

Association of retinol binding protein-4, cystatin C, homocysteine and high-sensitivity C-reactive protein levels in patients with newly diagnosed type 2 diabetes mellitus

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Abstract

Introduction: To investigate the serum retinol binding protein (RBP)-4, cystatin C (Cys C), homocysteine (HCY) and high-sensitivity C-reactive protein (hs-CRP) levels in newly diagnosed type 2 diabetes mellitus (NT2DM) patients, prediabetes mellitus (PDM) subjects and normal controls, as well as their correlation with clinical and laboratory indexes, such as blood pressure and lipoprotein.

Material and methods: A total of 242 subjects, including 141 NT2DM patients, 48 PDM subjects and 53 healthy controls, were recruited in the present study. Serum RBP-4, Cys C and hs-CRP concentrations were measured by enzyme-linked immunosorbent assay (ELISA). HCY concentration was determined by the chemical luminescence method.

Results: There were significant differences in Cys C and hs-CRP among NT2DM patients, PDM subjects and normal controls. In comparison to controls, there were significantly elevated Cys C and hs-CRP levels in PDM (both $p < 0.001$), and a significantly increased Cys C level in NT2DM ($p < 0.001$); however, there were no significant differences in Cys C and hs-CRP levels between NT2DM and PDM, and no significant differences of hs-CRP levels between NT2DM and normal controls. No significant differences of RBP-4 and HCY levels among NT2DM, PDM and normal control groups were observed.

Conclusions: Aberrant Cys C expression and its clinical associations in NT2DM suggest their important role in this disease.

Key words: retinol binding protein-4, cystatin C, homocysteine, high-sensitivity C-reactive protein, type 2 diabetes mellitus.

Introduction

Type 2 diabetes mellitus (T2DM) is one of the most common chronic metabolic diseases, featuring hyperglycemia resulting from resistance to insulin action and an inadequate compensatory insulin secretory response [1]. The chronic hyperglycemia of diabetes is associated with long-term damage, dysfunction and failure of various organs, of which the long-term complications, such as retinopathy, nephropathy, cardiovascular (CV) and cerebrovascular diseases, are considered to be the

dominant factors affecting the life span and the quality of life in T2DM [2–5].

Cystatin C (Cys C), a cysteine proteinase inhibitor, is a non-glycosylated single chain protein with a molecular weight of 1.3 kDa. It is secreted shortly after its synthesis and steadily produced by all nucleated cells [6, 7]. Investigations have demonstrated that serum Cys C improves estimates of glomerular filtration rate (GFR) more than serum creatinine (Cr)-based methods alone, and was associated with the presence of CV disease in subjects with mild renal impairment [8–10]. It has been reported that Cys C levels increase and cys-GFR levels decrease with increasing severity of glucose intolerance [11]. Moreover, in patients with diabetes, several studies have demonstrated that serum Cys C had better performance compared with serum Cr in evaluation of renal function, suggesting that it is a sensitive serum marker for the early assessment of kidney disease with diabetes [11, 12].

Retinol binding protein (RBP)-4, as a 21 kDa protein, was initially known as a specific carrier for the delivery of retinol in circulation [13]. In the liver and adipose tissue, there was overexpression of RBP-4. Insulin resistance is a conspicuous characteristic of prediabetes mellitus (PDM) states (impaired fasting glucose (IFG) and/or impaired glucose tolerance (IGT)), and plays a crucial role in the development of T2DM. Increased serum RBP-4 levels have been reported in connection with IGT [14]. The relationship between RBP-4 and diabetes has been revealed in some studies; serum RBP-4 concentrations were negatively correlated with insulin resistance and development of T2DM [15–17]. However, the expression of RBP-4 in newly diagnosed T2DM and PDM and its association with insulin secretion and pancreas beta function are still unknown.

Homocysteine (HCY) is a sulfur-containing amino acid derived from methionine [18]. Previous studies have reported that increased HCY levels are a risk factor in relation to multiple diseases, including CV disease, neurodegenerative disease, and renal and vascular complications of diabetes [19–21]. Furthermore, there was an association of HCY and non-cardiac vascular diseases [22]. *In vivo*, hyperhomocysteinemia mice manifested glucose intolerance, insulin resistance, and impaired insulin signaling pathway [23]. As a risk factor, HCY is strongly linked to CV complications in T2DM. Nevertheless, there are still some controversial reports regarding the relationship of circulating HCY levels with insulin resistance in T2DM.

High-sensitivity C-reactive protein (hs-CRP), as one of the important biomarkers of inflammation, is mainly produced and released by the liver when stimulated by inflammatory factors

[24]. Serum concentration of hs-CRP was closely associated with systemic inflammatory conditions in the body. hs-CRP is closely linked to metabolic syndrome features; there was an increased hs-CRP concentration in the IGT group compared to normal controls [25, 26]. Studies have reported that hs-CRP concentrations were associated with insulin resistance and the development of diabetic nephropathy and coronary heart disease (CHD) risk [27–29].

In the present study, we aimed to investigate the expression levels of serum RBP-4, Cys C, HCY and hs-CRP in patients with newly diagnosed T2DM (NT2DM) by comparison with PDM subjects and healthy controls, and analyze their correlations with clinical manifestations and laboratory indexes.

Material and methods

Study subjects and methods

From June 2016 to December 2017, a total of 242 subjects (141 NT2DM patients, 48 PDM subjects and 53 normal controls) were recruited for the measurement of serum RBP-4, Cys C, HCY and hs-CRP concentrations. NT2DM patients and PDM subjects came from the Department of Endocrinology at the Second Hospital of Hefei City, when they first visited the DM clinic. The diagnosis of T2DM was established according to the American Diabetes Association diagnostic criteria 2018 [30]. PDM was defined as those without DM but a fasting plasma glucose (FPG) value ≥ 5.6 mmol/l and FPG < 6.9 mmol/l or a 2-h plasma glucose (2-h PG) value ≥ 7.8 mmol/l and 2-h PG < 11.1 mmol/l during a 75-g oral glucose tolerance test (OGTT) or glycated hemoglobin (HbA_{1c}) $\geq 5.7\%$ and $\text{HbA}_{1c} < 6.4\%$. Fifty-three healthy volunteers from a medical examination center, without liver or metabolic diseases, acute CV or cerebrovascular accidents, were included as normal controls. None of the study subjects were taking any medications during the study. Demographics, clinical manifestations and routine laboratory results were obtained from hospital medical records and then reviewed by experienced physicians.

Standard protocol approvals and patient consent

This study was approved by the Ethical Committee of the Second Hospital of Hefei City (Hefei, Anhui, China). All the study subjects provided informed consent to participate in this study.

All studies on humans described in the present manuscript were carried out with the approval of the responsible ethics committee and in accordance with national law and the Helsinki Declaration of 1975 (in its current, revised form).

Laboratory analyses

After a fasting period of at least 10–12 h overnight, venous blood samples were collected. Fasting blood glucose (FBG), total cholesterol (TC), triglycerides (TGs), high-density lipoprotein cholesterol (HDL-C), low-density lipoprotein cholesterol (LDL-C), very low-density lipoprotein cholesterol (VLDL-C), alanine aminotransferase (ALT), alkaline phosphatase (ALP), aspartate aminotransferase (AST), γ -glutamyltransferase (GGT), lactate dehydrogenase (LDH), total bilirubin (TBIL), indirect bilirubin (IBIL), direct bilirubin (DBIL), Cr and uric acid (UA) were measured using a model 7600 automated bio-analyzer (Hitachi, Tokyo, Japan).

Blood pressure was measured after 15 min rest at the left arm in a sitting position using a sphygmomanometer. Body mass index (BMI) was calculated by dividing subjects' weight by their squared height (kg/m^2).

Apolipoprotein (Apo)-A1 and Apo-B were measured by immunoturbidimetry (Roche/Cobas Integra Tinaquant, Roche Diagnostics).

HCY concentration was determined by the chemical luminescence method (Abbott Laboratories, Abbott Park, Illinois, USA).

The OGTT, with a standard oral glucose load (75 g), was performed after a 10–12-hour fast with venous blood sampling in the fasting state, then fasting blood glucose (FBG), 30 min blood glucose (BG), 60 min BG, 120 min (2 h) BG, fasting insulin, 30 min insulin, 60 min insulin, 120 min insulin, fasting C-peptide, 30 min insulin C-peptide, 60 min C-peptide and 120 min insulin C-peptide were measured. HbA_{1c} was measured using cation-exchange column chromatography on an automatic analyzer (Bio-Rad Company, Hercules, California, USA).

Surrogate markers of insulin sensitivity and secretion were computed, including homeostasis model of assessment of insulin resistance ($\text{HOMA-IR} = (\text{fasting serum insulin (mmol/ml)} \times \text{FPG (mmol/l)})/22.5$) [31], HOMA β cell function index ($\text{HOMA-}\beta = (\text{FINS } (\mu\text{U/ml}) \times 20/\text{FPG (mmol/l)} - 3.5)$) [32] and the Matsuda index of insulin sensitivity ($\text{Matsuda ISI} = 10000/(\text{Glu0} \times \text{Ins0})^{1/2} \times (\text{Glumean} \times \text{Insmean})^{1/2}$) [33].

Enzyme-linked immunosorbent assay (ELISA) for serum RBP-4, Cys C and hs-CRP

Blood samples were obtained from 5 ml of whole blood of all study subjects and then frozen at -80°C immediately after collection until assayed. Serum RBP-4, Cys C, and hs-CRP concentrations were determined by specific ELISA kits according to the manufacturer's recommendation (R&D Systems, Inc).

Statistical analysis

Collected data were presented as mean \pm standard deviation (SD), or median (interquartile range, IQR) if they were not in normal distribution. One-way ANOVA or a nonparametric test (Kruskal-Wallis test) was applied to compare the statistical differences of continuous variables among study groups. The χ^2 test or Fisher's exact test was used to analyze categorical variables. Correlation analyses were computed using Spearman's rank correlation coefficient. Receiver operating characteristic (ROC) analysis was performed and the area under the curve (AUC) was applied to assess specificity and sensitivity of using serum Cys C and hs-CRP as biomarkers for T2DM or PDM. Statistical analysis was performed with the Statistical Package of Social Sciences (SPSS) 23.0 (SPSS Inc., Chicago, IL, USA) and MedCalc, version 11.4.2.0 (Mariakerke, Belgium). All results with a two-tailed $p < 0.05$ were considered to be statistically significant.

Results

Characteristics of the study population

Table I shows the demographic and clinical characteristics of the study subjects. The distributions of age and gender in NT2DM, PDM and normal controls were not significantly different (all $p > 0.05$). However, there were significant differences in BMI, waist circumference (WC), hip circumference, waist-to-hip ratio (WHR), systolic blood pressure (SBP), diastolic blood pressure (DBP), UA, IBIL, ALT, GGT, LDH, TC, TG, LDL-C, HDL-C and VLDL-C levels among NT2DM, PDM and NC groups. The pancreatic function was assessed in accordance to different time points of blood glucose, insulin, C-peptide, and the computed homeostasis model assessment of insulin resistance (HOMA-IR), homeostasis model assessment of β cell function (HOMA- β) and Matsuda ISI (Table II).

Comparison of RBP-4, Cys C, HCY and hs-CRP concentrations between NT2DM patients, PDM subjects and normal controls

There were significant differences in serum Cys C and hs-CRP levels among the three groups (both $p < 0.05$). Compared to normal controls, there were significantly higher Cys C and hs-CRP levels in the PDM group (both $p < 0.001$), a significantly increased Cys C level in the NT2DM group ($p < 0.001$), but no significant differences in Cys C and hs-CRP levels between NT2DM and PDM groups, and no significant differences in hs-CRP levels between NT2DM and normal controls. In addition, the results indicated that there were no sig-

Table I. Clinical characteristics of study subjects

Parameters	NT2DM (n = 141)	PDM (n = 48)	NC (n = 53)	P-value
Age [years]	50.8 ±10.1	49.7 ±13.0	47.0 ±8.0	0.078
Gender (female/male)	69/72	23/25	25/28	0.974
BMI [kg/cm ²]	26.5 (25.4, 27.7)	25.7 (24.1, 26.8)	23.6 (22.1, 24.6)	< 0.001
WC [cm]	88.72 ±9.94	91.87 ±9.00	83.08 ±7.81	< 0.001
HIP [cm]	92.06 ±8.14	98.63 ±8.39	95.19 ±6.96	< 0.001
WHR	0.95 (0.92, 1.04)	0.90 (0.88, 0.94)	0.86 (0.83, 0.88)	< 0.001
SBP [mm Hg]	135 ±16	136 ±16	125 ±14	0.001
DBP [mm Hg]	83 ±11	83 ±11	77 ±10	0.001
Urea nitrogen [mmol/l]	5.10 ±1.48	5.09 ±1.56	4.82 ±1.18	0.464
Cr [μmol/l]	57.97 ±14.37	59.98 ±14.20	62.85 ±13.80	0.101
UA [μmol/l]	342.04 ±63.36	319.25 ±75.46	294.15 ±54.47	< 0.001
TBIL [μmol/l]	15.9 (13.2, 21.2)	15.7 (11.5, 20.5)	13.6 (11.3, 16.6)	0.074
DBIL [μmol/l]	4.2 (3.3, 5.5)	3.7 (2.7, 5.1)	4.2 (3.0, 5.1)	0.540
IBIL [μmol/l]	11.9 (9.6, 16.3)	12.0 (8.4, 15.1)	9.6 (7.9, 12.5)	0.045
ALP [U/l]	76 ±21	76 ±21	71 ±22	0.289
AST [U/l]	20 (16, 28)	24 (18, 29)	19 (16, 24)	0.019
ALT [U/l]	35 (32, 38)	31 (26, 36)	18 (13, 27)	< 0.001
GGT [U/l]	36 (26, 54)	34 (19, 68)	23 (16, 34)	0.007
LDH [U/l]	166 ±33	189 ±38	184 ±32	< 0.001
TG [mmol/l]	2.4 ±0.6	1.9 ±0.7	1.4 ±0.6	< 0.001
TC [mmol/l]	5.4 ±0.3	4.8 ±0.2	4.1 ±0.4	< 0.001
HDL-C [mmol/l]	1.4 ±0.3	1.5 ±0.3	1.6 ±0.4	< 0.001
LDL-C [mmol/l]	3.3 ±0.4	2.9 ±0.7	2.2 ±0.2	< 0.001
VLDL-C [mmol/l]	0.4 (0.3, 0.6)	0.3 (0.2, 0.5)	0.2 (0.1, 0.4)	0.003
Apo-B [g/l]	0.9 (0.7, 1.1)	0.9 (0.7, 1.0)	0.9 (0.8, 1.0)	0.403
Apo-A1 [g/l]	1.1 ±0.2	1.2 ±0.2	1.2 ±0.2	< 0.001

Apo-A1 – apolipoprotein-A1, Apo-B – apolipoprotein-B, ALP – alkaline phosphatase, AST – aspartate aminotransferase, ALT – alanine transaminase, BMI – body mass index, Cr – creatine, DBIL – direct bilirubin, DBP – diastolic blood pressure, FBG – fasting blood glucose, GGT – γ -glutamyltransferase, HDL-C – high-density lipoprotein cholesterol, IBIL – indirect bilirubin, LDH – lactate dehydrogenase, LDL-C – low-density lipoprotein cholesterol, NT2DM – newly diagnosed type 2 diabetes mellitus, NC – normal control, PDM – prediabetes mellitus, SBP – systolic blood pressure, TC – total cholesterol, TG – triglycerides, TBIL – total bilirubin, UA – uric acid, VLDL-C – very low-density lipoprotein cholesterol, WC – waist circumference, WHR – waist-to-hip ratio.

nificant differences in serum RBP-4 and HCY levels among NT2DM, PDM and NC groups (all $p > 0.05$) (Table III).

Correlations of Cys C levels with clinical and laboratory parameters among study groups

The results of univariate correlation analysis showed that Cys C concentrations were significantly correlated with age, WHR, FBG, C-peptide (0 min, 120 min), urea nitrogen, Cr, UA, TBIL, IBIL,

AST, HDL-C, VLDL-C and HOMA- β in the NT2DM group (all $p < 0.05$), and correlated with SBP, Cr, UA and LDL-C in the PDM group (all $p < 0.05$). Moreover, there was a significant association of serum Cys C levels with age, BMI, SBP, FBG, C-peptide (0 min, 120 min), Cr, UA, DBIL, ALT, AST, GGT, TG, TC, HDL-C, VLDL-C, HOMA-IR and Matsuda ISI in normal controls (all $p < 0.05$). However, no significant correlations of Cys C with other clinical and quantitative laboratory parameters were observed (all $p > 0.05$) (Table IV).

Table II. Evaluation of pancreatic function of study population

Parameter	NT2DM (n = 141)	PDM (n = 48)	Normal control (n = 53)	P-value
FBG [mmol/l]	7.44 ±1.07	5.54 ±0.65	4.84 ±0.62	< 0.001
30 min BG (OGTT) [mmol/l]	13.19 ±1.71	10.57 ±1.80	8.78 ±1.63	< 0.001
60 min BG (OGTT) [mmol/l]	15.77 ±1.85	11.69 ±2.15	8.06 ±2.49	< 0.001
120 min BG (OGTT) [mmol/l]	16.90 (14.00, 17.88)	8.93 (8.05, 9.75)	6.05 (5.34, 6.71)	< 0.001
Insulin (OGTT 0 min) [mU/l]	6.31 (5.24, 8.47)	7.77 (6.05, 8.70)	6.63 (5.54, 8.26)	0.021
Insulin (OGTT 30 min) [mU/l]	16.81 (10.41, 28.04)	43.99 (30.14, 63.56)	51.80 (33.60, 71.89)	< 0.001
Insulin (OGTT 60 min) [mU/l]	23.88 (16.22, 38.08)	65.67 (47.54, 80.36)	55.08 (38.46, 75.87)	< 0.001
Insulin (OGTT 120 min) [mU/l]	28.86 (17.24, 48.99)	64.72 (42.09, 89.68)	34.17 (21.53, 47.12)	< 0.001
0 min C-peptide (OGTT) [ng/ml]	1.48 ±0.70	1.76 ±0.66	1.26 ±0.56	0.002
30 min C-peptide (OGTT) [ng/ml]	2.46 (1.73, 3.77)	5.14 (3.77, 6.36)	4.70 (3.15, 5.99)	< 0.001
60 min C-peptide (OGTT) [ng/ml]	3.61 (2.29, 5.31)	7.78 (5.88, 8.77)	6.20 (4.56, 8.59)	< 0.001
120 min C-peptide (OGTT) [ng/ml]	5.50 ±3.26	8.69 ±2.95	5.59 ±2.33	< 0.001
HbA _{1c} (%)	7.91 ±0.42	6.06 ±0.28	5.12 ±0.31	< 0.001
HOMA-IR	2.20 (1.73, 2.78)	1.82 (1.45, 2.22)	1.44 (1.19, 1.86)	< 0.001
HOMA-β	32.89 (24.32, 45.60)	81.09 (59.16, 109.50)	117.11 (74.55, 158.77)	< 0.001
Matsuda ISI	93.38 ±38.31	78.76 ±24.14	114.64 ±44.28	< 0.001

BG – blood glucose, FBG – fasting blood glucose, HOMA-IR – homeostasis model assessment of insulin resistance, HOMA-β – HOMA beta cell function index, HbA_{1c} – glycosylated hemoglobin, OGTT – oral glucose tolerance test, PDM – prediabetes mellitus, NT2DM – newly diagnosed type 2 diabetes mellitus.

Table III. Comparison of RBP-4, Cys C, HCY and hs-CRP levels among three groups

Parameter	NT2DM (n = 141)	PDM (n = 48)	NC (n = 53)	P-value
RBP-4 [µg/ml]	49.00 (39.95, 56.95)	47.55 (40.65, 63.63)	50.90 (43.45, 60.05)	0.461
Cys C [mg/ml]	0.87 (0.75, 1.00)*	0.83 (0.77, 1.05)**	0.76 (0.65, 0.86)	< 0.001
HCY [µmol/l]	11.30 (9.40, 14.35)	11.60 (10.50, 14.03)	10.90 (8.65, 12.95)	0.350
hs-CRP [µg/ml]	0.55 (0.40, 0.74)	0.61 (0.50, 0.99)**	0.50 (0.40, 0.57)	0.006

*Significant difference in NT2DM versus NC. **Significant difference in PDM versus NC. RBP-4 – retinol binding protein-4, Cys C – cystatin C, HCY – homocysteine, Hs-CRP – high-sensitivity C-reactive protein, PDM – prediabetes mellitus, NT2DM – newly diagnosed type 2 diabetes mellitus, NC – normal control.

Correlations of hs-CRP levels with clinical and laboratory parameters among study groups

Univariate correlation analysis demonstrated that serum hs-CRP levels positively associated with BMI, ALP and HOMA-β in the NT2DM group (all $p < 0.05$), and positively associated with 2 h BG, but negatively associated with GGT in the PDM group (all $p < 0.05$). In addition, in normal controls, there was a significant positive association of serum hs-CRP levels with C-peptide (0 min, 120 min), UA, GGT, and ALP and a negative association with HDL-C and Apo-A1 (all $p < 0.05$). However, no significant correlations of hs-CRP with other clinical and quantitative

laboratory parameters were found (all $p > 0.05$) (Table V).

Correlations of HCY levels with clinical and laboratory parameters among study groups

In the NT2DM group, HCY level correlated with Cr, UA, TBIL, DBIL, IBIL, glutamic-pyruvic transaminase (GPT), AST, LDH and Apo-A1 (all $p < 0.05$), while in the PDM group, HCY level positively associated with WHR, but negatively associated with ALP (all $p < 0.05$). Furthermore, we found that HCY level positively correlated with SBP, DBP, UA, ALP, LDH and HDL-C in normal controls (all $p < 0.05$). No significant correlations of HCY with other clinical and quantitative laboratory parameters were observed (all $p > 0.05$) (Table VI).

Table IV. Association of Cys C level with demographic and clinical characteristics

Parameter	NT2DM (n = 141)		PDM (n = 48)		Normal control (n = 53)	
	r	p	r	p	r	p
Age	0.196	0.020	0.056	0.708	0.297	0.031
BMI	0.117	0.166	0.177	0.228	0.361	0.008
WHR	0.261	0.002	0.254	0.082	0.260	0.060
SBP	0.115	0.175	0.346	0.016	0.542	< 0.001
DBP	0.087	0.303	0.196	0.181	0.268	0.052
FBG	-0.246	0.003	0.080	0.591	0.303	0.027
120 min BG (OGTT)	-0.141	0.095	-0.029	0.843	0.166	0.234
Insulin (OGTT 0 min)	0.159	0.060	0.186	0.207	0.261	0.060
Insulin (OGTT 120 min)	0.129	0.129	0.134	0.363	-0.004	0.979
C-peptide 0 min (OGTT)	0.253	0.002	0.169	0.250	0.583	< 0.001
C-peptide 120 min (OGTT)	0.173	0.040	0.190	0.195	0.339	0.013
Urea nitrogen	0.186	0.027	0.131	0.376	0.220	0.113
Cr	0.462	< 0.001	0.467	0.001	0.424	0.002
UA	0.192	0.023	0.409	0.004	0.468	< 0.001
TBIL	0.210	0.013	0.181	0.219	-0.160	0.253
DBIL	0.157	0.063	0.168	0.253	-0.330	0.016
IBIL	0.201	0.017	0.185	0.208	-0.007	0.963
ALT	0.144	0.088	0.160	0.277	0.506	< 0.001
AST	0.246	0.003	0.245	0.093	0.438	0.001
GGT	0.108	0.203	0.271	0.063	0.359	0.008
ALP	0.164	0.052	0.020	0.895	0.222	0.110
LDH	0.072	0.397	0.188	0.200	0.136	0.330
TG	0.021	0.805	0.006	0.970	0.481	< 0.001
TC	-0.021	0.808	0.022	0.883	0.365	0.007
HDL-C	-0.255	0.002	-0.126	0.393	-0.334	0.015
LDL-C	0.024	0.781	0.293	0.043	0.103	0.461
VLDL-C	0.172	0.042	0.096	0.516	0.478	< 0.001
Apo-B	-0.091	0.284	0.268	0.065	0.113	0.420
Apo-A1	-0.089	0.294	-0.192	0.192	-0.244	0.079
HOMA-IR	0.082	0.332	0.205	0.163	0.354	0.009
HOMA-β	0.250	0.003	0.039	0.791	-0.120	0.392
ISI Matsudo	-0.123	0.146	-0.153	0.300	-0.301	0.028

Apo-A1 – apolipoprotein-A1, Apo-B – apolipoprotein-B, ALP – alkaline phosphatase, ALT – alanine transaminase, AST – aspartate aminotransferase, BMI – body mass index, BU – blood urea, Cr – creatine, Cys C – cystatin C, DBIL – direct bilirubin, DBP – diastolic blood pressure, FBG – fasting blood glucose, GGT – γ -glutamyltransferase, HDL-C – high-density lipoprotein cholesterol, IBIL – indirect bilirubin, LDH – lactate dehydrogenase, LDL-C – low-density lipoprotein cholesterol, NT2DM – newly diagnosed type 2 diabetes mellitus, NC – normal control, PDM – prediabetes mellitus, SBP – systolic blood pressure, TC – total cholesterol, TG – triglycerides, TBIL – total bilirubin, UA – uric acid, VLDL-C – very low-density lipoprotein cholesterol, WC – waist circumference, WHR – waist-to-hip ratio.

Table V. Association of hs-CRP level with demographic and clinical characteristics

Parameters	NT2DM (n = 141)		PDM (n = 48)		Normal control (n = 53)	
	r	p	r	p	r	p
Age	-0.073	0.393	0.046	0.756	-0.094	0.504
BMI	0.258	0.002	0.267	0.067	0.130	0.352
WHR	0.117	0.169	0.098	0.506	0.235	0.091
SBP	0.009	0.913	0.100	0.501	0.160	0.253
DBP	0.113	0.180	0.156	0.290	-0.031	0.824
FBG	-0.088	0.297	0.031	0.837	0.041	0.769
120 min BG (OGTT)	0.055	0.520	-0.290	0.046	-0.076	0.591
Insulin (OGTT 0 min)	0.139	0.101	0.103	0.485	0.242	0.081
Insulin (OGTT 120 min)	0.034	0.691	-0.029	0.844	0.088	0.531
C-peptide 0 min (OGTT)	0.124	0.142	0.279	0.055	0.430	0.001
C-peptide 120 min (OGTT)	0.057	0.503	0.224	0.127	0.291	0.034
Urea nitrogen	-0.024	0.780	-0.028	0.848	-0.080	0.567
Cr	-0.025	0.765	0.047	0.749	0.110	0.433
UA	0.130	0.125	0.251	0.085	0.273	0.048
TBIL	0.017	0.841	-0.026	0.859	-0.002	0.989
DBIL	-0.079	0.355	0.000	0.997	-0.190	0.172
IBIL	0.047	0.581	-0.048	0.747	0.043	0.762
ALT	0.136	0.109	0.057	0.699	-0.005	0.970
AST	0.019	0.823	0.174	0.238	-0.034	0.808
GGT	0.113	0.181	0.312	0.031	0.302	0.028
ALP	0.179	0.033	0.074	0.616	0.412	0.002
LDH	0.086	0.312	0.105	0.477	0.151	0.281
TG	-0.039	0.649	-0.023	0.879	0.243	0.080
TC	-0.035	0.680	0.247	0.091	0.080	0.568
HDL-C	-0.100	0.237	-0.194	0.187	-0.351	0.010
LDL-C	-0.040	0.640	0.261	0.073	-0.084	0.549
VLDL-C	0.086	0.309	0.084	0.572	0.206	0.139
Apo-B	-0.010	0.910	-0.026	0.862	-0.174	0.212
Apo-A1	-0.161	0.056	-0.243	0.096	-0.340	0.013
HOMA-IR	0.114	0.176	0.119	0.422	0.216	0.121
HOMA-β	0.181	0.032	0.054	0.717	0.053	0.706
ISI	-0.104	0.220	0.022	0.883	-0.268	0.053

Apo-A1 – apolipoprotein-A1, Apo-B – apolipoprotein-B, ALP – alkaline phosphatase, ALT – alanine transaminase, AST – aspartate aminotransferase, BMI – body mass index, BU – blood urea, Cr – creatine, DBIL – direct bilirubin, DBP – diastolic blood pressure, FBG – fasting blood glucose, GGT – γ-glutamyltransferase, hs-CRP – high-sensitivity C-reactive protein, HDL-C – high-density lipoprotein cholesterol, IBIL – indirect bilirubin, LDH – lactate dehydrogenase, LDL-C – low-density lipoprotein cholesterol, NT2DM – newly diagnosed type 2 diabetes mellitus, NC – normal control, PDM – prediabetes mellitus, SBP – systolic blood pressure, TC – total cholesterol, TG – triglycerides, TBIL – total bilirubin, UA – uric acid, VLDL-C – very low-density lipoprotein cholesterol, WC – waist circumference, WHR – waist-to-hip ratio.

Table VI. Association of HCY level with demographic and clinical characteristics

Parameter	NT2DM (n = 141)		PDM (n = 48)		Normal control (n = 53)	
	r	p	r	p	r	p
Age	0.086	0.312	-0.039	0.793	0.137	0.329
BMI	0.060	0.478	0.018	0.901	0.148	0.289
WHR	0.160	0.057	0.405	0.004	0.009	0.950
SBP	-0.015	0.857	0.089	0.545	0.395	0.003
DBP	0.045	0.597	0.107	0.471	0.377	0.005
FBG	-0.103	0.223	-0.091	0.539	0.015	0.915
120 min BG (OGTT)	0.003	0.968	-0.073	0.621	-0.024	0.866
Insulin (OGTT 0 min)	0.075	0.375	0.062	0.674	-0.051	0.716
Insulin (OGTT 120 min)	0.068	0.420	0.076	0.608	-0.127	0.364
C-peptide 0 min (OGTT)	-0.014	0.872	-0.027	0.854	0.163	0.243
C-peptide 120 min (OGTT)	0.001	0.989	0.168	0.255	0.104	0.457
Urea nitrogen	0.069	0.413	-0.032	0.828	-0.026	0.852
Cr	0.302	< 0.001	0.233	0.111	0.185	0.185
UA	0.189	0.025	0.185	0.209	0.337	0.014
TBIL	0.229	0.006	0.050	0.735	-0.041	0.768
DBIL	0.238	0.004	0.119	0.419	-0.095	0.500
IBIL	0.206	0.014	0.017	0.910	-0.029	0.835
GPT	0.207	0.014	0.180	0.222	0.210	0.131
AST	0.219	0.009	0.149	0.312	0.265	0.056
ALT	0.044	0.607	0.280	0.054	0.178	0.202
ALP	0.160	0.058	-0.394	0.006	0.365	0.007
LDH	0.195	0.021	0.278	0.056	0.429	0.001
TG	0.005	0.949	-0.133	0.367	0.230	0.097
TC	0.024	0.777	0.169	0.250	0.254	0.067
HDL-C	-0.265	0.001	-0.097	0.512	-0.273	0.048
LDL-C	-0.134	0.114	0.015	0.921	0.069	0.623
VLDL-C	0.012	0.883	-0.005	0.974	0.183	0.190
Apo-B	-0.093	0.273	0.188	0.200	0.060	0.669
Apo-A1	-0.166	0.049	-0.009	0.950	-0.184	0.188
HOMA-IR	0.030	0.724	0.047	0.749	-0.057	0.687
HOMA-β	0.108	0.204	0.121	0.413	-0.040	0.774
ISI	-0.082	0.336	-0.084	0.572	0.037	0.792

Apo-A1 – apolipoprotein-A1, Apo-B – apolipoprotein-B, ALP – alkaline phosphatase, ALT – alanine transaminase, AST – aspartate aminotransferase, BMI – body mass index, BU – blood urea, Cr – creatine, DBIL – direct bilirubin, DBP – diastolic blood pressure, FBG – fasting blood glucose, GPT – glutamic-pyruvic transaminase, HCY – homocysteine, HDL-C – high-density lipoprotein cholesterol, IBIL – indirect bilirubin, LDH – lactate dehydrogenase, LDL-C – low-density lipoprotein cholesterol, NT2DM – newly diagnosed type 2 diabetes mellitus, NC – normal control, PDM – prediabetes mellitus, SBP – systolic blood pressure, TC – total cholesterol, TG – triglycerides, TBIL – total bilirubin, UA – uric acid, VLDL-C – very low-density lipoprotein cholesterol, WC – waist circumference, WHR – waist-to-hip ratio.

Correlations of RBP-4 levels with clinical and laboratory parameters among study groups

The correlation analysis demonstrated that serum RBP-4 concentration was correlated positively with Cr, UA, ALT, VLDL-C and Apo-B in the NT2DM group (all $p < 0.05$), positively associated with Cr, UA, DBIL and ALT in the PDM group (all $p < 0.05$), and there was a positive association with urea nitrogen, IBIL, GPT, AST and VLDL-C in the normal control group (all $p < 0.05$). However, no significant correlations of RBP-4 with other clinical and quantitative laboratory parameters were observed (all $p > 0.05$) (Table VII).

Predictive accuracy of Cys C and hs-CRP for diagnostic biomarkers of NT2DM

An ROC curve was drawn to determine the diagnostic performance of Cys C and hs-CRP as NT2DM biomarkers. The Cys C had an area under the curve (AUC) value of 0.683 (95% CI: 0.612–0.747), with 58.16% sensitivity and 75.47% specificity (Figure 1 A). For hs-CRP, the AUC value was 0.578 (95% CI: 0.505–0.649), with 46.81% sensitivity and 77.36% specificity (Figure 1 B).

Predictive accuracy of Cys C and hs-CRP for diagnostic biomarkers of PDM

The diagnostic performance of Cys C and hs-CRP as PDM biomarkers was assessed by ROC curve and AUC. The AUC value of Cys C is 0.696 (95% CI: 0.597–0.784), with 75.00% sensitivity and 54.72% specificity (Figure 2 A). Moreover, hs-CRP had an AUC value of 0.694 (95% CI: 0.595–0.782), with 81.13% sensitivity and 56.25% specificity (Figure 2 B).

Discussion

Although RBP-4, Cys C, HCY and hs-CRP have been studied in many metabolic diseases, there are very few studies regarding the expression of RBP-4, Cys C, HCY and hs-CRP and their association with NT2DM. In the present study, we investigated the serum RBP-4, Cys C, HCY and hs-CRP concentrations in NT2DM and their relations with clinical and laboratory features. The current study demonstrated that, in comparison to healthy controls, there were significantly increased Cys C levels in NT2DM and significantly elevated Cys C and hs-CRP levels in PDM. Both the Cr and UA were correlated with Cys C concentration among NT2DM, PDM and normal controls. Furthermore, hs-CRP level was correlated with BMI, ALP and HOMA- β in the NT2DM group, and was associated with 2 h BG and GGT in the PDM group. In normal controls, serum hs-CRP level cor-

related with C-peptide (0 min, 120 min), UA, GGT, ALP, HDL-C and Apo-A1. However, we did not find any differences of RBP-4 and HCY concentration among NT2DM, PDM and normal controls.

As we know, patients with T2DM have a high incidence of renal involvement [34, 35]. Cys C is a cysteine protease inhibitor that is produced at a constant rate by all nucleated cells and freely filtered by glomeruli owing to its low molecular weight; thus, it can freely pass through the glomeruli and be reabsorbed in proximal tubules, followed by complete catabolism without being affected by features influencing Cr level [11, 36]. Previous studies have revealed that there was a higher Cys C level in T2DM patients than in controls, and it was associated with renal dysfunction [12, 37]. Consistent with prior studies, we found that in NT2DM patients, there was a higher Cys C level than in healthy controls, and serum Cys C concentrations were positively associated with age, WHR, C-peptide (0 min, 120 min), urea nitrogen, Cr, UA, TBIL, IBIL, AST, HDL-C, VLDL-C and HOMA- β , and were negatively associated with FBG and HDL-C. In addition, there was a higher Cys C level in PDM than in the control group. The increased Cys C level was correlated with SBP, Cr, UA and LDL-C. It has been reported that the Cys C level is able to increase at the stage of PDM, and a study of 83 patients with T2DM showed that Cys C is a good marker of incipient renal disease, which may be because minimal glomerular damage could result in a significant increase of serum Cys C level, causing disease progression [38]. Therefore, early detection of Cys C not only helps to predict the possible potential diabetes mellitus risk, but also benefits in identification of early renal function disorder of T2DM.

Hs-CRP has been applied in clinical settings to monitor chronic and acute inflammatory conditions [39]. As a systemic marker, it is extremely sensitive to inflammatory condition and tissue damage and plays a crucial part in atherosclerosis and thus causes CV disease [40, 41]. Serum hs-CRP is positively associated with the metabolic syndrome and has been acknowledged to be an independent risk factor for development of diabetic neuropathy, diabetic foot ulcers and CV complications [42, 43]. Recently, Aryan *et al.* performed a larger sample population-based study to evaluate the predicted value of hs-CRP for complications of T2DM. The results showed that serum hs-CRP associated with the risk of coronary heart disease (CHD) (hazard ratio (HR) = 1.028, 95% CI: 1.024–1.032), diabetic neuropathy (HR = 1.025, 95% CI: 1.021–1.029), diabetic retinopathy (HR = 1.037, 95% CI: 1.030–1.043) and diabetic kidney disease (HR = 1.035, 95% CI: 1.027–1.043) [44]. In the present study, our results showed a signifi-

Table VII. Association of RBP-4 level with demographic and clinical characteristics

Parameter	NT2DM (n = 141)		PDM (n = 48)		Normal control (n = 53)	
	r	p	r	p	r	p
Age	0.013	0.876	0.058	0.695	0.028	0.844
BMI	0.110	0.195	0.098	0.509	-0.182	0.192
WHR	-0.013	0.874	0.051	0.733	-0.076	0.590
SBP	-0.089	0.295	0.085	0.565	0.015	0.917
DBP	-0.026	0.760	0.254	0.082	-0.028	0.844
FBG	0.065	0.443	0.052	0.726	-0.077	0.582
120 min BG (OGTT)	-0.043	0.610	0.033	0.822	-0.008	0.956
Insulin (OGTT 0 min)	0.076	0.372	-0.042	0.777	-0.151	0.280
Insulin (OGTT 120 min)	0.068	0.421	-0.177	0.230	-0.007	0.959
C-peptide 0 min (OGTT)	0.022	0.794	-0.172	0.243	-0.127	0.364
C-peptide 120 min (OGTT)	-0.017	0.844	-0.263	0.071	-0.045	0.752
Urea nitrogen	0.102	0.227	0.258	0.076	0.282	0.041
Cr	0.296	< 0.001	0.469	0.001	0.423	0.002
UA	0.211	0.012	0.415	0.003	0.083	0.553
TBIL	-0.010	0.902	0.213	0.146	0.142	0.310
DBIL	0.039	0.642	0.319	0.027	-0.061	0.666
IBIL	-0.037	0.661	0.176	0.231	0.361	0.008
GPT	0.051	0.551	0.084	0.571	0.289	0.036
AST	0.070	0.408	0.119	0.422	0.306	0.026
ALT	0.196	0.020	0.449	0.001	0.050	0.720
ALP	-0.124	0.142	-0.235	0.107	0.251	0.070
LDH	0.065	0.446	0.056	0.708	0.188	0.178
TG	-0.088	0.297	0.059	0.689	0.135	0.334
TC	0.023	0.784	0.046	0.755	-0.062	0.660
HDL-C	0.129	0.129	0.062	0.677	0.116	0.408
LDL-C	0.035	0.680	-0.227	0.121	0.215	0.122
VLDL-C	0.345	< 0.001	0.159	0.280	0.411	0.002
Apo-B	0.375	< 0.001	0.256	0.079	-0.030	0.829
Apo-A1	0.127	0.134	0.064	0.666	-0.034	0.811
HOMA-IR	0.082	0.333	-0.020	0.893	0.026	0.853
HOMA-β	0.034	0.692	-0.074	0.619	0.061	0.663
ISI	-0.103	0.225	0.024	0.869	0.028	0.844

Apo-A1 – apolipoprotein-A1, Apo-B – apolipoprotein-B, ALP – alkaline phosphatase, ALT – alanine transaminase, AST – aspartate aminotransferase, BMI – body mass index, BU – blood urea, Cr – creatine, DBIL – direct bilirubin, DBP – diastolic blood pressure, FBG – fasting blood glucose, GPT – glutamic-pyruvic transaminase, HDL-C – high-density lipoprotein cholesterol, IBIL – indirect bilirubin, LDH – lactate dehydrogenase, LDL-C – low-density lipoprotein cholesterol, NT2DM – newly diagnosed type 2 diabetes mellitus, NC – normal control, RBP-4 – retinol binding protein-4, PDM – prediabetes mellitus, SBP – systolic blood pressure, TC – total cholesterol, TG – triglycerides, TBIL – total bilirubin, UA – uric acid, VLDL-C – very low-density lipoprotein cholesterol, WC – waist circumference, WHR – waist-to-hip ratio.

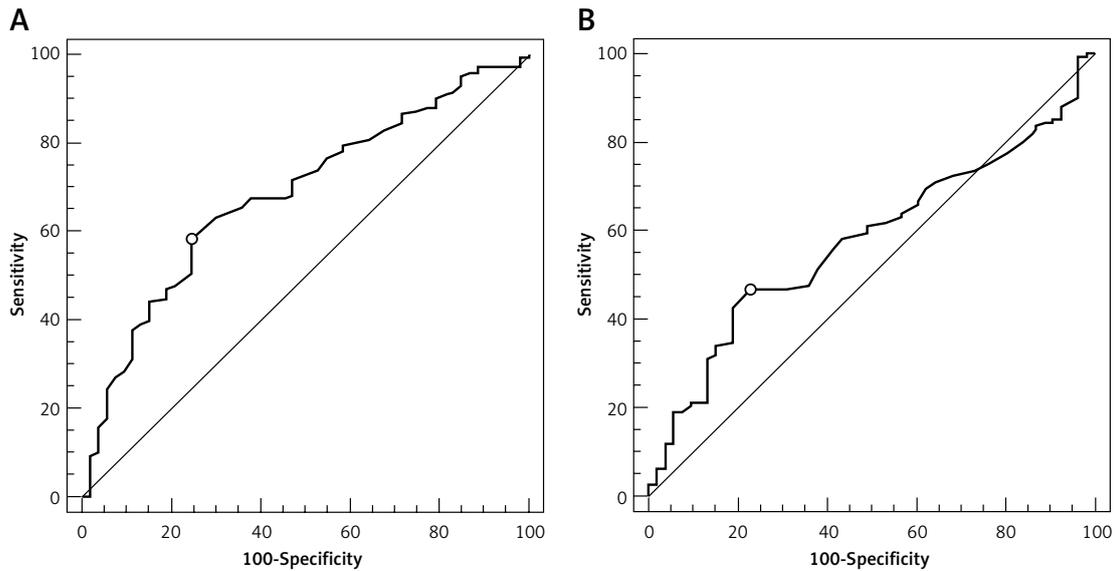


Figure 1. **A** – Receiver operating characteristic (ROC) curve analysis of Cys C for the discriminative ability of NT2DM patients versus healthy controls. **B** – Receiver operating characteristic (ROC) curve analysis of hs-CRP for the discriminative ability of NT2DM patients versus healthy controls

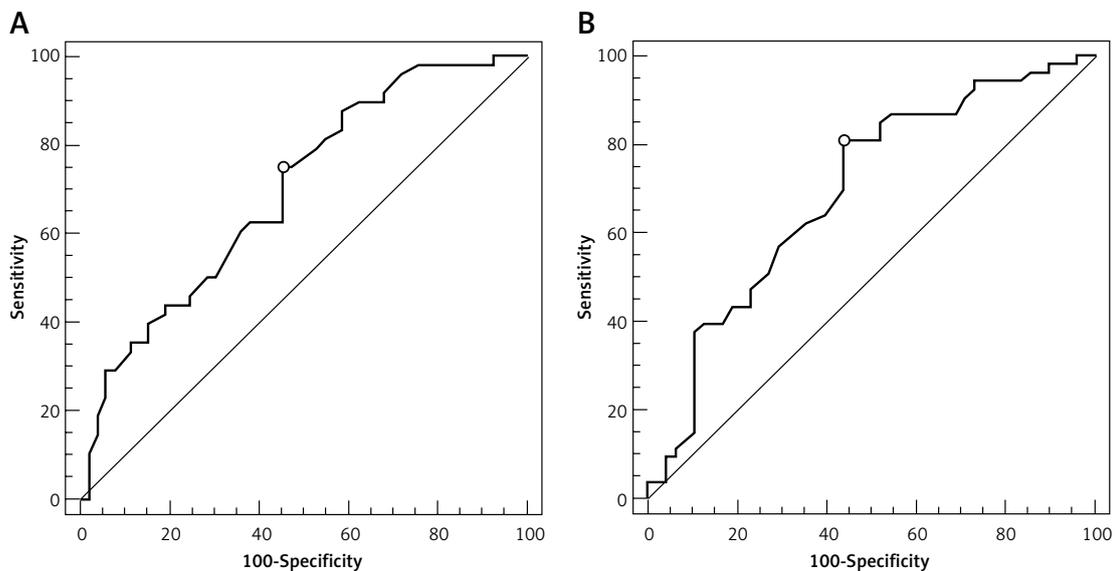


Figure 2. **A** – Receiver operating characteristic (ROC) curve analysis of Cys C for the discriminative ability of PDM subjects versus healthy controls. **B** – Receiver operating characteristic (ROC) curve analysis of hs-CRP for the discriminative ability of PDM subjects versus healthy controls

cantly higher hs-CRP level in the PDM group than in normal controls, but we did not observe any significant differences of hs-CRP between NT2DM and normal controls, as well as NT2DM and PDM groups. It is not consistent with previous studies, which suggested that there was an elevated hs-CRP concentration in T2DM patients compared to healthy controls, and this elevated level of hs-CRP might be an increased risk of diabetes as well [42, 45–47]. The differences in selection of the study population (NT2DM and PDM subjects in our study versus T2DM patients with or without diabetic complications) may cause this discrepancy in hs-CRP concentration. In addition, it has also been

reported that hs-CRP has an association with insulin resistance, of which CRP may contribute to vascular inflammation and cause injury of vascular cells and further contribute to the development of insulin resistance.

Circulating HCY concentrations have been studied in different populations, and the results are not consistent, including lower, higher or unchanged HCY levels in T2DM patients compared to healthy controls [48–51]. However, a meta-analysis of case-control studies demonstrated that there was a significantly higher pooled HCY level in T2DM patients than in healthy controls [52]. In the present study, we did not find any significant

differences in NT2DM, PDM and normal controls. Therefore, the relationships between circulating HCY concentrations and T2DM or PDM remain to be further studied.

Several studies have shown that there was an association of RBP-4 with abnormal glucose tolerance, and an elevated RBP-4 concentration in the T2DM group compared to the healthy control group [13, 17, 53]. In addition, it has been reported that RBP-4 is associated with obesity and insulin resistance. Perseghin *et al.* found that RBP-4 concentration was associated with peripheral insulin sensitivity and correlated with the soleus intramyocellular lipid (IMCL) content and with the intrahepatic lipid (IHL) content [54]. Shaker *et al.* reported that there an increased RBP-4 concentration correlated with BMI, waist/hip ratio, insulin and HOMA-IR. Furthermore, RBP-4 correlated with visceral fat and liver fat in diabetic patients [55]. The present analysis did not reveal differences of RBP-4 concentration among NT2DM, PDM and normal control groups, but the correlation analysis revealed an association between RBP-4 and some clinical and laboratory indexes in NT2DM and PDM groups. The difference of RBP-4 concentration may be attributed to the possible ethnic variation. In addition, the measurement methods used in this study might have influenced the RBP-4 level, compared with methods such as quantitative western blotting (WB), and this could lead to the varied results reported [56]. Therefore, further studies are still needed to study ethnic variations of RBP-4 among different populations.

Nevertheless, several other methodological issues and limitations of our study need to be acknowledged. First, the study design was a case-control, observational study, lacking a clear time structure, and a causal relationship between RBP-4, Cys C, HCY and hs-CRP and NT2DM could not be proved. Second, the variations of different ELISA kits when performing detection of the same plasma factor may lead to unstable measurement results. Furthermore, due to a relatively small sample size and single human race, it may restrict the generalizability of our results. Hence, further studies with a large sample size or in multiple human races are still required to confirm our results.

In conclusion, decreased serum Cys C levels and their association with some clinical and laboratory parameters in NT2DM suggest their potential roles in T2DM. In addition, the results of ROC showed relatively high specificity of Cys C and hs-CRP for identifying the T2DM high-risk population, indicating useful biomarkers for the differential diagnosis of T2DM. However, further mechanism studies and longitudinal large-cohort studies are still needed to further reveal the role of Cys C and hs-CRP in the pathogenesis of T2DM.

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Conflict of interest

The authors declare no conflict of interest.

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