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Serum Levels of Meteorin-Like (Metrnl) Are Increased in Patients with Newly Diagnosed Type 2 Diabetes Mellitus and Are Associated with Insulin Resistance

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Background: Meteorin-like (Metrnl) is a novel adipomyokine that may improve glucose tolerance and affect insulin resistance. This study aimed to investigate the association between serum levels of Metrnl with blood glucose status and to its association with insulin resistance.

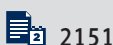
Material/Methods: The study included 160 subjects with normal glucose tolerance (NGT) (n=40), impaired fasting glucose (IFG) (n=40), impaired glucose tolerance (IGT) (n=40), and newly diagnosed type 2 diabetes mellitus (T2DM) (n=40). An enzyme-linked immunosorbent assay (ELISA) was used to measure the serum levels of Metrnl. Partial correlation analysis was used to analyze the relationship between serum levels of Metrnl and metabolic parameters. Multiple logistic regression analysis was performed to identify the association between serum levels of Metrnl with the risk of diabetes.

Results: Serum levels of Metrnl was highest in patients with T2DM and significantly increased in patients with prediabetes compared with individuals with NGT. After adjusting for age, gender, and body mass index (BMI), serum Metrnl level was significantly correlated with lipid profile, glucose profile, and insulin resistance. Multiple logistic regression analysis showed that Metrnl significantly increased the risk of T2DM (OR=1.727; P=0.008) before adjusting for the homeostatic model assessment of insulin resistance (HOMA-IR). When further adjusted for HOMA-IR, Metrnl was no longer associated with an increased OR for T2DM (OR=1.491; P=0.066), while the HOMA-IR significantly increased the risk of T2DM (OR=1.935; P=0.008).

Conclusions: Serum levels of Metrnl were significantly increased in patients with T2DM and may increase the risk of T2DM independent of insulin resistance.

MeSH Keywords: **Cytokines • Diabetes Mellitus, Type 2 • Insulin Resistance**

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Background

Insulin resistance is a risk factor for the development of type 2 diabetes mellitus (T2DM). Improving insulin resistance continues to be an important area of research into the treatment of T2DM and metabolic syndrome, as new insulin-sensitizing treatments are needed [1,2]. Secreted proteins, including adipokines and myokines, have been shown to be closely associated with the development of insulin resistance and have the potential for drug development in the treatment of obesity and T2DM [3–10].

Meteorin-like (Metrnl), a secreted adipomyokine, was recently identified and shown to be induced by exercise or cold exposure [5,11,12]. Metrnl increases energy expenditure and improves insulin sensitivity by inducing the expression of genes associated with brown fat thermogenesis in mice [5]. This finding raises the possibility that Metrnl has a role in the pathophysiological processes of metabolic diseases, including T2DM. The findings of previously published studies have shown that serum levels of Metrnl are increased in patients with T2DM, but the findings are controversial. Lee et al. [13] and Zheng et al. [14] reported lower serum levels of Metrnl in patients with newly diagnosed T2DM, while Chung et al. [15] identified increased serum levels of Metrnl. Also, in preclinical studies, the effects of Metrnl on adipose tissue in mice showed that expression of Metrnl by adipocytes could increase insulin sensitivity by stimulating the peroxisome proliferator-activated receptor gamma (PPAR γ) pathway [16]. The overexpression of Metrnl has been shown to reduce lipogenesis and inhibit the expression of PPAR γ in human adipocytes, which may lead to hyperinsulinemia and insulin resistance [17].

Given the possible therapeutic role of Metrnl in patients with T2DM, the aim of this study was to investigate the association between serum levels of Metrnl with blood glucose status and to investigate its association with insulin resistance. Serum levels of Metrnl were measured in patients with newly diagnosed T2DM and compared with patients with normal glucose tolerance (NGT), impaired fasting glucose (IFG), and impaired glucose tolerance (IGT).

Material and Methods

Patients and control subjects

This observational study included 160 subjects. Four groups were studied that included subjects with normal glucose tolerance (NGT) or healthy volunteers (n=40), a group with impaired fasting glucose (IFG) (n=40), a group with impaired glucose tolerance (IGT) (n=40), and a group with newly diagnosed type 2 diabetes mellitus (T2DM) (n=40). The study was conducted at

the Qilu Hospital of Shandong University from November 2017 to May 2018 and was approved by the Ethics Committee of Shandong University, Qilu Hospital (Approval No. 2017-014).

Patients with previously diagnosed T2DM, other types of diabetes, severe cardiovascular or cerebrovascular diseases, severe liver or renal disease, and medication history including the use of antidiabetics, statins, diuretics, corticosteroids, estrogen, and progestin, which influence glucose tolerance and insulin sensitivity, were excluded. T2DM was diagnosed as a fasting blood glucose (FBG) ≥ 7.0 mmol/L and/or 2-hour postprandial blood glucose (PBG) ≥ 11.1 mmol/L. IFG was diagnosed as an FBG of 6.1–7.0 mmol/L and a PBG < 7.8 mmol/L. IGT was diagnosed as an FBG < 6.1 mmol/L and a PBG of 7.8–11.1 mmol/L. These diagnostic criteria were in accordance with the 2006 World Health Organization (WHO) criteria [18]. Overweight and obesity was diagnosed by a body mass index (BMI) ≥ 25.0 kg/m² based on the criteria recommended by the WHO [19].

Clinical data

Detailed clinical and demographic data, including past medical history of the study participants in the four study groups, were collected by trained clinical investigators. Height and weight measurements were used to calculate BMI. The waist circumference (WC) was measured from the midpoint between the lower border of the ribcage and the anterior superior iliac spine. After sitting for at least 5 min, blood pressure (BP) was measured consecutively three times on the left arm using an Omron Model HEM-752 sphygmomanometer (Omron Company, Dalian, China), and the average value was recorded.

Laboratory tests

After a 10-hour overnight fast, a sample of fasting venous blood was collected at 7:00 am and the serum was used for the measurement of fasting blood glucose (FBG), fasting insulin, total cholesterol (TC), triglyceride (TG), low-density lipoprotein cholesterol (LDL-C), high-density lipoprotein cholesterol (HDL-C), alanine aminotransferase (ALT), aspartate aminotransferase (AST), uric acid (UA), and creatinine by using an Architect ci16200 integrated automatic analyzer (Abbott Labs, Abbott Park, Ill, USA). Glycated hemoglobin (HbA1c) was measured by high-performance liquid chromatography (HPLC) using a VARIANT™ II TURBO (Bio-Rad, Hercules, CA, USA). PBG was measured after all subjects had completed an oral glucose tolerance test (OGTT) of 75 g of glucose.

Measurement of serum Metrnl levels

Serum Metrnl levels were measured using enzyme-linked immunosorbent assay (ELISA) kits (Code: CSB-EL013718HU) (CusaBio Biotech Co., Ltd., Wuhan, China). The homeostatic

Table 1. Characteristics of the subjects in the four study groups with normal glucose tolerance (NGT), impaired fasting glucose (IFG), impaired glucose tolerance (IGT), and type 2 diabetes mellitus (T2DM).

Characteristic	NGT (n=40)	IFG (n=40)	IGT (n=40)	T2DM (n=40)	P-value ^a
Female (%)	18 (45.0)	16 (40.0)	20 (50.0)	16 (40.0)	0.773
Age (years)	50.68±5.40	51.50±5.27	50.65±6.27	52.13±7.91	0.676
BMI (kg/m ²)	24.73±2.57	26.02±3.09	26.35±3.38	27.58±3.10 ^b	0.001
WC (cm)	84.28±8.45	88.52±8.99	88.41±9.37	92.11±7.38 ^b	0.001
Systolic BP (mmHg)	128.35±14.99	132.33±12.00	132.28±13.13	141.20±14.00 ^{bcd}	<0.001
Diastolic BP (mmHg)	78.13±8.63	82.55±7.34	82.55±8.12	85.68±6.44 ^b	<0.001
FBG (mmol/L)	5.23±0.30	6.38±0.20 ^b	5.53±0.32 ^c	8.44±2.44 ^{bcd}	<0.001
PBG (mmol/L)	5.80±0.91	6.90±0.48	8.99±0.83 ^{bc}	14.85±3.92 ^{bcd}	<0.001
HbA1c (%)	4.89±0.35	5.35±0.34	5.44±0.39 ^b	7.08±1.55 ^{bcd}	<0.001
Fasting insulin (μU/mL)	4.55 (3.32–6.05)	8.05 (5.98–9.83) ^b	8.35 (5.30–10.85) ^b	7.75 (4.69–10.10) ^b	<0.001
HOMA-IR (index)	1.08 (0.80–1.41)	2.31 (1.63–2.78) ^b	2.13 (1.33–2.65) ^b	2.59 (1.66–3.64) ^{bcd}	<0.001
TC (mmol/L)	4.64±0.55	4.99±0.72	5.02±1.09	5.05±0.80	0.080
TG (mmol/L)	0.94 (0.71–1.07)	1.25 (0.96–2.00) ^b	1.35 (0.97–1.98) ^b	1.63 (1.10–2.17) ^{bc}	<0.001
LDL-C (mmol/L)	2.90±0.61	3.22±0.64	3.18±0.77	3.33±0.64 ^b	0.032
HDL-C (mmol/L)	1.43±0.27	1.39±0.43	1.40±0.44	1.31±0.29	0.489
ALT (U/L)	18.04±4.50	17.30±5.24	17.89±5.57	23.31±8.50 ^{bcd}	<0.001
AST (U/L)	19.91±3.83	18.80±4.34	19.05±4.37	22.33±7.45 ^{bcd}	0.011
Creatinine (μmol/L)	61.92±12.81	62.56±9.23	62.00±10.24	63.32±10.26	0.943
UA (umol/L)	269.20±76.06	273.12±51.69	260.35±63.87	273.75±75.65	0.801
MetrnI (pg/mL)	126.86 (103.44–193.94)	176.76 (135.89–246.20) ^b	192.69 (116.84–311.12) ^b	256.79 (193.97–384.53) ^{bcd}	<0.001

Data are presented as the mean ± standard deviation (SD) or the median (interquartile range). NGT – normal glucose tolerance; IFG – impaired fasting glucose; IGT – impaired glucose tolerance; DM – diabetes mellitus; BMI – body mass index; WC – waist circumference; BP – blood pressure; FBG – fasting blood glucose; PBG – postprandial blood glucose; HOMA-IR – homeostatic model assessment of insulin resistance; TC – total cholesterol; TG – triglyceride; LDL-C – low-density lipoprotein cholesterol; HDL-C – high-density lipoprotein cholesterol; ALT – alanine aminotransferase; AST – aspartate aminotransferase; UA – uric acid. ^a Difference between the four groups; ^b P<0.05 compared with the NGT group; ^c P<0.05 compared with the IFG group; ^d P<0.05 compared with IGT group.

model assessment of insulin resistance (HOMA-IR) index was calculated as the fasting insulin concentration (mIU/L)×FBG concentration (mmol/L)/22.5. The estimated glomerular filtration rate (eGFR), based on the serum creatinine level was calculated using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) equation [20].

Statistical analysis

Statistical analysis was performed using SPSS version 22.0 software (SPSS Inc., Chicago, Ill, USA). The continuous variables with a normal distribution were expressed as the mean ± standard

deviation (SD), while those with a non-normal distribution were expressed as the median and interquartile range (IQR). The Kolmogorov-Smirnov test was performed to analyze the normality of the data. Between-group differences were analyzed using one-way analysis of variance (ANOVA) using Tukey's honestly significant difference (HSD) test or the chi-squared (χ^2) test. Partial correlation analysis was used to evaluate the relationship between serum levels of MetrnI and metabolic parameters. Multiple logistic regression analysis was used to determine the association of serum MetrnI levels and the risk of T2DM. A P-value <0.05 was considered to be statistically significant.

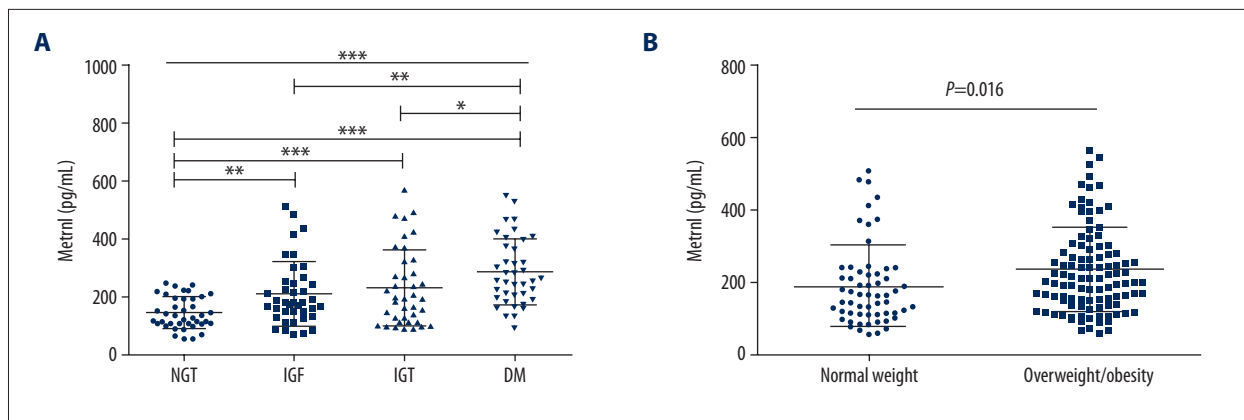


Figure 1. Comparison of serum meteorin-like (Metrn) levels according to blood glucose status and weight in subjects in the four study groups. **(A)** Serum meteorin-like (Metrn) levels in subjects with normal glucose tolerance (NGT), impaired fasting glucose (IFG), impaired glucose tolerance (IGT), and newly diagnosed type 2 diabetes mellitus (T2DM). **(B)** Serum Metrn levels according to weight status (normal weight vs. overweight and obese). * $P<0.05$; ** $P<0.01$; *** $P<0.001$. Data are shown as the mean \pm standard deviation (SD).

Results

Clinical and laboratory characteristics of the study participants

Table 1 shows the comparison of the demographic, clinical, and laboratory data between the four study groups, subjects with normal glucose tolerance (NGT) or healthy volunteers, the group with impaired fasting glucose (IFG), the group with impaired glucose tolerance (IGT), and the group with newly diagnosed type 2 diabetes mellitus (T2DM). Compared with the other groups, the group with T2DM had significantly higher systolic blood pressure (BP), glycated hemoglobin (HbA1c), fasting blood glucose (FBG), postprandial blood glucose (PBG), homeostatic model assessment of insulin resistance (HOMA-IR) index, alanine aminotransferase (ALT), and aspartate aminotransferase (AST). Serum levels of Metrn were also highest in patients with T2DM (Figure 1A), and the serum levels of Metrn level were higher in the prediabetes groups than in the NGT subjects, although there was no significant difference between the IFG and IGT groups. Serum levels of Metrn level between subjects with normal weight were significantly different when compared with levels in overweight and obese subjects (Figure 1B).

Serum levels of Metrn and metabolic parameters

Partial correlation analysis was used to investigate the relationship between serum levels of Metrn and metabolic parameters (Table 2). Before adjusting for age, gender, and body mass index (BMI), serum Metrn levels were positively correlated with BMI, waist circumference (WC), total cholesterol (TC), triglyceride (TG), low-density lipoprotein cholesterol (LDL-C), FBG, PBG, HbA1c, fasting insulin, and HOMA-IR and was negatively correlated with high-density lipoprotein cholesterol (HDL-C) in all

the subjects studied. After adjusting for age, gender, and BMI, these correlations remained statistically significant except for WC, systolic BP, TC, and LDL-C.

Serum levels of Metrn and T2DM

The relationship between Metrn and T2DM was explored by multiple logistic regression analysis (Table 3). Before adjusting for confounders (model 1), serum Metrn levels were associated with significantly increased odds ratios (ORs) for T2DM (OR=1.892; $P<0.001$). After adjusting for age, gender, and BMI (model 2), the association between increased serum Metrn levels and an increased risk for T2DM was also found (OR=1.780, $P=0.001$). When further adjusted for systolic BP, TC, TG, ALT (model 3), the estimated glomerular filtration rate (eGFR), and uric acid (UA) and increased serum levels of Metrn remained significantly correlated with an increased the risk of T2DM (OR=1.727; $P=0.008$). However, when further statistical adjustment was made for HOMA-IR (model 4), serum levels of Metrn were no longer significantly correlated with an increased OR for T2DM (OR=1.491; $P=0.066$), while the HOMA-IR significantly increased the risk of T2DM (OR=1.935; $P=0.008$).

Discussion

Meteorin-like (Metrn) is a novel adipomyokine that shows high expression levels in skeletal muscle and white adipose tissue [5,21]. In mice, Metrn can promote the activation of adipose tissue macrophages and increases the expression of anti-inflammatory and thermogenic genes in white adipose tissue [5]. Therefore, Metrn might have a potential therapeutic role in inflammatory and metabolic diseases by regulating tissue energy homeostasis [5,22–25].

Table 2. Correlation analysis of serum levels of meteorin-like (Metrnl), and metabolic parameters.

Variable	Metrnl		Metrnl (age, gender, BMI adjusted)	
	Coefficient (r)	P-value	Coefficient (r)	P-value
Age (years)	-0.022	0.784		
BMI (kg/m ²)	0.283	<0.001		
WC (cm)	0.194	0.014	0.014	0.862
Systolic BP (mmHg)	0.086	0.278	0.090	0.263
Diastolic BP (mmHg)	0.087	0.274	0.157	0.049
TC (mmol/L)	0.157	0.048	0.154	0.054
TG (mmol/L)	0.503	<0.001	0.457	<0.001
LDL-C (mmol/L)	0.200	0.011	0.142	0.076
HDL-C (mmol/L)	-0.394	<0.001	-0.221	0.005
FBG (mmol/L)	0.379	<0.001	0.222	0.005
PBG (mmol/L)	0.420	<0.001	0.313	<0.001
HbA1c (%)	0.379	<0.001	0.273	0.001
Fasting insulin (μU/mL)	0.338	<0.001	0.231	0.004
HOMA-IR (index)	0.433	<0.001	0.267	0.001

BMI – body mass index; WC – waist circumference; BP – blood pressure; TC – total cholesterol; TG – triglyceride; LDL-C – low-density lipoprotein cholesterol; HDL-C – high-density lipoprotein cholesterol; FBG – fasting blood glucose; PBG – postprandial blood glucose; HOMA-IR – homeostatic model assessment of insulin resistance.

Table 3. Multiple logistic regression analysis of the association between serum levels of meteorin-like (Metrnl) with type 2 diabetes mellitus (T2DM).

Model	Independent variable	beta coefficient	Standard error of beta coefficient	Wald statistic	Odds ratio (95% CI)	P-value
Model 1	Metrnl, per 100 pg/mL	0.638	0.161	15.757	1.892 (1.381–2.593)	<0.001
Model 2	Metrnl, per 100 pg/mL	0.577	0.171	11.345	1.780 (1.273–2.491)	0.001
Model 3	Metrnl, per 100 pg/mL	0.546	0.207	6.931	1.727 (1.150–2.593)	0.008
Model 4	Metrnl, per 100 pg/mL	0.399	0.217	3.387	1.491 (0.974–2.281)	0.066
	HOMA-IR, per 1 unit	0.660	0.248	7.080	1.935 (1.190–3.146)	0.008

Model 1: not adjusted; Model 2: adjusted for age, gender, and body mass index (BMI); Model 3: Model 2 plus systolic blood pressure (BP), total cholesterol (TC), triglyceride (TG), alanine aminotransferase (ALT), estimated glomerular filtration rate (eGFR), and uric acid (UA); Model 4: Model 3 plus the homeostatic model assessment of insulin resistance (HOMA-IR).

Although there have been several reported studies on the association between serum levels of Metrnl in type 2 diabetes mellitus (T2DM), the findings have been conflicting. Lee et al. [13] and Zheng et al. [14] reported lower serum levels of Metrnl in patients with newly diagnosed T2DM, whereas Chung et al. [15] reported increased serum levels of Metrnl, which was consistent with the findings of the present study. Because of the limited number of studies on the association between T2DM and serum levels of Metrnl, it is difficult to explain these conflicting findings. However, preclinical studies in mice have shown that

Metrnl in adipocytes could increase insulin sensitivity by stimulating the peroxisome proliferator-activated receptor gamma (PPAR γ) pathway [16], which has a regulatory role in insulin resistance and adipocyte differentiation [26]. A study conducted in human adipocytes showed that overexpression of Metrnl inhibited the expression of PPAR γ , which might lead to hyperinsulinemia and insulin resistance [17]. Therefore, increased serum levels of Metrnl might increase the risk of T2DM by promoting insulin resistance, which is consistent with the findings of the present study.

The findings of this study showed that increased serum levels of Metrnl were significantly associated with T2DM (OR=1.727; P=0.008), in model 3 that used multiple logistic regression analysis after adjusting for confounding factors, except for the homeostatic model assessment of insulin resistance (HOMA-IR). However, when further adjusted for HOMA-IR, in model 4, serum levels of Metrnl were no longer associated with an increased OR for T2DM (OR=1.491; P=0.066). Another explanation for the increased serum levels of Metrnl in patients with T2DM may be attributed to a protective compensatory response against metabolic overload, including insulin resistance [15]. On the basis of these results, the effect of Metrnl on insulin resistance should be confirmed by further studies as at this time, studies on serum Metrnl and its association with T2DM are limited and the findings have been contradictory. To our knowledge, the present study was the first to identify a significant correlation between serum levels of Metrnl with insulin resistance and to report that Metrnl might contribute to the development of T2DM dependent of insulin resistance in human subjects.

Because Metrnl is highly expressed in white adipose tissue, serum levels are likely to be correlated with parameters that include body mass index (BMI) and waist circumference (WC). The findings of this study showed that serum Metrnl levels were increased in overweight and obese subjects when compared with subjects of normal weight (Figure 1B) and was significantly correlated with BMI and WC. Löffler et al. also reported increased Metrnl expression in adipocytes in obese children compared with lean children [17]. However, Chung et al. [15] found that serum Metrnl levels were not associated with body weight, BMI, WC, and mass of adipose tissue. A preclinical study showed that when compared with control mice, neither Metrnl adipose-specific knockout nor transgenic-overexpression mice

exhibited a change in body weight, lean to fat body mass, or distribution of adipose tissue [16]. It is clear that the relationship between serum levels of Metrnl and obesity requires further clinical study.

This study also had several limitations. This was an observational study that did not investigate the causal relationship between serum levels of Metrnl and the development of T2DM. Also, this study included only Chinese subjects, and the findings require validation in subjects of different ethnicity. In this study, it was not possible to exclude other potential confounding factors, especially exercise, as Metrnl expression may be induced in muscle after exercise. Also, as Metrnl is mainly produced by muscle and adipose tissue, the percentage of body fat and skeletal muscle mass should also be assessed in future studies, and statistical analysis of patient data should also be adjusted for these parameters. Finally, the sample size of this study was quite small, and further studies with larger numbers of subjects should be undertaken in the future.

Conclusions

The findings from an observational study conducted at a single center in China showed that serum levels of meteorin-like (Metrnl) were significantly increased in patients with newly diagnosed type 2 diabetes mellitus (T2DM) and were significantly correlated with the lipid profile, glucose profile, and insulin resistance. Further logistic regression analysis showed that serum Metrnl levels were significantly correlated with an increased risk of T2DM, and this association might be due to insulin resistance. Future mechanism research studies are required to explore the mechanism for Metrnl in insulin resistance and T2DM.

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