

Increased chitotriosidase activity in plasma of patients with type 2 diabetes

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Abstract

Introduction: Chitotriosidase (CHIT1) is a chitinolytic enzyme involved mainly in the immune and inflammatory response. It shows increased activity in many pathologies, including in newly diagnosed type 2 diabetes (T2D). This study aimed to investigate this enzyme's activity in plasma of patients with ongoing T2D and indicate factors related to the increased activity of this enzyme.

Material and methods: Ninety-one patients and 46 control subjects without abnormalities in carbohydrate metabolism and inflammatory states were enrolled in the study. Plasma CHIT1 activity was measured by a spectrofluorometric method. Routine laboratory parameters such as blood glucose, total cholesterol and HDL fraction, triglyceride, glycated hemoglobin, white blood cell count and C-reactive protein were measured by standard methods.

Results: We found that the chitotriosidase activity was significantly higher ($p < 0.001$) in type 2 diabetic patients and positively associated with parameters of glycemic control (levels of glucose and glycated hemoglobin) and blood pressure. Plasma glucose level and systolic blood pressure were independent determinants of increased CHIT1 activity in T2D patients, even after adjustment for disease duration, body mass index, parameters of inflammation and lipid metabolism. We also found that increased CHIT1 activity was associated with occurrence of diabetic angiopathies.

Conclusions: This investigation indicates a possible role of chitotriosidase in the course of T2D, especially in relation to development of diabetic angiopathies.

Key words: chitinase, type 2 diabetes, diabetic angiopathies, glucose, systolic blood pressure.

Introduction

Chitotriosidase (EC 3.2.1.14), also named chitinase-1 (CHIT1), belongs to the 18 family of chitinases and shows hydrolytic activity against chitin. Although vertebrates do not synthesize chitin, the enzyme was in the 1990s unexpectedly discovered in humans [1–3]. The exact function of human CHIT1 remains unclear, but its participation in immune defense is postulated. The enzyme's ability to cleave the β -(1,4)-glycosidic linkages in chitin determines its antibacterial, antifungal and antiparasitic action [4]. Increased serum CHIT1 activity, resulting from macrophage stimulation, has been recognized as a useful marker in diagnosing and moni-

toring Gaucher disease [5]. Recently it has been reported that increased activity of chitotriosidase may also be connected with the inflammatory state, and therefore a number of investigations on the role of this enzyme in different pathological units, mainly with inflammatory background, have been carried out [6]. Some authors have postulated participation of chitotriosidase in the development of atherosclerosis, which creates a possible link to the course of type 2 diabetes (T2D) [7].

Type 2 diabetes, the most common form of diabetes, results from insulin resistance and/or deficiency and is connected with many metabolic abnormalities, such as hyperglycemia, obesity, hypertension, and dyslipidemia. These are main systemic factors, which trigger proinflammatory events, leading to endothelial dysfunction and increased prevalence of diabetic angiopathies (DA) in T2D diabetic patients. Diabetic angiopathies are the major cause of decreased quality of life and increased mortality of T2D, remaining a constant target of diagnosis, monitoring and treatment [8–12].

The aim of our study was to investigate plasma CHIT1 activity in patients with ongoing type 2 diabetes and evaluate its possible relationship with anthropometric and metabolic parameters, describing the diabetic state. Moreover, we assessed whether this enzyme could provide predictive information for the risk of occurrence of diabetic angiopathies.

Material and methods

Initially, 91 patients with type 2 diabetes mellitus, of average duration 11 years, were recruited from the Clinic of Angiology, Hypertension and Diabetology of Wrocław Medical University. Type 2 diabetes was diagnosed according to the standards set by the Polish Diabetes Association [13]. Diabetic angiopathies were recognized in 44 patients (27% of them had microangiopathy, 30% had macroangiopathy and 43% had both micro- and macroangiopathies). Patients were treated continuously with routine medications (including hypoglycemic, hypotensive and hypolipidemic agents if necessary) for 3 months before examination. The control group consisted of 46 subjects without abnormalities in carbohydrate metabolism (glucose level < 5.5 mmol/l) and a history of inflammatory states during the last 3 months, matched for sex and age. All participants were informed about the aim of the study and their written permission was given. The use of human blood was approved by the Local Bioethics Committee of Wrocław Medical University.

Venous blood was collected after overnight fasting into standard vacuum tubes with heparin (16 IU/ml) to obtain plasma. Routine biochemical parameters were measured on admission to

the hospital by the standard methods using Cobas Mira Plus (ROCHE). Subjects from the control group were examined using the same methods. Plasma samples were stored at -80°C until assayed. To measure chitotriosidase activity a 5 μl plasma sample was incubated with 100 μl of substrate solution (22 μM 4-methylumbelliferyl- β -N-N'-N"-triacetylchitotrioside (Sigma Chemical Co, USA) in McIlvaine's buffer, pH 5.0) for 15 min at 37°C (modified from Hollak *et al.*, 1994). The reaction was terminated using 2 ml of 0.3 M glycine-NaOH buffer (pH 10.5) and fluorescence of the product, 4-methylumbelliferone, was measured with a Perkin Elmer LS 50B fluorometer at excitation $\lambda = 365$ nm and emission $\lambda = 445$ nm. Intra- and interassay coefficients of variation were 4.9% and 5.3%, respectively. Thirteen participants with CHIT1 activity at zero level were excluded from further analysis. Finally, 82 patients and 42 control subjects were evaluated, and their anthropometric and clinical features are presented in Table I.

Participants were divided into subgroups based on: a) body mass index (BMI ≥ 30 kg/m² – obese, ≥ 25 kg/m² and ≤ 29.9 kg/m² – overweight, ≤ 24.9 kg/m² – normal weight), b) presence of dyslipidemia (defined as total cholesterol ≥ 4.9 mmol/l and/or LDL cholesterol (LDL-Ch) ≥ 3.4 mmol/l or HDL cholesterol (HDL-Ch) < 1.0 mmol/l in males and < 1.3 mmol/l in females or triglyceride (TG) ≥ 1.7 mmol/l), c) quality of short-term glycemic control (plasma glucose ≥ 7 mmol/l), d) blood pressure control ($\geq 140/90$), e) presence of inflammation (recognized if white blood cell (WBC) count $\geq 10.9 \times 10^9/\text{l}$ or C-reactive protein (CRP) concentration > 6 mg/l), f) smoking status [13, 14]. Based on the pharmacological treatment of hyperglycemia, the patients were subdivided into those treated with insulin alone ($n = 21$) and those treated with insulin and oral agents (mainly metformin, $n = 61$).

Statistical analysis

The statistical analysis was performed using Statistica PL for Windows, version 10.0. The non-parametric Mann-Whitney *U* test was used for the comparison between continuous variables and the χ^2 test for categorical variables. Univariate and multiple regression analysis was performed to evaluate the association of CHIT1 activity (as the dependent variable) with the explanatory variables in the examined population. Stepwise logistic linear regression was used to evaluate the predictive value of the increased CHIT1 levels (as the independent variable). A *p*-value below 0.05 was considered statistically significant. All results were expressed as a mean value and a standard deviation.

Table I. Anthropometric and clinical characteristics of participants

Parameter	Diabetic subjects	Control subjects
N (male/female)	82 (38/44)	42 (18/24)
Age [years]	59.73 ±5.58	57.37 ±5.51
Disease duration [years]	11.01 ±5.13	–
BMI [kg/m ²]	30.23 ±4.51	27.75 ±3.91*
SBP [mm Hg]	140.00 ±14.00	122.00 ±11.00***
DBP [mm Hg]	80.50 ±9.11	76.50 ±8.50
CRP [mg/l]	10.38 ±9.41	4.4 ±3.49***
WBC [$\times 10^9/l$]	7.64 ±4.91	7.05 ±1.75***
Glucose [mmol/l]	8.93 ±2.91	5.00 ±0.72***
HbA _{1c} (%)	7.8 ±1.54	5.67 ±0.27***
Total-Ch [mmol/l]	5.08 ±1.07	5.19 ±2.36*
HDL-Ch [mmol/l]	1.16 ±0.29	1.45 ±0.38**
LDL-Ch [mmol/l]	2.95 ±0.85	2.95 ±0.61
TG [mmol/l]	2.34 ±0.81	1.56 ±0.38***
Smoking (%)	10	12

Data are presented as mean ± standard deviation or number/percent, statistical significance: * $p < 0.5$, ** $p < 0.01$, *** $p < 0.001$, BMI – body mass index, SBP – systolic blood pressure, DBP – diastolic blood pressure, CRP – C-reactive protein, WBC – white blood cells, HbA_{1c} – glycated hemoglobin, total-Ch – total cholesterol, HDL-Ch – HDL cholesterol, LDL-Ch – LDL cholesterol, TG – triglyceride.

Results

Plasma chitotriosidase activities in patients with type 2 diabetes mellitus and control subjects are presented in Figure 1 (first two bars). A statistically significant ($p < 0.001$) increase of CHIT1 activity (over 70%) was observed in type 2 diabetic

patients compared with the control subjects. No significant differences in CHIT1 activity were observed in subgroups of diabetic patients divided according to BMI value, as well as in subgroups created on the basis of presence of dyslipidemia and inflammation. Neither tobacco use nor ap-

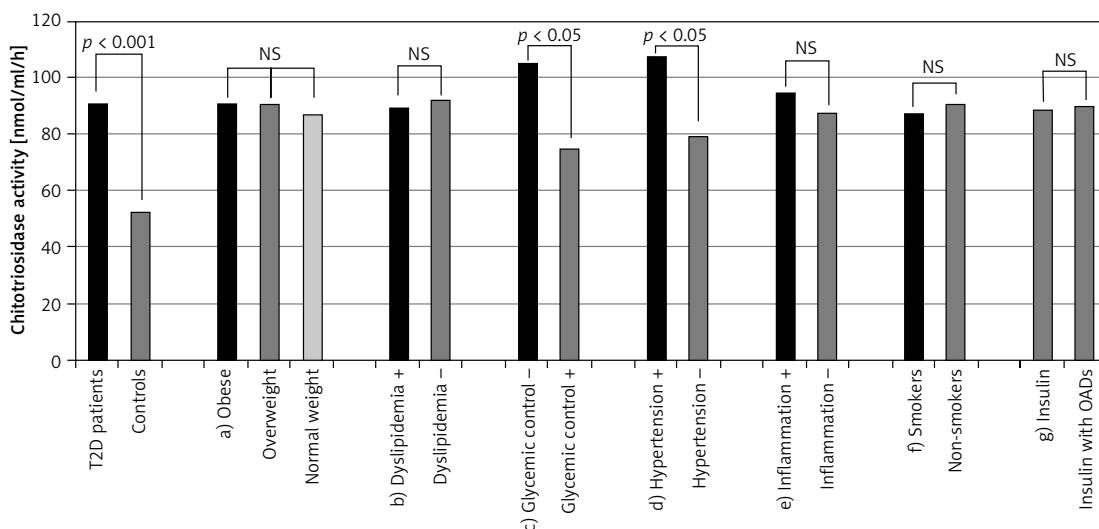


Figure 1. Chitotriosidase activity in examined population and in different subgroups of diabetic patients. First two bars illustrate activity of chitotriosidase in all patients with type 2 diabetes and controls. Next bars illustrate activity of chitotriosidase in distinguished subgroups of diabetic patients: a) patients with obesity, overweight and normal weight, b) patients with and without dyslipidemia, c) patients with poor and good short-term glycemic control, d) patients with poorly controlled hypertension and normalized blood pressure, e) patients with and without symptoms of inflammation, f) smokers and non-smokers, g) patients treated with insulin or insulin with oral antidiabetic drugs (OADs); NS – not statistically significant

plied hypoglycemic treatment had an influence on activity of this enzyme.

To identify the factors associated with increased CHIT1 activity in examined participants (divided or not according to presence of T2D) univariate regression analysis was performed at first, and the results are shown in Table II. Plasma CHIT1 activity was the dependent variable, and the explanatory variables were duration of diabetes, obesity (described by BMI value), parameters of glycemic control (glucose and glycosylated hemoglobin), inflammation (CRP and WBC), lipid metabolism (HDL-Ch, LDL-Ch and TG) and systolic and diastolic blood pressure (SBP, DBP). In the total population CHIT1 activity was significantly associated with plasma glucose and HDL-Ch concentration, SBP and DBP, whereas in type 2 diabetic patients CHIT1 activity was additionally associated with glycosylated hemoglobin (HbA_{1c}), but not with HDL-Ch. However, in the control group CHIT1 activity was associated with CRP and SBP. To further determine the independent factors associated with the increased plasma CHIT1 activity, multiple regression analysis was performed, and the results are given in Table III. Body mass index, plasma glucose, HbA_{1c}, CRP, SBP, DBP, LDL-Ch, HDL-Ch and TG were included in the analyses, provided that the *p* value was below 0.1 in the univariate model. In the present study we focused on T2D patients. As shown in Table III, only plasma glucose and SBP were still independent predictors of increased plasma CHIT1 activity in T2D patients, and 14% of the variance of CHIT1 activity could be predicted by these variables. Plasma glucose has stronger predictive value ($\beta = 0.37$, $p < 0.01$) than SBP ($\beta = 0.27$, $p < 0.05$).

As the last step, we assessed whether the increased CHIT1 activity may predispose to occurrence of diabetic angiopathies. CHIT1 activity was transformed into categorical data, according to the mean value (52.3 nmol/ml/h) observed in the control group, also corresponding to the first quartile of enzyme activity in diabetic patients. Values equal to or above 52.3 nmol/ml/h were considered as increased enzyme activity (signed as yes = 1). Among the individuals with T2D, the proportion of patients with angiopathies was significantly higher in those with CHIT1 activity ≥ 52.3 nmol/ml/h (36/43, which is 86%), compared to patients with CHIT1 activity < 52.3 nmol/ml/h (8/39, which is 20%). A logistic linear regression analysis with the presence of DA (yes/no) as a dependent variable and CHIT1 activity along with other well-known risk factors as predictors was performed, and the results are presented in Table IV. As revealed in the univariate analysis, increased CHIT1 activity, hyperglycemia, presence of inflammation, hypertension and hypertriglyceridemia were predictors of

occurrence of DA. In the multivariate analysis, increased CHIT1 activity remained the independent predictor for occurrence of diabetic angiopathies in the constructed model, and the highest odds ratios were obtained for the presence of inflammation, increased CHIT1 activity and poor short-term glycemic control.

Discussion

In the current study we firstly revealed a significant ($p < 0.001$), almost 2-fold, increase of chitotriosidase activity in plasma of patients with ongoing type 2 diabetes mellitus in comparison to the control group. About 75% of diabetic patients had higher CHIT1 activity than in the control group, which is not fully explicable. To our knowledge, in the scientific literature only a single study describes CHIT1 activity in patients with newly diagnosed, untreated, and uncomplicated T2D [15]. The authors reported increased activity of chitotriosidase in these patients and its significant correlation with age, plasma glucose level and asymmetric dimethylarginine concentration, although the strength of the correlations was rather weak. On the basis of these findings, the authors postulated the relevance of CHIT1 for endothelial functions and insulin resistance in the examined population and concluded that increased CHIT1 activity may be a predictor of endothelial dysfunction. However, reported associations pertained to the whole examined population (with and without diabetes), and no information was provided about CHIT1 specifically in the group of diabetic patients. Moreover, the influence of other parameters, which may also affect endothelial function, e.g. hypercholesterolemia, hypertension, or inflammation, was not taken into consideration [8, 16, 17].

In this work, we undertook research to evaluate the influence of metabolic disturbances accompanying diabetes on CHIT1 activity. First of all, we found that neither obesity nor dyslipidemia significantly affected enzyme activity in the examined patients (Figure 1). Regression analysis assessed the lack of relationship between plasma CHIT1 activity and parameters of lipid metabolism in type 2 diabetic patients and revealed that the enzyme activity was not associated with duration of the disease or BMI value. Alanbay *et al.* [18] also demonstrated that elevation of CHIT1 activity was independent of obesity in women with polycystic ovary syndrome. In contrast to Sonmez *et al.* [15], we did not find any association with age. It may result from the adjustment for age in our study, although the results obtained by Ramanathan *et al.* [19] indicate that elevation of CHIT1 activity may be caused by normal age-related chronic macrophage activation, not their acute

Table II. Univariate regression analysis in examined population with plasma chitotriosidase activity as dependent variable

Covariate	Diabetic subjects			Control subjects			Total (diabetic and control subjects)		
	β	95% CI	P-value	β	95% CI	P-value	β	95% CI	P-value
Disease duration	0.105	(-0.037)-0.248	0.148	-	-	-	-	-	-
BMI	-0.126	(-0.266)-0.013	0.076	0.081	(-0.202)-0.364	0.013	-0.002	(-0.126)-0.123	0.998
Glucose	0.151	0.016-0.285	0.028	0.165	(-0.057)-0.388	0.143	0.153	(-0.062)-0.168	0.036
HbA _{1c}	-0.289	(-0.561)-0.017	0.037	-0.103	(-0.242)-0.036	0.147	0.013	(-0.111)-0.137	0.840
CRP	-0.160	(-0.367)-0.047	0.127	0.264	(-0.010)-0.538	0.048	-0.044	(-0.210)-0.122	0.601
WBC	0.038	(-0.106)-0.181	0.606	-0.126	(-0.370)-0.118	0.306	0.053	(-0.069)-0.175	0.369
SBP	0.204	0.064-0.344	0.005	0.401	0.140-0.661	0.003	0.339	0.221-0.458	< 0.001
DBP	0.168	0.027-0.309	0.020	0.250	(-0.250)-0.525	0.074	0.178	0.054-0.302	0.005
LDL-Ch	0.089	(-0.059)-0.237	0.236	0.176	(-0.103)-0.456	0.211	0.124	0.050-0.253	0.059
HDL-Ch	-0.013	(-0.158)-0.132	0.859	0.009	(-0.245)-0.263	0.942	-0.162	(-0.285)-(-0.040)	0.009
TG	-0.001	(-0.146)-0.143	0.988	-0.243	(-0.447)-(-0.039)	0.061	0.035	(-0.087)-0.157	0.576

Table III. Multiple regression analysis in examined population with plasma chitotriosidase activity as dependent variable

Covariate	Diabetic subjects			Control subjects			Total (diabetic and control subjects)		
	β	95% CI	P-value	β	95% CI	P-value	β	95% CI	P-value
Adjustment:									
Glucose	0.367	0.082-0.652	0.009	-	-	-	0.149	(-0.001)-0.299	0.040
SBP	0.271	0.041-0.502	0.030	0.334	0.076-0.592	0.010	0.239	0.063-0.415	0.004
Multiple correlation coefficient, significance of model			$r^2 = 0.139, p = 0.012$			$r^2 = 0.206, p = 0.001$			$r^2 = 0.173, p < 0.001$

Data are presented as standardized regression coefficient \pm 95% confidence interval with statistical significance, Table II – covariates with statistical significance < 0.05 are bolded, Table III – statistically significant covariates only are given, β – standardized regression coefficient, CI – confidence interval, p – value of statistical significance, BMI – body mass index, HbA_{1c} – glycated hemoglobin, CRP – C-reactive protein, WBC – white blood cells, SBP – systolic blood pressure, DBP – diastolic blood pressure, LDL-Ch – LDL cholesterol, HDL-Ch – HDL cholesterol, TG – triglyceride.

Table IV. Stepwise logistic regression analysis with variables related to presence of diabetic angiopathies in type 2 diabetic patients

Covariate	Univariate			Multivariate		
	OR (95% CI)	Wald	P-value	OR (95% CI)	Wald	P-value
CHIT1 ≥ 52.3 nmol/ml/h	0.212 (0.102–0.442)	17.13	0.001	2.58 (1.165–3.35)	6.37	0.007
BMI ≥ 25 kg/m ²	1.351 (0.512–3.562)	0.37	0.535			
Glucose ≥ 100 mg/dl	0.230 (0.111–0.476)	15.65	0.002	2.48 (1.3–3.75)	9.76	0.002
HbA _{1c} ≥ 7.5%	0.446 (0.212–0.937)	4.54	0.028			NS
WBC ≥ 10.9 × 10 ⁹ /l or CRP > 6 mg/l	0.342 (0.163–0.718)	8.03	0.004	2.96 (1.77–5.14)	16.53	0.001
SBP > 140 or DBP > 90 mm Hg	0.424 (0.205–0.879)	5.32	0.016			NS
Total-Ch ≥ 200 mg/dl	0.568 (0.292–1.108)	2.75	0.093	1.57 (1.053–3.321)	4.31	0.037
TG ≥ 150 mg/dl	2.163 (1.042–4.490)	4.28	0.032	0.46 (0.247–0.849)	6.16	0.013
Smoking	1.065 (0.498–2.236)	0.02	0.887			

Data are presented as odds ratio (±95% confidence interval) with Wald coefficient and statistical significance. OR – odds ratio, CI – confidence interval, p – value of statistical significance, BMI – body mass index, HbA_{1c} – glycated hemoglobin, CRP – C-reactive protein, WBC – white blood cells, SBP – systolic blood pressure, DBP – diastolic blood pressure, total-Ch – total cholesterol, TG – triglyceride.

activation or an inflammatory state. A stepwise multiple regression analysis revealed that plasma CHIT1 activity was independently correlated with glucose level in the total population and in diabetic patients, which may indicate its participation in the course of diabetes. We also found that CHIT1 activity was additionally associated with the SBP value in all three analyzed groups (stratified or not according to presence of diabetes). So far, a significant positive correlation of this enzyme's activity and blood pressure (but with DBP) has only been found in pre-eclamptic pregnant women [20]. Since hypertension is a known risk factor for cardiovascular diseases, we may speculate that it may play an important role in the development and/or progression of unfavorable vascular changes, which is linked with increased risk for mortality and morbidity in diabetes. Interestingly, although chitotriosidase has been proposed as an inflammatory marker, we did not find any association with CRP in diabetic patients, which was, by contrast, observed in the control group. So far, a lack of significant correlation between serum CHIT activity and CRP, as well as IL-6, has been reported in several studies concerning obese children and adolescents [21, 22]. Therefore, it may be suggested that CHIT1 activity does not reflect the chronic inflammatory state occurring in type 2 diabetes, but is connected with inflammatory episodes of other etiology.

It is well known that diabetes mellitus is associated with multifaceted metabolic disorders. High prevalence of hypertension and dyslipidemia is demonstrated among patients with type 2 diabetes. Thus, these factors could not be ignored when analyzing various metabolic interplays in the disease course. Hypertension and dyslipidemia (alongside hyperglycemia, obesity and tobacco use) are influential factors for future development of vascular diseases, which lead to failure of various organs, significantly lowering quality of life and shortening the lifespan of diabetic patients [8, 16, 23]. It is important in our study to estimate their influence on chitotriosidase activity in the plasma of type 2 diabetic patients in the context of its possible participation in the development of diabetic angiopathies. Boot *et al.* [24] firstly demonstrated an association between CHIT1 expression and lipid-laden macrophages in the atherosclerotic vessel wall, and Artieda *et al.* [7] suggested a role for CHIT1 as a marker of atherosclerotic extension and showed its predictive value in determining the risk of new cardiovascular events in a population with stable coronary heart disease during 4-year follow-up studies. In the present study we found that patients with increased CHIT1 activity had higher prevalence of diabetic angiopathies compared to those with

enzyme activity similar to the control group. We suggest that chitotriosidase may be another causal factor that contributes to the development of vascular changes in diabetes. Multiple logistic regression analysis shows a relationship between increased CHIT1 activity and occurrence of DA. After adjustment for hyperglycemia, presence of inflammation, hypertension, hypercholesterolemia and hypertriglyceridemia, CHIT1 activity remained independently correlated with the presence of diabetic angiopathies. In the constructed model, the CHIT1 odds ratio for presence of diabetic angiopathies was almost the same as for glucose, which indicates the significance of increased CHIT1 activity in the pathomechanism of DA. However, it is known that there are also other mechanisms which trigger the inflammatory response in T2D. For example, oxidative stress is considered as an important factor that promotes endothelial dysfunction and further contributes to development of vascular abnormalities. As diabetes progresses, other metabolic disturbances also develop, creating a vicious circle of metabolic decline. It is not easy to evaluate whether increased CHIT1 activity is a cause or rather a consequence of DA, but this work shows an evident relationship between these events. Unfortunately, in the present study we did not distinguish subgroups with micro- and macrovascular complications, which could explain whether the observed increase in CHIT1 activity is independent of vessel size and, indirectly, of the pathomechanism of diabetic angiopathies (which is different for micro- and macroangiopathies). Further studies are needed to establish its diagnostic value (specificity and sensitivity) in development of different types of vascular late complications.

In conclusion, chitotriosidase activity in plasma of patients with ongoing type 2 diabetes was significantly higher than in the control group, and was positively associated with hyperglycemia and hypertension (independently from other metabolic risk factors of diabetes progression) as well as with higher risk of occurrence of diabetic angiopathies, which indicates its potential significance in the course of type 2 diabetes.

Conflict of interest

The authors declare no conflict of interest.

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