of the normal control group and MetS group.

Mean BMI level in control and MetS groups is 27.75±3.67 and 35.14±4.25, respectively (Table 1 and Fig. 1). Mean serum FBS level in control and MetS groups is 84.8±12.5 and 108.8±34.69, respectively (Table 1 and Fig. 2). Mean serum TGL level in control and MetS groups is 96.5±60.13 and 239.76±51.21, respectively (Table 1 and Fig. 3). Mean serum urea level in control and MetS groups is 17.52±5.2 and 21.4±5.9, respectively (Table 1 and Fig. 4). Mean serum creatinine level in control and MetS groups is 0.91±0.17 and 1.0±0.14, respectively (Table 1 and Fig. 5).

There was a significant difference between mean BMI, FBS, and TGL levels. That indicates right selection of individuals in both the groups. There was a statistically significant difference between mean serum urea (p=0.007) and mean serum creatinine (p=0.047) levels also.

Our results provide evidence for a positive correlation between urea, creatinine levels in control and MetS individuals. That indicates MetS individuals are prone to develop CKD [16]. Many studies show renal dysfunction becomes apparent long before the appearance of hypertension or diabetes in MetS.

In increased BMI (obesity), there is increased flux of free fatty acids into the liver leading to excessive hepatic production of TGL which leads to hypertriglyceridemia. And also, adipocytes secrete inflammatory cytokines such as tumor necrosis factor- α , interleukin-6, and C-reactive protein [17,18]. Hypertriglyceridemia itself causes for insulin resistance. In diabetes mellitus, increased fasting glucose level leads to hyperinsulinemia and insulin resistance, which leads to excessive release of inflammatory cytokines, which cause glomerular membrane thickening, endothelial dysfunction, renin-aldosterone system activation, oxidative stress, and elevated plasminogen activator inhibitor-1. Ultimately leads to glomerulosclerosis, tubulointerstitial injury, and CKD [19-21].

CONCLUSION

The present study shows urea and creatinine levels were consistently higher in MetS individuals. These data have indicated that MetS increases the risk of CKD through multiple pathways. Hence, early identification and treatment of individual components of MetS may reduce the risk of the development of CKD. Hence, MetS could be emerging risk factor for the development of CKD.

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