



Close association between circulating high-sensitivity cardiac troponin I and metabolic syndrome in the general population

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Abstract

Individuals with metabolic syndrome reportedly have an increased risk of cardiovascular disease, although the association between asymptomatic myocardial damage and metabolic syndrome has not been sufficiently investigated. The present study investigated possible associations between circulating cardiac troponin and metabolic syndrome or related factors. Subjects undergoing their annual health checkups were enrolled in the study ($n = 1242$). Laboratory measurements included serum high-sensitivity cardiac troponin I (hs-cTnI) and plasma B-type natriuretic peptide (BNP). Individual salt intake was estimated by calculating 24-h urinary sodium excretion from spot urine. Subjects whose electrocardiograms revealed ST-T segment abnormalities or who had renal insufficiency or a history of cardiovascular events were excluded. Subjects with metabolic syndrome had higher hs-cTnI levels than those without, but their BNP levels were equivalent. hs-cTnI levels were significantly associated with the presence and components of metabolic syndrome. Logistic regression analysis with the endpoint of hs-cTnI levels higher than the median value identified metabolic syndrome as an independent determinant of increased hs-cTnI levels. Additionally, urinary salt excretion levels were increased in subjects with metabolic syndrome or any of its components. Logistic regression analysis with the endpoint of metabolic syndrome revealed that hs-cTnI levels were independently associated with the presence of metabolic syndrome. A close association between hs-cTnI levels and the presence of metabolic syndrome, at least partially mediated by increased salt intake, was confirmed to exist in the general population. The findings support the idea that patients with metabolic syndrome develop asymptomatic myocardial damage without obvious ischaemic findings, which leads to increased cardiovascular risk.

Keywords Metabolic syndrome · High-sensitivity cardiac troponin I · Myocardial damage · Blood pressure · Urinary salt excretion

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Introduction

Metabolic syndrome is diagnosed based on body weight gain and accumulation of visceral fat accompanied by multiple cardiovascular risk factors [1–3]. The diagnostic criteria for metabolic syndrome comprise the following four components: visceral fat obesity indicated by elevated waist circumference, lipid metabolism disorder, elevated blood pressure (BP) and abnormal fasting glucose due to insulin resistance [4, 5]. Approximately 20–40% of adults in Western countries have been shown to have metabolic syndrome, although only ~10% of adults in Japan have been diagnosed with that condition [6–9]. The basic pathophysiology of metabolic syndrome is based on insulin resistance and inflammation that lead to sympathetic nerve activation, increased fluid retention and metabolic disorder and contribute to macrovascular and microvascular disease

[10–16]. This may be one of the reasons why subjects with metabolic syndrome reportedly face an increased risk of cardiovascular disease, including coronary artery disease and heart failure, as well as all-cause mortality [9, 15–18].

Myocardial damage is apparent in patients with myocardial infarction, including acute coronary syndrome, and is often confirmed to be present in both acute and non-acute ischaemic myocardial disease [19, 20]. Alternatively, myocardial damage can progress asymptotically through non-ischaemic mechanisms such as cardiac overload and inflammation [21, 22], with diagnosis based solely on circulating levels of constitutively expressed myocardial proteins such as the cardiac troponins detected incidentally in clinical settings [23, 24]. Elevated levels of cardiac troponins are also reported to occur with left ventricular overload caused by elevated BP [25, 26]. However, there are limited data on persistent myocardial damage in metabolic syndrome without obvious ischaemic conditions [27, 28]. We thus hypothesized that metabolic syndrome could be a cause of asymptomatic myocardial damage among patients in stable condition with no obvious ischaemic findings. To this end, the present study investigated possible associations between circulating high-sensitivity cardiac troponin I (hs-cTnI) and the presence of metabolic syndrome or related factors in individuals without obvious ischaemic findings on resting electrocardiograms.

Methods

The present study enrolled subjects attending their annual physical checkups for a study protocol approved by the ethics committees of Nagoya City University Graduate School of Medical Sciences and Enshu Hospital. The study was performed in accordance with the principles of the Declaration of Helsinki. Written informed consent was obtained from all subjects prior to their participation in the study.

Subjects

Subjects who visited Enshu Hospital in 2015–16 for an annual health checkup were screened for eligibility to participate in the present study. Subjects with renal dysfunction (creatinine ≥ 1.5 mg/dL), cancer, active inflammatory disease, or a history of cardiovascular events (stroke, myocardial infarction and heart failure) were excluded, as were subjects with obvious ST segment or T wave abnormality, implanted pacemakers or frequent arrhythmia (including atrial fibrillation and atrial flutter) on a standard 12-lead electrocardiogram. Since high-intensity physical activity can influence hs-cTnI levels,

subjects who engaged in hard physical labour or high-intensity exercise were also excluded.

Participants were instructed to collect overnight urine in a paper cup and to submit a sample of the urine in a plastic tube. Salt intake was assessed by estimating 24-h urinary sodium excretion, calculated using the Kamata formula [29], which estimates 24-h sodium excretion based on lean body mass and the composition of overnight urine in the Japanese population, achieving a correlation coefficient of 0.78 between measured and estimated sodium excretion [29]. Then, blood samples were taken early in the morning after an overnight fast for laboratory measurements, including serum concentrations of hs-cTnI. Systolic and diastolic BPs were measured in the non-dominant arm of seated subjects using a validated oscillometric technique (HEM-7070; Omron Corporation, Kyoto, Japan). Three consecutive BP measurements were taken at 2-min intervals, and the mean of the second and third measurements was recorded as the BP. Subjects taking antihypertensive medications or having systolic BP ≥ 140 mmHg and diastolic BP ≥ 90 mmHg were defined as having hypertension [30]. Subjects taking lipid-lowering medications or having high-density lipoprotein cholesterol (HDL-C) levels < 40 mg/dL, low-density lipoprotein cholesterol (LDL-C) levels ≥ 140 mg/dL or triglycerides ≥ 150 mg/dL were defined as having dyslipidaemia [31]. Subjects taking antihyperglycaemic medication or presenting a fasting plasma glucose (FPG) level ≥ 126 mg/dL were defined as having diabetes [32]. Metabolic syndrome was defined based on the Japanese diagnostic criteria (abdominal obesity, with waist circumference ≥ 85 cm for men or ≥ 90 cm for women, and two or more of the following three criteria: (1) triglycerides ≥ 150 mg/dL and/or HDL-C < 40 mg/dL; (2) systolic BP ≥ 130 mmHg and/or diastolic BP ≥ 85 mmHg; and (3) FPG ≥ 110 mg/dL) [4].

Biochemical analysis

Biochemical tests including determination of serum total cholesterol, LDL-C, HDL-C, and triglycerides were performed using standard laboratory assays. Plasma B-type natriuretic peptide (BNP) concentrations were determined using a commercially available chemiluminescence enzyme immunoassay (MI02 Shionogi BNP Kit; Shionogi, Osaka, Japan). Circulating levels of hs-cTnI were measured by the ARCHITECT high-sensitive troponin I assay according to the manufacturer's instructions (Abbott, Tokyo, Japan).

Statistical analysis

Data were analysed using IBM SPSS Statistics 19 (IBM Corp., Chicago, IL, USA). Dichotomous variables (gender, smoking status and presence or absence of metabolic

syndrome) were assigned values of 0 (female, non-smoker and absence of metabolic syndrome) or 1 (male, smoker and presence of metabolic syndrome). Data with a normal distribution are expressed as the mean \pm standard deviation. Data that are not normally distributed (BNP and hs-cTnI) are expressed as the median with interquartile range and are evaluated after log transformation. Comparative analyses of continuous variables were performed using *t* tests. Differences in continuous variables among more than three groups were tested using analysis of variance followed by a post hoc Scheffe test. Logistic regression analysis determined the independent variables associated with increased hs-cTnI and the presence of metabolic syndrome. Receiver operating characteristic (ROC) curve analyses were used to select a cut-off level for hs-cTnI. The median value of hs-cTnI was used as the cut-off value defining increased hs-cTnI. A two-tailed $P < 0.05$ value was considered significant.

Results

Of the 1242 subjects enrolled in the study, the numbers (percentages of the total) with hypertension, dyslipidaemia, diabetes mellitus and obesity (body mass index >25 kg/m²) were 390 (31.4%), 617 (49.7%), 95 (7.6%) and 238 (19.2%), respectively. In addition, 148 subjects (11.9%), comprising 115 (9.3%) males and 33 (2.7%) females, fulfilled the diagnostic criteria for metabolic syndrome (Table 1).

Subjects with metabolic syndrome had higher levels of hs-cTnI than those without, while BNP levels were equivalent in those with and without metabolic syndrome (Table 1). Similarly, subjects with any given component of metabolic syndrome or with at least one of the components of metabolic syndrome and abdominal obesity (the preliminary conditions for metabolic syndrome) had significantly higher levels of hs-cTnI than those without (Figs. 1 and 2a), and subjects with the preliminary conditions for metabolic syndrome had higher levels of hs-cTnI than normal subjects even after the exclusion of hypertensive subjects (Fig. 2b). Logistic regression analysis examined how the presence of metabolic syndrome affected the elevation of hs-cTnI, with the endpoint of hs-cTnI levels higher than the median value. The analysis showed that the presence of metabolic syndrome was an independent determinant of increased hs-cTnI levels after adjustment for age and gender (Table 2, adjusted Model 1) and further adjustment for smoking status and creatinine (Table 2, adjusted Model 2). Additional analysis showed that the presence of metabolic syndrome was significantly associated with increased hs-cTnI levels after adjustment for medications (Table 2, adjusted Model 3).

Table 1 Characteristics of subjects

Variable	Total subjects (<i>n</i> = 1242)	Subjects without metabolic syndrome (<i>n</i> = 1094)	Subjects with metabolic syndrome (<i>n</i> = 148)
Age (years)	58 \pm 12	58 \pm 12	62 \pm 10*
Male gender, <i>n</i> (%)	791 (63.7)	676 (61.8)	115 (77.7) [†]
Current smoker, <i>n</i> (%)	239 (19.2)	205 (18.7)	34 (23.0)
BMI (kg/m ²)	22.6 \pm 3.1	22.2 \pm 2.9	25.9 \pm 3.0*
Waist circumference (cm)	83.8 \pm 8.4	82.5 \pm 7.8	93.4 \pm 6.0*
Systolic BP (mmHg)	123 \pm 14	122 \pm 14	132 \pm 11*
Diastolic BP (mmHg)	75 \pm 9	74 \pm 9	79 \pm 9*
Creatinine (mg/dL)	0.79 \pm 0.16	0.79 \pm 0.17	0.83 \pm 0.15 [†]
FPG (mg/dL)	97 \pm 16	95 \pm 14	108 \pm 25*
HDL-C (mg/dL)	61 \pm 16	62 \pm 16	49 \pm 12*
LDL-C (mg/dL)	123 \pm 28	123 \pm 27	123 \pm 29
Triglyceride (mg/dL)	104 \pm 56	97 \pm 49	159 \pm 73*
BNP (pg/mL)	16.5 [8.5–29.8]	16.7 [8.6–29.9]	15.9 [8.3–28.2]
Urinary salt excretion (g/day)	9.1 \pm 2.0	9.0 \pm 2.0	10.1 \pm 2.3*
hs-cTnI (pg/mL)	2.4 [1.6–3.8]	2.3 [1.5–3.7]	3.4 [2.2–4.6]*
Medications, <i>n</i> (%)			
ACE-I or ARB	184 (14.8)	114 (10.4)	70 (47.3)*
β -blockers	25 (1.4)	15 (1.4)	10 (6.8)*
Calcium channel blocker	199 (16.0)	139 (12.7)	60 (40.5)*
Diuretics	24 (1.9)	19 (1.7)	5 (3.4)
Statin	187 (15.1)	108 (9.9)	79 (53.4)*
Hypoglycaemia agent	74 (6.0)	59 (4.1)	29 (19.6)*
Antithrombotic agent	55 (4.4)	34 (3.1)	21 (14.2)*
Complications, <i>n</i> (%)			
Hypertension	390 (31.4)	278 (25.4)	112 (75.7)*
Dyslipidemia	617 (49.7)	469 (42.9)	148 (100)*
Diabetes mellitus	95 (7.6)	59 (5.4)	36 (24.3)*
Obesity (BMI >25 kg/m ²)	238 (19.2)	157 (14.4)	81 (54.7)*

Data are presented as the mean \pm standard deviation, median [interquartile range] or as *n* (%)

BMI body mass index, BP blood pressure, FPG fasting plasma glucose, HDL-C high-density lipoprotein cholesterol, LDL-C low-density lipoprotein cholesterol, BNP B-type natriuretic peptide, hs-cTnI high-sensitivity cardiac troponin I, ACE-I angiotensin-converting enzyme inhibitor, ARB angiotensin receptor blocker

* $P < 0.0001$, [†] $P < 0.01$ vs. subjects without metabolic syndrome

We then examined the effects of urinary salt excretion levels on metabolic syndrome based on the reported association of salt intake with insulin resistance and fluid retention. The analysis showed that urinary salt excretion

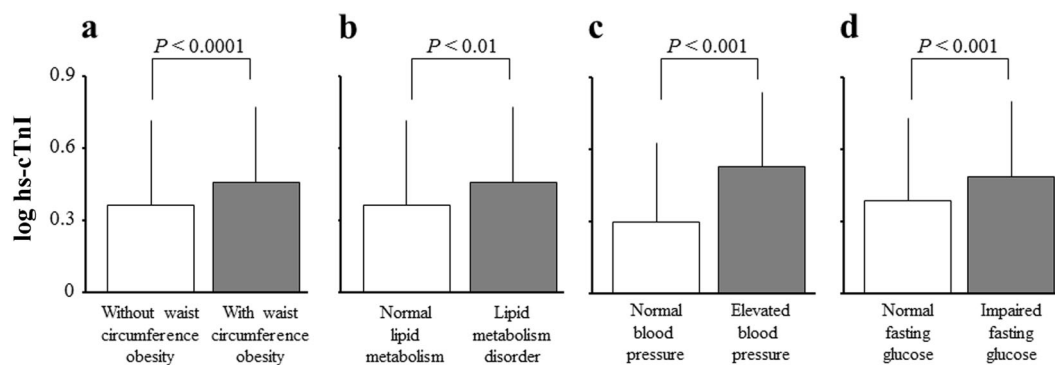
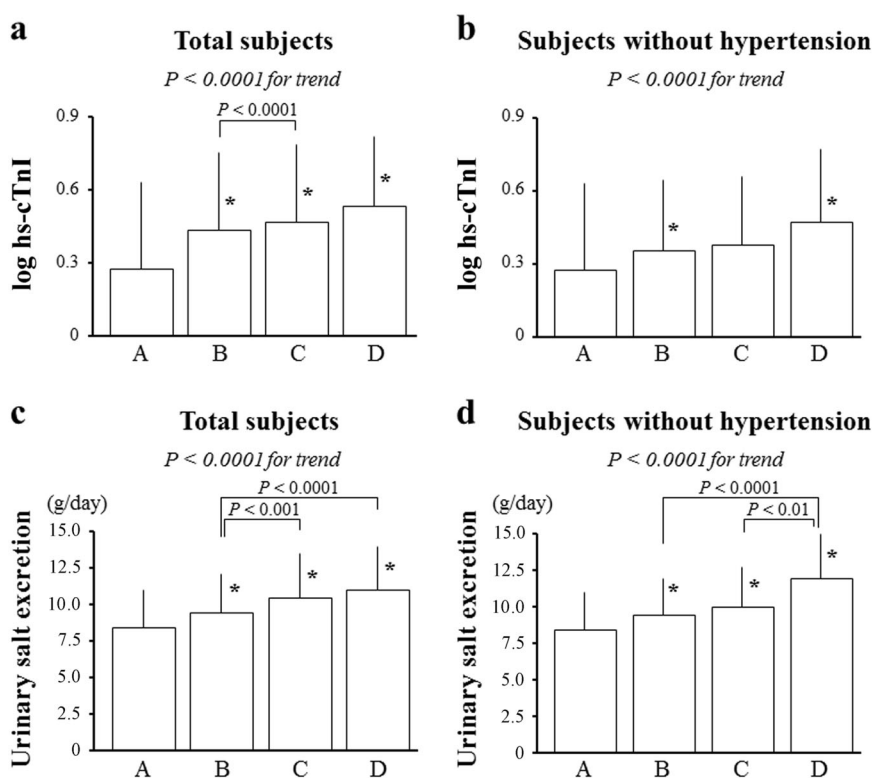


Fig. 1 Log-transformed circulating high-sensitivity cardiac troponin I (hs-cTnI) levels in subjects with and without the following components of metabolic syndrome: **a** abdominal obesity (waist circumference ≥ 85 cm for men and ≥ 90 cm for women), **b** lipid metabolism disorder (triglycerides ≥ 150 mg/dL and/or high-density

lipoprotein cholesterol < 40 mg/dL), **c** elevated blood pressure (systolic pressure ≥ 130 mmHg and/or diastolic pressure ≥ 85 mmHg), and **d** abnormal fasting plasma glucose (fasting plasma glucose ≥ 110 mg/dL). Data are shown as the mean \pm standard deviation

Fig. 2 Log-transformed circulating high-sensitivity cardiac troponin I (hs-cTnI) levels in normal subjects without any of the components of metabolic syndrome (Group A), subjects with at least one of the components of metabolic syndrome excluding abdominal obesity (Group B), subjects with at least one of the components of metabolic syndrome including abdominal obesity (Group C), and subjects with metabolic syndrome (Group D), assessed **a** regardless of blood pressure and **b** in subjects without hypertension. Log-transformed urinary salt excretion levels in Group A, Group B, Group C and Group D assessed **c** regardless of blood pressure and **d** in subjects without hypertension. Data are shown as the mean \pm standard deviation. * $P < 0.0001$ vs. normal subjects (Group A)



levels were higher in subjects with metabolic syndrome or the preliminary conditions for metabolic syndrome than in normal subjects (Fig. 2c), even after excluding subjects with hypertension from the analysis (Fig. 2d). ROC curve analysis yielded 2.6 pg/ml as the cut-off hs-cTnI level indicating metabolic syndrome (area under the curve 0.658, $P < 0.01$, 95% confidence interval 0.630–0.684), sensitivity 66.9% (95% confidence interval 58.7–74.4), specificity 59.0% (95% confidence interval: 56.0–61.9) (Supplementary Figure). Logistic regression analysis with the presence of metabolic syndrome as the endpoint showed that hs-cTnI levels were independently associated with metabolic

syndrome after adjustment for age and gender (Table 3, adjusted Model 1) and further adjustment for smoking status and creatinine (Table 3, adjusted Model 2). Urinary salt excretion was added to the analysis in adjusted Model 2 and was also identified as an independent determinant of metabolic syndrome (Table 3, adjusted Model 3).

Discussion

The main findings of the present study are as follows: (i) subjects with metabolic syndrome had higher levels of

Table 2 Results of logistic regression analysis demonstrating factors possibly associated with higher concentrations of high-sensitivity cardiac troponin I than the median value

Variable	OR (95% CI)	P value
Unadjusted		
Metabolic syndrome	2.61 (1.80–3.78)	<0.0001
Adjusted Model 1		
Metabolic syndrome	1.75 (1.16–2.63)	<0.01
Age (years)	1.09 (1.09–1.11)	<0.0001
Male gender	2.44 (1.86–3.19)	<0.0001
Adjusted Model 2		
Metabolic syndrome	1.75 (1.16–2.64)	<0.01
Age (years)	1.10 (1.08–1.11)	<0.0001
Male gender	2.17 (1.52–3.12)	<0.0001
Current smoking	0.90 (0.63–1.27)	0.53
Creatinine (mg/dL)	2.05 (0.71–5.89)	0.18
Adjusted Model 2		
Metabolic syndrome	1.75 (1.16–2.64)	<0.01
Age (years)	1.10 (1.08–1.11)	<0.0001
Male gender	2.17 (1.52–3.12)	<0.0001
Current smoking	0.90 (0.63–1.27)	0.53
Creatinine (mg/dL)	2.05 (0.71–5.89)	0.18
Adjusted Model 3		
Metabolic syndrome	1.73 (1.05–2.86)	<0.05
Age (years)	1.09 (1.07–1.11)	<0.0001
Male gender	1.90 (1.30–2.78)	<0.01
Current smoking	0.92 (0.64–1.33)	0.66
Creatinine (mg/dL)	2.39 (0.76–7.51)	0.14
ACE-I or ARB	1.25 (0.73–2.12)	0.42
β-blockers	0.76 (0.19–3.15)	0.71
Calcium channel blocker	1.31 (0.79–2.16)	0.30
Diuretics	2.18 (0.56–8.57)	0.26
Statin	0.93 (0.61–1.44)	0.76
Hypoglycaemia agent	0.99 (0.52–1.87)	0.97
Antithrombotic agent	1.50 (0.39–5.77)	0.56

OR odds ratio, CI confidence interval, ACE-I angiotensin-converting enzyme inhibitor, ARB angiotensin receptor blocker

Adjusted Model 1 included age and male gender. Adjusted Model 2 was further adjusted for current smoking and creatinine besides Model 1. Adjusted Model 3 was further adjusted for medications besides Model 2

hs-cTnI, but not BNP, than those without; (ii) subjects with any given component of metabolic syndrome or at least one component of metabolic syndrome showed significantly higher levels of hs-cTnI and urinary salt excretion than those without; (iii) logistic regression analysis with the endpoint of hs-cTnI levels higher than the median value showed that the presence of metabolic syndrome was an independent determinant of increased hs-cTnI; and (iv) logistic regression analysis with the endpoint of

Table 3 Results of logistic regression analysis demonstrating factors possibly associated with metabolic syndrome

Variable	OR (95% CI)	P value
Unadjusted		
hs-cTnI (pg/ml)	3.55 (2.18–5.79)	<0.0001
Adjusted Model 1		
hs-cTnI (pg/ml)	2.16 (1.22–3.82)	<0.01
Age (years)	1.03 (1.01–1.05)	<0.01
Male gender	1.92 (1.27–2.91)	<0.01
Adjusted Model 2		
hs-cTnI (pg/mL)	2.15 (1.21–3.82)	<0.01
Age (years)	1.03 (1.01–1.05)	<0.01
Male gender	1.82 (1.10–3.01)	<0.05
Current smoking	1.27 (0.81–1.99)	0.31
Creatinine (mg/dL)	0.97 (0.29–3.25)	0.95
Adjusted Model 3		
hs-cTnI (pg/ml)	1.93 (1.08–3.45)	<0.05
Age (years)	1.02 (1.004–1.042)	<0.05
Male gender	1.67 (0.99–2.78)	0.051
Current smoking	1.61 (1.01–2.56)	<0.05
Creatinine (mg/dL)	1.16 (0.34–3.91)	0.81
Urinary salt excretion (g/day)	1.26 (1.16–1.38)	<0.0001

OR odds ratio, CI confidence interval, hs-cTnI high-sensitivity cardiac troponin I

Adjusted Model 1 included age and male gender. Adjusted Model 2 was further adjusted for current smoking and creatinine besides Model 1. Adjusted Model 3 was further adjusted for urinary salt excretion besides Model 2

meeting the diagnostic criteria for metabolic syndrome showed that hs-cTnI was independently associated with the presence of metabolic syndrome after adjustment for possible confounding factors including urinary salt excretion. These results indicate that the presence of metabolic syndrome is significantly associated with asymptomatic myocardial damage through increased salt intake.

Metabolic syndrome is associated with high risks of cardiovascular disease and mortality [14–18], and although the underlying mechanism is not fully understood, the increased incidence of macrovascular disease, mainly coronary artery disease, is thought to be related [14]. Since insulin resistance and vascular inflammation promote atherosclerotic cardiovascular disease, metabolic syndrome is associated with the progression of atherosclerotic plaque and instability of the plaque, leading to obstructive coronary artery disease and acute coronary syndrome [14–18]. Although significant associations between cardiac troponin and metabolic syndrome were reported previously, subjects with ST-T segment abnormalities in the electrocardiogram had not been excluded from the previous studies; therefore, subjects with coronary artery disease may have been included [27, 28]. In contrast, microvascular disease relevant to metabolic syndrome could

be considered a small vessel disorder caused by microvascular endothelial dysfunction [11–14]. Such microvascular disease could develop silently in metabolic syndrome without increasing cardiac load and could thus cause asymptomatic cardiovascular disease. The independent association of hs-cTnI, but not BNP, with the presence of metabolic syndrome as demonstrated in the present study may be caused by microvascular disorder and suggests that subjects with metabolic syndrome should be treated carefully even in the absence of elevated BNP levels. In addition, if possible, hs-cTnI levels should be assessed repeatedly in individuals with metabolic syndrome. On the other hand, the median level of hs-cTnI was significantly higher in subjects with metabolic syndrome than in those without (3.4 vs. 2.3 pg/ml). The cut-off level of hs-cTnI (2.6 pg/ml) suggested by ROC curve analysis as a threshold for metabolic syndrome was only slightly higher than the median value in subjects without metabolic syndrome but was not sufficiently robust as a diagnostic indicator for single tests in a clinical setting. These results also suggest that attention should be paid to subjects with obesity or cardiovascular risk factors and hs-cTnI levels higher than the cut-off level, even in the absence of metabolic syndrome.

We previously found that left ventricular voltage measured by electrocardiography was independently associated with hs-cTnI levels even after adjustment for BNP [33]. In the present study, hs-cTnI levels were increased in subjects with at least one of the components of metabolic syndrome even when hypertensive subjects were excluded from the analysis. Moreover, the presence of metabolic syndrome was an independent determinant of increased hs-cTnI levels after adjustment for systolic and diastolic BP. Additionally, to adjust for possible confounding factors including all components of metabolic syndrome, we conducted further logistic regression analysis with the endpoint of hs-cTnI levels higher than the median value and found that the presence of metabolic syndrome was an independent determinant of high hs-cTnI (Supplementary Table 1). Moreover, similar logistic regression analysis with the endpoint of metabolic syndrome showed that hs-cTnI was independently associated with the presence of metabolic syndrome (Supplementary Table 2). Thus, metabolic syndrome significantly affects hs-cTnI levels independently of hypertension. These findings support the hypothesis that both cardiac load and metabolic disorder are causally associated with asymptomatic myocardial damage. Indeed, insulin resistance and metabolic disorder have been implicated in cardiovascular disease through the mechanisms of inflammation, oxidative stress, fibrosis and alteration in cardiac function [34–37]. Although an association between metabolic syndrome and systemic inflammation has been reported previously, the association between myocardial damage and inflammation during metabolic syndrome

remains to be intensively investigated [38, 39]. Evaluating the relationship between hs-cTnI and high-sensitivity C-reactive protein was beyond the scope of the present study; however, we speculate that systemic inflammation is one of the pathological pathways leading to myocardial injury in metabolic syndrome. Previous animal studies have implied that obesity and insulin resistance cause mitochondrial dysfunction and an increase in fatty acids, leading to increased oxidative stress, release of inflammatory cytokines such as tumour necrosis factor- α and interleukin-6, and a change in the balance of adipokines [40–42]. These disorders affecting the heart could contribute to asymptomatic myocardial damage in metabolic syndrome. On the other hand, urinary salt excretion levels were assessed with respect to metabolic syndrome in the present study, since obesity and insulin resistance might be associated pathophysiologically [13–16]. As expected, urinary salt excretion levels were increased in subjects with metabolic syndrome or its components even if hypertensive subjects were excluded, indicating that an increase in salt intake is an independent determinant of the presence of metabolic syndrome. Importantly, these results are in line with previous reports that sodium intake affects all components of metabolic syndrome [15, 16]. Moreover, sodium restriction was reported to reduce cardiac injury by reducing inflammation and oxidative stress in a rat model of metabolic syndrome [43, 44]. Therefore, metabolic syndrome might occur through a pathway relevant to sodium and volume retention through insulin resistance and inflammation, leading to asymptomatic myocardial damage. Conversely, hs-cTnI was an independent determinant of metabolic syndrome after adjustment for possible confounding factors. These findings emphasize the strong association between metabolic syndrome and asymptomatic myocardial damage.

Cardiac troponin levels are the mainstay of diagnosing acute coronary syndrome [18, 19] but are also used for differentiating myocardial damage from coronary artery disease [22, 45]. Indeed, the usefulness of cardiac troponins as predictors of cardiovascular events in non-ischaemic heart disease has been demonstrated [46, 47]. Thus, since a mild increase in cardiac troponins even within the normal range has been associated with future cardiovascular events, assessment of myocardial damage is meaningful in the general population, whose cardiac troponin levels would not be high [48]. Cardiac troponin is a complex of three types of regulatory proteins, called troponin T, troponin I, and troponin C. Cardiac troponin T and I are often used as cardiac biomarkers, whereas cardiac troponin C has complete amino acid homology with skeletal muscle troponin C and thus is not used as a cardiac biomarker [49]. In addition, cardiac troponin T levels can be elevated in patients with reduced renal function, and cardiac troponin I is superior to troponin T in the detection of electrocardiographic cardiac

injury [50]. Hence, we focused on troponin I to detect myocardial damage when investigating the relationship between hs-cTnI and metabolic syndrome. Therefore, findings obtained from the present study suggest that detecting patients with both metabolic syndrome and increased hs-cTnI levels is meaningful for identifying individuals with high cardiovascular risk and mortality.

The present study has several limitations, and the findings should thus be interpreted with caution. First, the present study was a cross-sectional study, and the background of the subjects enrolled in the study was heterogeneous. Second, the causal relationship between hs-cTnI levels and metabolic syndrome was not investigated, and biological assessments to demonstrate the mechanisms underlying the close association between hs-cTnI levels and metabolic syndrome *in vivo* are needed in the future. Third, enrolled subjects were asymptomatic and without ST-T abnormalities on their electrocardiograms, but neither the exercise stress test nor coronary angiography was performed to exclude ischaemic heart disease. Fourth, other variables that might affect metabolic syndrome or hs-cTnI, such as diet, physical activity, sleep, and genetic variance, were not taken into account. Further investigations with a larger population, a longitudinal design and detailed examination are also necessary for definite conclusions to be drawn.

In conclusion, a close association between circulating hs-cTnI levels and the presence of metabolic syndrome, at least partially mediated by increased salt intake, in the general population without electrocardiographic ST-T segment abnormalities. The significant association between asymptomatic myocardial damage and metabolic syndrome might be causally involved in high-risk cardiovascular disease and mortality in metabolic syndrome without obvious ischaemic findings.

Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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