

Relationship between N-terminal pro-B-type natriuretic peptide levels and metabolic syndrome

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Abstract

Introduction: Previous studies have shown that obese individuals have reduced natriuretic peptide levels. But conflicting data exist on the relation of natriuretic peptide levels to other metabolic risk factors.

Material and methods: We investigated the relationship between plasma N-terminal pro-B-type natriuretic peptide levels (NT-proBNP) and metabolic syndrome (MetS) and metabolic risk factors in 469 patients free of heart failure. Two hundred thirty diagnosed MetS cases and 239 non-MetS cases were included in this study. Echocardiography examinations were performed and left ventricular mass index was calculated according to the Devereux correction formula. NT-proBNP was measured by electrochemiluminescence. The log-transformed NT-proBNP levels were used for abnormal distribution. Multiple linear regression analysis was performed to assess the association between levels of NT-proBNP and metabolic factors. Covariance analysis was used for group comparisons.

Results: Log NT-proBNP levels were independently related to age, gender, body mass index, systolic blood pressure, diastolic blood pressure, fasting blood glucose, triglyceride, high density lipoprotein cholesterol, low density lipoprotein cholesterol, left ventricular mass index and left ventricular ejection fraction in multiple linear regression analysis ($p < 0.05$). Adjusted log NT-proBNP levels were lower in persons with MetS compared with those without MetS ($p < 0.05$). Individuals with hyperlipidaemia, elevated body mass index, diastolic blood pressure and fasting plasma glucose had lower levels of log NT-proBNP than those without MetS ($p < 0.05$).

Conclusions: There is a relationship between metabolic components and lower plasma NT-proBNP concentration. These findings raise the possibility that reduced plasma NT-proBNP levels are a manifestation of MetS, which might possess significant clinical and pathophysiological implications.

Key words: natriuretic peptide, hyperlipidaemia, left ventricular mass index, risk factors.

Introduction

The metabolic syndrome (MetS) is a worldwide problem, which refers to a constellation of coronary heart disease (CHD) risk factors including obesity and abdominal fat distribution, disorders of glucose and lipid metabolism, and hypertension. The burden of MetS is likely to continue to rise, largely due to decreased physical activity and increases in obesity in our society.

Obesity is associated with depressed N-terminal pro-B-type natriuretic peptide (NT-proBNP), a cardiac hormone that was recognized as a new and useful biochemical marker for both diagnosis and ruling out of congestive heart failure [1]. Despite the well-documented association between NT-proBNP levels and obesity, data on relations with other metabolic risk factors are mixed [2, 3]. Other studies have not found an association between plasma natriuretic peptide levels and hyperlipidaemia [4] or hyperglycaemia [5]. However, Olsen *et al.* [6] reported an inverse association between NT-proBNP levels and plasma lipids and glucose.

Thus, we sought to elucidate the relations between plasma NT-proBNP levels and metabolic risk factors and the metabolic syndrome in a Chinese study. We hypothesized that plasma NT-proBNP levels would be lower in the presence of the metabolic syndrome or its components.

Material and methods

Study population

We retrospectively evaluated the 250 MetS patients and 250 non-MetS cases who were admitted to the department of endocrinology of Huashan Hospital affiliated to Fudan University from July 2005 to August 2009. Participants were excluded for the following reasons: unavailable NT-proBNP levels ($n = 20$), hepatic (total bilirubin level > 5 mg/dl; $n = 0$) or renal insufficiency (creatinine > 2.0 mg/dl; $n = 13$), a history of heart failure ($n = 11$), and missing covariates ($n = 3$). After these exclusions, 469 subjects (94% of attendees) remained eligible. Written consent was obtained from all patients before the study.

Measurements

The subjects were interviewed for the documentation of medical histories (hypertension and diabetes) and medication, history of smoking habits (current, former or never), laboratory assessment of cardiovascular disease risk factors, and standardized echocardiographic examination. The body weight of the subjects, wearing light clothing and without their shoes on, was measured with 0.1 kg precision. Height was measured to the nearest 0.5 cm. Body mass index was calculated as the weight in kilograms divided by the square of height in metres. Systolic and diastolic blood pressure values were the means of two physician-obtained measurements on the left arm of the seated participant. Diabetes was defined by either a history of fasting glucose ≥ 126 mg/dl (7.0 mmol/l) or the use of insulin or hypoglycaemic medications.

Laboratory assays

Peripheral venous blood samples were taken into tubes in the fasting state in all subjects. It was

centrifuged at 3000 rpm for 10 min for plasma separation and immediately used to measure NT-proBNP and other parameters.

Fasting plasma glucose and postprandial blood glucose concentration (glucose oxidase method, GOX0560), serum triglycerides (GPO-PAP method, TGP0560), high density lipoprotein cholesterol and low density lipoprotein cholesterol (IRC method), and serum creatinine (Sarcosine Oxidase-PAP method, S708) of all the patients were estimated on an automated chemistry analyser (Hitachi 7600-020, Hitachi High-Technologies Corp, Shanghai, China). Glycated haemoglobin was estimated by high-pressure liquid chromatography using a machine (HLC-723G7, Tosoh, Shanghai, China). Plasma concentrations of NT-proBNP were measured by the electrochemiluminescence immunoassay "Roche Diagnostics Elecsys 2010 system" (USA), and the reference range was 0 pg/ml to 600 pg/ml.

MetS definition

We used the definition of MetS recommended by the Chinese Diabetes Society of the Chinese Medical Association for the Chinese population. Thus, individuals were defined as having MetS if they satisfied ≥ 3 of the following 4 criteria: (1) overweight or obese or elevated body mass index (≥ 25.0 kg/m²), (2) high fasting plasma glucose (≥ 6.1 mmol/l) and/or postprandial blood glucose (≥ 7.8 mmol/l) and/or use of hypoglycaemic medications, (3) high blood pressure systolic/diastolic $\geq 140/90$ mmHg or use of antihypertensive medications, (4) high fasting triglycerides (≥ 1.7 mmol/l) and/or reduced high density lipoprotein cholesterol (< 0.9 mmol/l in men, < 1.0 mmol/l in women).

Echocardiography

Echocardiography examinations were performed with a Vingmed System 5 Doppler echocardiographic unit (GE Vingmed Ultrasound, Horten, Norway). Conventional echocardiography measurements were performed according to American Society of Echocardiography guidelines. Left ventricular mass (LVM) was calculated using the Devereux formula as follows: $LVM (g) = 1.04 \times [(LVIDd + LVPWd + IVSd)^3 - LVIDd^3] - 13.6$, where LVIDd represents left ventricular diameter in end diastole (cm), LVPWd represents left ventricular posterior wall thickness in diastole (cm), and IVSd represents interventricular septal thickness in end diastole (cm). Left ventricular mass was corrected for body surface area (BSA) to give LVM index (LVMI). Left atrial diameter (LAD) and aortic root dimension (AOD) were also measured. Left ventricular systolic function was assessed by

calculation of left ventricular ejection fraction (LVEF). Diastolic function was assessed by determining the E-to-A ratio (E/A), where E and A represent the early and late velocities respectively.

Statistical analysis

We used natural logarithmic (Log) transformations of peptide levels because NT-proBNP followed approximately lognormal distributions. Furthermore, the association between the elevation of NT-proBNP and potential metabolic factors was investigated by multiple linear regression analysis. Multiple linear regression analysis was performed to relate Log NT-proBNP to metabolic variables that included gender, body mass index, high density lipoprotein cholesterol, low density lipoprotein cholesterol, systolic blood pressure, diastolic blood pressure, fasting plasma glucose, and postprandial blood glucose. This model also included age, serum creatinine, AOD, LAD, LVMI, LVEF and E/A.

Next, considering differences of NT-proBNP levels between MetS and non-MetS groups with disturbance of the possible confounding variables, we performed covariance analysis with adjustment for age, gender, serum creatinine, LAD, AOD, LVMI and LVEF which have been identified by multiple regression analysis.

In order to better investigate the association of NT-proBNP and the components of MetS, we compared NT-proBNP levels of different subgroups by covariance analysis.

Results were analysed using the Statistical Package for Social Sciences for Windows ver. 13.0 (SPSS, Chicago, IL, USA). Continuous variables were expressed as mean \pm SD or median (range) values. Tests were two-sided and a *p*-value of < 0.05 was considered significant.

Results

Characteristics of the study sample ($n = 469$) are shown in Table I. There are 124 males and 106 females (mean age, 65.2 ± 12.0 years) in the MetS group and 142 males and 97 females (mean age, 66.0 ± 12.6 years) in the non-MetS group. As expected, body mass index, systolic blood pressure, diastolic blood pressure, fasting blood glucose, postprandial blood glucose, and lipid and lipoprotein profile were significantly different between the MetS group and the non-MetS group ($p < 0.001$). Hypertension and diabetes were more prevalent in the patients with MetS ($p < 0.001$).

NT-proBNP and metabolic factors

Results of multiple linear regression models relating Log NT-proBNP with metabolic risk factors are shown in Table II. In all cases, a negative association was observed between plasma

NT-proBNP levels and body mass index ($p = 0.016$), as previously described. Additionally, NT-proBNP levels were inversely associated with diastolic blood pressure ($p = 0.015$), fasting blood glucose ($p = 0.046$), triglycerides ($p = 0.009$), low density lipoprotein cholesterol ($p = 0.036$), and LVEF ($p \leq 0.001$). NT-proBNP was positively associated with age ($p < 0.001$), systolic blood pressure ($p = 0.004$), high density lipoprotein cholesterol ($p = 0.013$) and LVMI ($p < 0.001$). Plasma NT-proBNP was negatively associated with systolic blood pressure ($p = 0.001$) and fasting blood glucose ($p = 0.015$) in women but not in men. However, the association for triglycerides ($p = 0.002$) and high density lipoprotein cholesterol ($p = 0.014$) was significant in men but not in women.

Log NT-proBNP levels in patients with MetS and without MetS

The differences between the two groups of patients in terms of their NT-proBNP levels are displayed in Figure 1 and Table III. Plasma levels of log NT-proBNP are lower in all participants with MetS after adjustment for age, gender (for total patients), serum creatinine, LAD, AOD, LVMI, and LVEF ($p = 0.001$). Similarly, the decrease of NT-proBNP levels in patients with MetS remained significant for men ($p = 0.038$) and for women ($p = 0.022$) after the same adjustment for age, and clinical and echocardiographic characteristics.

Log NT-proBNP levels in patients of MetS subgroups and non-MetS

Next, we divided the subjects with MetS into subgroups according to their body mass index, systolic blood pressure, diastolic blood pressure, fasting blood glucose, triglycerides or high density lipoprotein cholesterol. In men, the cut points are 25.0 kg/m^2 , 140 mmHg , 90 mmHg , 6.1 mmol/l , 1.7 mmol/l or 0.9 mmol/l respectively. In women, the cut points are similar to those of men except that high density lipoprotein cholesterol is 1.0 mmol/l . Covariance analyses were conducted and adjusted for age, gender (for total patients), serum creatinine, LAD, AOD, LVMI, and LVEF. In addition, for example, accounting for the other MetS components, such as dyslipidaemia, hyperglycaemia and elevated blood pressure, may influence comparison of log NT-proBNP levels between MetS subgroups stratified by BMI and the non-MetS group. So here the analyses of covariance were also adjusted for the other MetS components except for the categorical exposure variable (like BMI in the above-mentioned example). The MetS components above referred to body mass index, triglycerides, high density lipoprotein cholesterol, low density lipoprotein cholesterol, fasting blood glucose,

Table I. Clinical characteristics of the study participants with and without MetS

Parameters	MetS(+) n = 230	MetS(-) n = 239	Value of p
Sex, no., male/female	124/106	142/97	
Age [year]	65.2 ±12.0	66.0 ±12.6	0.689
Smoking [%]	42	43	0.624
Hypertension [%]	53	11	< 0.001
Diabetes mellitus [%]	72	30	< 0.001
Body mass index [kg/m ²]	27.0 ±4.5	23.3 ±4.2	< 0.001
FBG [mmol/l]	6.6 ±1.2	5.2 ±1.0	< 0.001
PBG [mmol/l]	9.3 ±3.0	7.3 ±2.8	< 0.001
Glycated haemoglobin [%]	6.8 ±0.8	5.8 ±0.7	< 0.001
Systolic blood pressure [mmHg]	144 (85)	130 (75)	< 0.001
Diastolic blood pressure [mmHg]	89 (45)	77 (35)	< 0.001
LDL cholesterol [mmol/l]	2.61 ±0.8	2.59 ±0.8	0.301
HDL cholesterol [mmol/l]	0.9 ±0.2	1.2 ±0.3	< 0.001
Triglycerides [mmol/l]	2.2 ±1.3	1.3 ±0.6	< 0.001
Serum creatinine [μmol/l]	79.5 ±25.1	79.3 ±22.7	0.175
LAD [mm]	40.0 ±7.1	38.2 ±5.9	0.238
AOD [mm]	33.0 ±3.4	33.0 ±4.5	0.642
LVM [g]	228.2 ±64.1	195.5 ±61.5	< 0.001
LVMI [g/m ²]	127.3 ±31.9	119.1 ±40.5	0.023
EF [%]	62.7 ±11.7	64.5 ±9.5	0.091
E/A ratio	1.3 ±0.2	1.4 ±0.2	0.689
Medication [%]			
Calcium antagonists	25.3	23.2	0.331
β-blockers	50.8	49.6	0.782
ACE inhibitors	35.4	33.1	0.534

FBG – fasting blood glucose, PBG – postprandial blood glucose, HDL cholesterol – high density lipoprotein cholesterol, LDL cholesterol – low density lipoprotein cholesterol, LAD – left atrial diameter, AOD – aortic root dimension, EF – ejection fraction, E – transmitral early diastolic velocity, A – transmitral late diastolic velocity, E/A – ratio of transmitral early (E) and late (A) diastolic velocities, IVS – interventricular septum, PW – posterior wall, LVM – left ventricular mass, LVMI – left ventricular mass index, values are mean ± SD when appropriate

Table II. Multivariable association of metabolic risk factors with log NT-proBNP levels

Independent variables	Total		Men		Women	
	β	p	β	p	β	p
Age	0.337	< 0.001	0.342	< 0.001	0.387	< 0.001
Gender	0.221	0.004				
Body mass index	-0.173	0.007	-0.252	0.003	-0.223	0.018
Systolic blood pressure	0.217	0.004	0.114	0.225	0.372	0.001
Diastolic blood pressure	-0.179	0.015	-0.186	0.034	-0.397	0.007
Fasting blood glucose	-0.150	0.046	-0.176	0.093	-0.276	0.015
Triglycerides	-0.265	0.009	-0.351	0.002	0.204	0.285
HDL cholesterol	0.240	0.013	0.317	0.014	0.014	0.922
LDL cholesterol	-0.423	0.036	-0.456	0.170	0.102	0.715
LVMI	0.315	< 0.001	0.292	0.001	0.452	< 0.001
LVEF	-0.377	< 0.001	-0.446	< 0.001	-0.372	< 0.001

FBG – fasting blood glucose, HDL cholesterol – high density lipoprotein cholesterol, LDL cholesterol – low density lipoprotein cholesterol, LVMI – left ventricular mass index, LVEF – left ventricular ejection fraction

systolic blood pressure and diastolic blood pressure when internalized as covariates.

The differences between patients of MetS subgroups and non-MetS in terms of their NT-proBNP levels are displayed in Figure 2 and Tables IV-VIII. For all subjects totally, Log NT-proBNP concentrations were lower in MetS subjects with dyslipidaemia, hyperglycaemia, elevated body mass index and diastolic blood pressure than those without MetS ($p < 0.05$), but did not reach significance in MetS subjects with normal blood fat, glucose, body mass index, systolic blood pressure and diastolic blood pressure compared to those without MetS ($p = \text{NS}$). However, Log NT-proBNP levels were higher in MetS patients with elevated systolic blood pressure than non-MetS patients ($p < 0.05$). Similar findings were obtained in men and women except for women with elevated body mass index.

Discussion

N-terminal pro-B-type natriuretic peptide (NT-proBNP), the inactive amino terminal fragment, and the biologically active form of B-type natriuretic peptide (BNP), are cleavage products of the precursor pre-proBNP which is synthesized by ventricular myocytes in response to physiological signals such as stretching of the ventricular wall or changes in systemic blood pressure, sodium levels or extracellular volume. B-type natriuretic peptide is metabolized by specific natriuretic peptide receptors found in the kidney, lung, liver, and along the vascular endothelium, while NT-proBNP is mainly cleared by the kidneys. It has recently been recognized as a new and useful biochemical marker for both diagnosis and ruling out of congestive heart failure (CHF) [1]. For the present, assessment of plasma NT-proBNP concentration is of potential value for risk of death and cardiovascular complications in the general population and may be useful for detection of a symptomatic left ventricular systolic dysfunction and systolic dysfunction in the general population [7, 8]. Recent clinical trials have also shown that NT-proBNP is elevated in patients with acute coronary syndrome (ACS) and exercise training in patients after myocardial infarction could decrease plasma NT-proBNP levels [9, 10]. Baghdady *et al.* observed an increase of NT-proBNP levels in ventricular septal defect with cardiac decompensation, and it is correlated with clinical score and echocardiography parameters [11].

NT-proBNP and metabolic syndrome

An inverse relationship between BNP and MetS has been described previously in the Framingham Heart Study [2]. In addition, levels of NT-proBNP

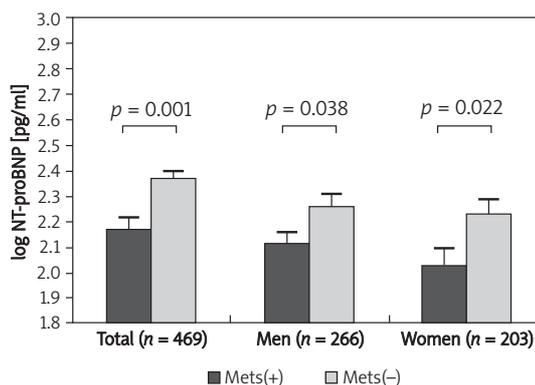


Figure 1 Adjusted log NT-proBNP levels in patients with Mets and without Mets

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

have also been detected to be significantly lower in MetS [6]. Consistent with the above-mentioned finding, differences of NT-proBNP plasma concentrations were established between MetS and non-MetS in our study by covariance analysis after adjustment for age, gender (for total patients), serum creatinine, LAD, AOD, LVMI, and LVEF. On the other hand, to the best of our knowledge, ours is the first investigation to demonstrate that plasma NT-proBNP is lower in patients with MetS defined by the Chinese Diabetes Society of the Chinese Medical Association.

NT-proBNP, body mass index and blood lipid

Consistent with prior experimental observations [12], we found that there is a strong relationship between serum NT-proBNP and plasma triglycerides, high density lipoprotein cholesterol, and low density lipoprotein cholesterol levels, suggesting a strong relationship between NT-proBNP and lipid metabolism. An inverse relationship between serum NT-proBNP and cholesterol has been described previously and has been hypothesized to be a potential link impairing the natural blood pressure regulation [6]. Our data support to some degree this hypothesis because of the inverse relationship between low density lipoprotein cholesterol and NT-proBNP. Log NT-proBNP levels were also found to be significantly lower in patients of MetS subgroups stratified by plasma triglycerides and high density lipoprotein cholesterol compared to those without MetS, suggesting a negative association between lower NT-proBNP concentration and hyperlipidaemia.

We also found that there is a strong relationship between serum NT-proBNP and body mass index. An inverse relationship between serum BNP and body mass index has been described recently [13],

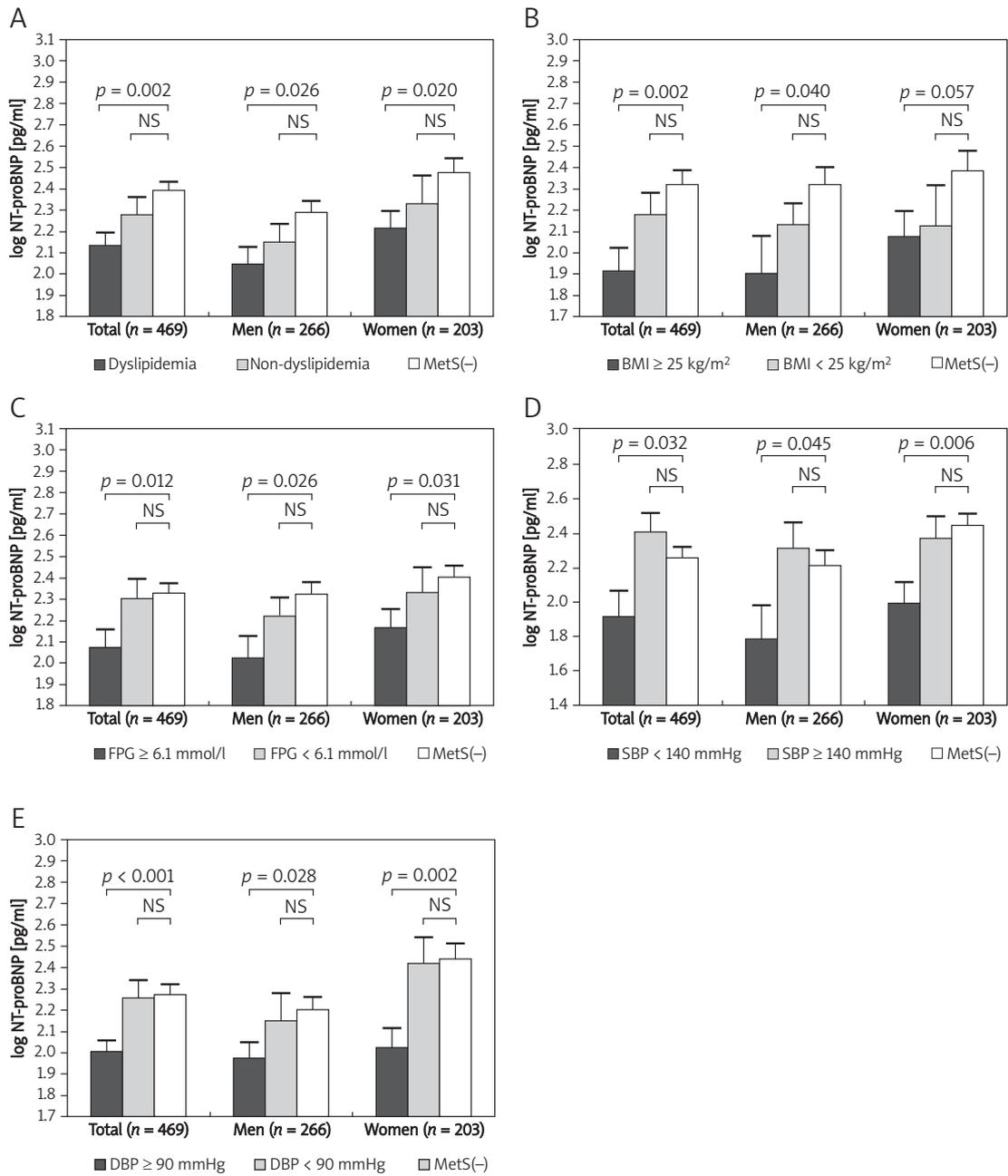


Figure 2. Adjusted log NT-proBNP levels in subgroups of MetS and patients without MetS
 MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Levels were obtained by log transformation of NT-proBNP levels. Covariance analysis between groups was carried out and adjusted for age, serum creatinine, left atrial diameter, left ventricular mass, left ventricular ejection fraction and the other MetS covariates. Data are presented as mean ± SD; NS – no significant

Table III. Adjusted log NT-proBNP levels in patients with MetS and without MetS

	Total		Men		Women	
	n	Log NT-proBNP	n	Log NT-proBNP	n	Log NT-proBNP
MetS(+)	230	2.173 ± 0.044	124	2.110 ± 0.056	106	2.022 ± 0.074
MetS(-)	239	2.367 ± 0.035	142	2.261 ± 0.045	97	2.232 ± 0.055
Value of p		0.001		0.038		0.022

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

Table IV. Adjusted log NT-proBNP levels in blood lipid subgroups of MetS and patients without MetS

	Total		Men		Women	
	n	Log NT-proBNP	n	Log NT-proBNP	n	Log NT-proBNP
MetS(-)	239	2.391 ±0.044	142	2.289 ±0.056	97	2.476 ±0.063
MetS(+)	230		124		106	
Non-dyslipidaemia	110	2.275 ±0.084 (p = 0.243)	51	2.145 ±0.093 (p = 0.213)	59	2.329 ±0.135 (p = 0.342)
Dyslipidaemia	120	2.133 ±0.059 (p = 0.002)	73	2.044 ±0.082 (p = 0.026)	47	2.213 ±0.084 (p = 0.020)

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

Table V. Adjusted log NT-proBNP levels in BMI subgroups of MetS and patients without MetS

	Total		Men		Women	
	n	Log NT-proBNP	n	Log NT-proBNP	n	Log NT-proBNP
MetS(-)	239	2.317 ±0.068	142	2.317 ±0.087	97	2.388 ±0.090
MetS(+)	230		124		106	
BMI < 25 kg/m ²	89	2.182 ±0.099 (p = 0.257)	46	2.130 ±0.100 (p = 0.147)	43	2.126 ±0.190 (p = 0.214)
BMI ≥ 25 kg/m ²	141	1.916 ±0.108 (p = 0.002)	78	1.904 ±0.175 (p = 0.040)	63	2.074 ±0.127 (p = 0.057)

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

Table VI. Adjusted log NT-proBNP levels in FPG subgroups of MetS and patients without MetS

	Total		Men		Women	
	n	Log NT-proBNP	n	Log NT-proBNP	n	Log NT-proBNP
MetS(-)	239	2.325 ±0.046	142	2.323 ±0.055	97	2.399 ±0.054
MetS(+)	230		124		106	
FPG < 6.1 mmol/l	105	2.296 ±0.100 (p = 0.795)	55	2.218 ±0.088 (p = 0.328)	50	2.328 ±0.119 (p = 0.602)
FPG ≥ 6.1 mmol/l	125	2.068 ±0.089 (p = 0.012)	69	2.021 ±0.104 (p = 0.026)	56	2.165 ±0.085 (p = 0.031)

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

Table VII. Adjusted log NT-proBNP levels in SBP subgroups of MetS and patients without MetS

	Total		Men		Women	
	n	Log NT-proBNP	n	Log NT-proBNP	n	Log NT-proBNP
MetS(-)	239	2.258 ±0.071	142	2.214 ±0.088	97	2.445 ±0.077
MetS(+)	230		124		106	
SBP < 140 mmHg	109	1.919 ±0.147 (p = 0.032)	56	1.795 ±0.188 (p = 0.045)	53	1.991 ±0.124 (p = 0.006)
SBP ≥ 140 mmHg	121	2.416 ±0.106 (p = 0.212)	68	2.316 ±0.150 (p = 0.560)	53	2.375 ±0.120 (p = 0.643)

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

Table VIII. Adjusted log NT-proBNP levels in DBP subgroups of MetS and patients without MetS

	Total		Men		Women	
	n	Log NT-proBNP	n	Log NT-proBNP	n	Log NT-proBNP
MetS(-)	239	2.273 ±0.045	142	2.200 ±0.058	97	2.439 ±0.073
MetS(+)	230		124		106	
DBP < 90 mmHg	107	2.250 ±0.091 (p = 0.824)	59	2.146 ±0.132 (p = 0.715)	48	2.420 ±0.123 (p = 0.906)
DBP ≥ 90 mmHg	123	2.001 ±0.059 (p < 0.001)	65	1.974 ±0.078 (p = 0.028)	58	2.024 ±0.092 (p = 0.002)

MetS – metabolic syndrome, NT-proBNP – N-terminal pro-B-type natriuretic peptide. Covariance analysis between groups was carried out. Levels were obtained by log transformation of NT-proBNP levels adjusted for age, gender, serum creatinine, LAD, AOD, LVMI, and LVEF. Data are presented as mean ± SD

which is also supported to some degree by our data. This is also supported by previous observations that obese subjects are more sensitive to sodium load on account of a reduced effect of atrial natriuretic peptide, partly attributable to increased removal of natriuretic peptide from the circulation by adipocytes through abundant natriuretic peptide clearance receptors in adipose tissue of obese people [14]. NT-proBNP may also relate to this, but the importance of the natriuretic clearance receptor for clearance of NT-proBNP is unknown [15]. Alternatively, NT-proBNP may, through lipolytic and lipomobilizing effects, change the metabolic state as demonstrated for atrial natriuretic peptide [16] and reduce the incidence of overweight and obesity. An association between natriuretic peptides and levels of the insulin-sensitizing hormone adiponectin has been reported, with low levels of BNP associated with low levels of adiponectin despite adjustment for body mass index [17]. Natriuretic peptides also stimulate lipolysis and release of triacylglycerols from adipose tissue [18]. Reduced natriuretic peptide signalling could have detrimental effects via the promotion of lipid accumulation in adipose tissue and skeletal muscle.

NT-proBNP and blood glucose

An inverse relationship between plasma NT-proBNP levels and fasting blood glucose was found in our study. Log NT-proBNP levels were also found to be significantly lower in patients of MetS subgroups stratified by fasting blood glucose compared to non-MetS subjects. Some experimental observations suggest that low natriuretic peptide levels could predispose to insulin resistance. Reduced natriuretic peptide activity leads to greater activation of the renin-angiotensin system [19]. Activation of the renin-angiotensin system promotes the development of insulin resistance via multiple mechanisms [20], including inhibition of intracellular insulin signalling, enhanced oxidative stress, inflammation, reduced adipocyte differentiation, and decreased perfusion to the skeletal muscle and pancreas.

Natriuretic peptides and blood pressure

Previous studies have reported a positive association between plasma natriuretic peptide levels and systolic blood pressure [21]. In conformity with these data, we also found that low NT-proBNP levels were associated with almost all component of the metabolic syndrome except elevated systolic blood pressure, which was associated with higher plasma NT-proBNP levels, indicating the haemodynamic influence of blood pressure on natriuretic peptide synthesis. The observation that plasma NT-proBNP levels relate differently to systolic blood pressure than to other metabolic components suggests that blood pressure segregates separately from other MetS components [22]. In the Insulin Resistance Atherosclerosis Study, Hanley *et al.* [22] used principal factor analysis to identify 2 "factors". One is a metabolic factor, consisting of body mass index, glycaemic measures, triglycerides, and high density lipoprotein cholesterol; the other is a blood pressure factor, comprising systolic and diastolic blood pressure. HOMA-IR and fasting insulin correlated with the metabolic factor but not with the blood pressure factor. These data suggest that blood pressure may have a different pathophysiological mechanism from other metabolic components, which may help to interpret the different association between systolic blood pressure and NT-proBNP from others. There is an inverse correlation between plasma NT-proBNP levels and diastolic blood pressure, which may reflect the influence of pulse pressure [6]. Increased pulse pressure is often accompanied by lower diastolic blood pressure, particularly in older individuals.

NT-proBNP and other risk factors

The association between left ventricular hypertrophy and plasma levels of NT-proBNP is consistent. Some studies have found an association between NT-proBNP levels and left ventricular hypertrophy, whereas others have not [8]. In the present study we found there was an independent

association between left ventricular mass index and NT-proBNP levels in subjects.

Age has been shown to influence circulating natriuretic peptide levels, and the magnitude of the effects and potential importance in the interpretation of NT-proBNP remain unclear [23]. A possible explanation for increased NT-proBNP levels with age may be increased age-related fibrosis, diastolic dysfunction and reduced renal clearance [24]. In this study we also found that NT-proBNP levels were positively related to age in all subjects.

Comparison with previous studies

An inverse association between plasma natriuretic peptide levels and obesity has previously been described in the Framingham [2] and other studies [13, 25]. However, few studies have investigated the relations between NT-proBNP levels and metabolic components, especially in MetS subjects diagnosed according to recommendations of the Chinese Diabetes Society. Inconsistent associations have been reported between plasma natriuretic peptides and glycaemic status [5] or plasma lipids [4]. In the population-based Danish cohort of Olsen *et al.* [6], plasma NT-proBNP levels were inversely associated with serum total cholesterol, serum triglycerides and plasma glucose. The present study differs from those findings by the use of covariance analyses and the ability to adjust for differences in age, gender, renal function with serum creatinine, LAD, AOD, LVMI, and LVEF. In addition, as gender is an independent factor influencing NT-proBNP levels, we divided all subjects into male and female groups. For men, fasting blood glucose was not an independent predictor of NT-proBNP, while for women NT-proBNP levels were not independently correlated with blood lipid. Notwithstanding the above, the similarity in study findings provides convincing evidence of an association between the natriuretic peptide and metabolic components.

Several limitations of the study deserve comment. First, the study results indicated that NT-proBNP is different in MetS subjects from non-MetS. The fact that NT-proBNP was related to metabolic components independently of age and clinical and echocardiographic characteristics indicates that NT-proBNP is a superior marker of metabolic syndrome and therefore suitable to detect subclinical MetS before it develops into clinical disease. However, the positive or negative relationship between NT-proBNP and metabolic components may be a problem when defining cut-off values for varying levels of obesity, dyslipidaemia, or hyperglycaemia. Here subgroups are established according to the Chinese criteria. Next, further studies are required to investigate

NT-proBNP levels in MetS subgroups by other traditional cut-off levels in patients. Second, the multiple regression models explained only a moderate influence of potential risk factors on NT-proBNP levels. Other environmental factors may contribute to the unexplained variation. Finally, it is important to mention that our study concerned Chinese individuals and our findings may not be relevant to those of other countries.

The diagnostic value of plasma natriuretic peptide measurements has been demonstrated in a variety of settings. The range of factors that may influence the biomarker needs to be understood for interpretation of the results. Previous studies have shown the use of lower natriuretic peptide cut points for diagnosing heart failure in obese individuals [25]. Our data raise the possibility that other metabolic components should be taken into account when plasma natriuretic peptide (NT-proBNP) measurements are interpreted. But whether diagnostic thresholds should be altered in individuals with the metabolic syndrome is unknown and needs further investigation in prospective studies. In addition to that, because natriuretic peptides play an important role in the counter-regulatory response to volume and pressure overload, it is possible that lower natriuretic peptide levels contribute to the susceptibility of individuals with the metabolic syndrome to hypertension and left ventricular hypertrophy.

In conclusion, plasma NT-proBNP levels were lower in patients with MetS, especially in MetS patients with dyslipidaemia, elevated body mass index, fasting blood glucose and diastolic blood pressure. Our data raise the possibility that reduced plasma NT-proBNP levels may be a marker for MetS, which might possess significant clinical and pathophysiological implications. More prospective studies are needed regarding the relations of lower NT-proBNP levels and sodium, insulin resistance, and the impact of reduced natriuretic peptide levels on cardiovascular events in MetS.

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