Role of Betatrophin and Inflammation Markers in Type 2 Diabetes Mellitus, Prediabetes and Metabolic Syndrome

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ABSTRACT

Objective: To determine the association of betatrophin and inflammation factors in metabolic diseases such as diabetes, impaired fasting glucose, impaired glucose tolerance and metabolic syndrome.

Study Design: A cross-sectional, analytical study.

Place and Duration of Study: Firat University Medical School between April 2017 and December 2020.

Methodology: The study included 20 patients with type 2 diabetes mellitus, 20 patients with impaired fasting glucose (IFG), 20 patients with impaired glucose tolerance (IGT), 20 patients with metabolic syndrome (MetS), and a control group consisting of 20 healthy individuals. Anthropometric, fasting serum biochemical data were collected. Circulating betatrophin, and inflammation markers were measured by enzyme-linked immunosorbent assay (ELISA). The association of betatrophin, TNF-alpha, and IL-6 levels between groups were performed with One-way ANOVA and post-hoc Tukey HSD test. **Results:** Significantly higher levels of circulating betatrophin were observed in IFG, IGT, And MetS groups compared to healthy controls (p=0.017). There were significantly difference TNF- α levels in IFG, IGT, and MetS groups compared to healthy controls (p<0.001). The levels of IL-6 were significantly higher in MetS group than healthy controls (p=0.007). **Conclusion:** The circulating betatrophin and TNF- α levels were increased in MetS, IFG and IGT. IL-6 was decreased in MetS compared to the healty controls. Further studies are needed to elucidate the role of betatrophin and inflammatory parameters in the development of T2DM and prediabetic syndromes, whether betatrophin could have clinical applications in the development of new antidiabetic agents.

Key Words: Betatrophin, IL-6, TNF-alpha, Metabolic syndrome, Insulin resistance, Diabetes mellitus type 2.

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INTRODUCTION

Type 2 diabetes mellitus (T2DM) is largely related to trends in obesity and sedentary lifestyle, and its frequency is increasing day by day.¹ Betatrophin is a peptide secreted from liver and adipose tissue. It is called as RIFL, lipacin, protein 8 like angiprotein ANPTL8, and consisting of 198 amino acids, increases beta cell proliferation in mice and improves glucose tolerance.² It has been reported that betatrophin stimulates pancreatic beta cells and increases endogenous insulin secretion.

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Received: April 28, 2021; Revised: November 24, 2021; Accepted: December 29, 2021 DOI: https://doi.org/10.29271/jcpsp.2022.03.303 As a result of a series of experiments performed on mouse models by Melton and Yi,³it has been revealed that betatrophin improves glycemic control by causing pancreatic beta cell rise and beta cell mass increase.⁴ With this important property of betatrophin, previous studies argue that it can replace insulin treatment and may be a new hope for diabetes treatment by increasing the number of endogenous insulin producing cells in diabetics.^{3,5} Contrary to the effects highlighted above, some previous studies have suggested that high betatrophin level causes the development of diabetic retinopathy.⁶⁻⁸ The relationship between betatrophin and T2DM is still controversial.

Several studies have suggested that adipokines, including TNF- α , and IL-6, mediate insulin resistance, and the inflammatory response.^{9,10} Disturbances in TNF- α metabolism play a role in metabolic disorders such as obesity and insulin resistance.¹¹ Interleukin-6 (IL-6) is a proinflammatory cytokine that induces inflammation and insulin resistance.¹² A previous study reported that the inflammatory process, which is induced by IL-6, caused chronic diseases such as diabetes, obesity, cancer, and atherosclerosis.¹³

As mentioned above, there are conflicting literature data on the effect of betatrophin and inflammatory parameters on the mechanism of insulin resistance. The current study was aimed to determine the association of betatrophin and inflammation factors in metabolic diseases such as diabetes, impaired fasting glucose (IFG), impaired glucose tolerance (IGT) and metabolic syndrome (MetS).

METHODOLOGY

The study was conducted between April 2017 and December 2020 at Firat University Medical School Elazig, Turkey. The study included 20 patients with type 2 diabetes mellitus, 20 patients with IFG, 20 patients with IGT, 20 patients with MetS, and a control group consisting of 20 healthy individuals. Individuals with signs of infection, those under anti-inflamatory and steroid treatment, malignant disease, rheumatic diseases, chronic renal failure, thyroid diseases, and type 1 diabetes mellitus, were excluded.

Diabetes and prediabetes were diagnosed according to the American Diabetes Association (ADA) criteria. T2DM was diagnosed based on either the fasting plasma glucose (FPG) of \geq 126 mg/dL (7.0 mmol/L) or 2-hour PG \geq 200 mg/dL (11.1 mmol/L) during OGTT or glycated hemoglobin (HbA1c) ≥6.5% (48 mmol/mol), or in a patient with classic symptoms of hyperglycemia or hyperglycemic crisis, a random plasma glucose ≥200 mg/dL (11.1 mmol/L). IFG is defined as FPG levels from 100 to 125 mg/dL (from 5.6 to 6.9 mmol/L) and IGT as 2-hour PG during 75-g OGTT levels from 140 to 199 mg/dL (from 7.8 to 11.0 mmol/L).¹⁴ In addition to fasting and postprandial plasma glucose to evaluate glycemic control, HbA1c levels, fasting serum insulin, Cpeptide concentration, metabolic parameters to evaluate pancreatic β cell function; serum lipid (total cholesterol, HDL, LDL, VLDL cholesterol, triglyceride) levels were measured. The insulin resistance of the participants was calculated with HOMA-IR formula.

Other blood samples were taken into aprotinin tubes. The blood samples taken into these tubes were centrifuged at 3500 rpm for 5 minutes within half an hour and their serums were separated.

Statistical analysis of the data was performed using the IBM SPSS 22 statistics package programme. The Shapiro-Wilk test was used to determine whether the data showed normal distribution. Descriptive statistics of the data were expressed as mean \pm standard deviation for variables with normal distribution in continuous data and median (IQR: 25th percentile-75th percentile) for variables with non-normal distribution, the frequency for categorical variables as percentage (n [%]). Categorical variables were compared using Chi-squared tests. In comparison of more than two independent groups, One-way ANOVA and Tukey HSD tests for post-hoc analysis were used for normally distributed continuous data and Kruskal-Wallis H test with Dunn-Bonferroni post-hoc analysis were used for non-normally distributed continuous data. A value of p <0.05 was considered statistically significant.

RESULTS

This study was conducted on 100 individuals including four patient groups and the control group. In the study, 68% of the individuals were women and 32% were men. There was a significant difference in fasting glucose and postprandial glucose, triglyceride, LDL and HbA1c levels between the patient groups and the control group (p < 0.05). There was no significant difference found among serum ALT, creatinine and urea levels (p > 0.05). The groups were found to be similar in terms of gender and age (p > 0.05). A significant difference found among groups in terms of BMI. Compared to the control group, there was a significant increase in BMI in T2DM, IGT and MS groups (p < 0.001). The groups were found to be similar in terms of educational status and family history of DM (p > 0.05, Table I).

A comparison of biochemical parameters among all groups was shown in Table II. HbA1c, FBG, and PPG levels were higher in MS and T2DM groups than in the control group (p < 0.001). Triglyceride and LDL levels were higher in the MS group than in the control group (p < 0.001). PPG level was higher in the IGT group than in the control group (p < 0.001). Triglyceride and LDL levels were higher in the MS group than in the control group (p < 0.001). The IGT group than in the control group (p < 0.001, Table II).

Significantly higher levels of circulating betatrophin were observed in IFG, IGT, and MetS groups, compared to healthy controls (p=0.017). There were significant difference in TNF- α levels in IFG, IGT, and MetS groups, compared to healthy controls (p < 0.001). The levels of IL-6 were significantly higher in MetS group than healthy controls (p=0.007, Table III).

DISCUSSION

The current study revealed that IFG and IGT patients had significantly higher serum betatrophin levels than healthy controls. This result suggests that betatrophin may be a predictive marker in prediabetes and T2DM. Prediabetes is the period before diabetes mellitus. The conversion rates of these patients to diabetes are quite high. In this study, an increased betatrophin levels are observed in stages such as IFG and IGT, which are the early stages of T2DM. However, betatrophin level decreased in patients with T2DM, compared to patients with IFG and IGT. Contrary to this study, it has been shown that serum betatrophin levels are high in patients with T2DM in most of the studies conducted to date.^{6,15,16} It was thought that this situation may be due to the use of anti-insulinemic metformin and insulin injections used patients with T2DM. Most of the patients in these patient groups were using metformin, while six patients were using insulin in addition to metformin. When compared with the literature, it has been stated that metformin decreases betatrophin levels and this result supports the present study.¹⁵ Serum betatrophin levels were found to be higher in patients with metabolic syndrome than healty controls. However, serum triglyceride levels were higher in MetS patients than control group. This finding suggests that there may be an interaction between serum triglyceride level and betatrophin level. Thus, future studies should focus on this relationship.

Table I: Distribution of gender, age and BMI values by groups.

	IFG n (%)	IGT n (%)	MetS n (%)	T2DM n (%)	Control n (%)	`p
Gender						
Female	14 (70.0)	15 (75.0)	15 (75.0)	8 (40.0)	16 (80.0)	0.050
Male	6 (30.0)	5 (25.0)	5 (25.0)	12 (60.0)	4 (20.0)	
Glucosuria		·	•	•		
Yes	0 (.0)	1 (5.0)	8 (40.0)	10 (50.0)	0 (.0)	<0.001
No	20 (100.0)	19 (95.0)	12 (60.0)	10 (50.0)	20 (100.0)	
Education status	·	·		· ·		
Literate	9 (45.0)	8 (40.0)	9 (45.0)	5 (25.0)	6 (30.0)	0.548
Primary school	6 (30.0)	3 (15.0)	6 (30.0)	2 (10.0)	3 (15.0)	
Middle school	2 (10.0)	3 (15.0)	3 (15.0)	4 (20.0)	6 (30.0)	
High school	2 (10.0)	3 (15.0)	2 (10.0)	6 (30.0)	4 (20.0)	
University	1 (5.0)	3 (15.0)	0 (.0)	3 (15.0)	1 (5.0)	
Family history of DM	·	·	· ·	· ·		
Yes	7 (35.0)	8 (40.0)	10 (50.0)	11 (55.0)	4 (20.0)	0.181
No	13 (65.0)	12 (60.0)	10 (50.0)	9 (45.0)	16 (80.0)	
Age (yıl). mean±sd	51.9±6.5	50.2±9.0	56.1±3.1	52.3±6.2	51.3±6.0	0.056**
BMI(kg/m ²) mean±sd	25.2±1.7	27.6±1.5°	34.9±3.4°	26.2±1.4°	23.5±1.0	< 0.001

Values are given as mean ± standard deviation and number (%). * Chi-square test; ** One-Way ANOVA test; °Compared to the control group, p<0.05, IFG: Impaired fasting glucose, IGT: Impaired glucose tolerance, MetS: Metabolic syndrome, T2DM: Type 2 diabetes mellitus, BMI: Body mass index.

Table II: Distribution of biochemical values by groups.

	IFG	IGT	MetS	T2DM	Control	n*	
	median±sd	median±sd	median±sd	median±sd	median±sd	þ	
HbA1c (%)	5.4±0.5	5.5±0.5	8.1±1.6°	8.7±1.7ª	5.1±0.4	<0.001	
FBG (mg/dL)	112.3±6.4	98.6±8.7	166.0±56.2°	151.6±46.5°	86.4±6.9	<0.001	
PPG (mg/dL)	133.2±19.2	166.4±14.1ª	258.2±71.2°	261.7±77.6°	98.4±11.3	<0.001	
Triglyceride (mg/dL)	172.7±167.6	140.7±60.9	222.4±101.9°	187.5±75.9	121.6±52.1	0.021	
LDL (mg/dL)	100.7±30.6	132.5±37.0	155.5±35.6°	131.1±26.0	109.0±27.6	<0.001	
ALT (U/L)	25.0±20.5	20.3±11.0	22.7±8.7	24.9±10.2	19.6±6.4	0.515	
Creatinine (mg/dL)	0.8±0.1	0.7±0.2	0.8±0.3	0.8±0.2	0.7±0.1	0.084	
Urea (mg/dL)	32.6±9.7	33.6±10.9	34.0±8.4	35.3±11.4	26.9±5.6	0.055	
$^{\circ}$ One-Way ANOVA test; $^{\circ}$ Compared to the control group, p<0.05; IFG; Impaired fasting glucose, IGT; Impaired glucose tolerance, MetS; metabolic							

One-Way ANOVA test; "Compared to the control group, p<0.05; IFG: Impaired fasting glucose, IGT: Impaired glucose tolerance, MetS: metabolic syndrome, T2DM: Type 2 diabetes mellitus, PPG: Post-prandial glucose.

	IFG median (IQR)	IGT median (IQR)	MetS median (IQR)	T2DM median (IQR)	Control median (IQR)	p*	
Betatrophin (ng/L)	334.4 (298.9-360.5) ^a	334.7 (292.9-357.9) ^a	295.6 (269.4-320.3)ª	320.1 (303.4-348.3)	292.4 (263.9-312.0)	0.017	
TNF-α (ng/ml)	1347.4 (363.9-11344.6) ^a	6873.9 (3644.1-8254.5) ^a	6621.6 (4351.4-7450.5) ^a	200.0 (122.2-1323.6)	1055.6 (731.9-3726.3)	<0.001	
IL-6 (ng/L)	202.9 (27.6-529.5)	312.4 (188.1-437.5)	517.6 (417.5-683.3)ª	373.5 (32.0-691.1)	248.4 (54.0-402.7)	0.007	
^a Compared to the control group, p<0.05; *Kruskal-Wallis H test; IFG: Impaired fasting glucose, IGT: Impaired glucose tolerance, MetS: Metabolic syndrome, T2DM: Type 2 diabetes mellitus. TNF-a: Tumor necrosis factor-alpha, IL-6: Interleukin-6.							

Table III: Distribution of betatrophin, TNF- α and IL-6 values by groups.

The current study showed higher BMI levels of patients with MetS and T2DM than of healthy controls. This finding is expected and consistent with literature.^{5,10} Increased serum betatrophin levels in the prediabetic period suggest that betatrophin may have a role in the pathogenesis of T2DM. Novel studies are needed to elucidate the possible role of betatrophin in the pathogenesis of T2DM.

Pro-inflammatory mediators play a crucial role to induce insulin resistance and T2DM *via* involvement of oxidative stress and the activation of various transcriptional mediated molecular and/or metabolic pathways. Experimental studies have found that pro-inflammatory mediators are interdependently involved to induce tissue-specific inflammation, which may lead to the development of insulin resistance and pathogenesis of T2DM.¹⁷ The current study showed that TNF- α levels were higher in IFG, IGT, and MetS patients than healthy controls. The previous studies showed that there is low-grade inflammation in the prediabetes stage.¹⁸ The current study data suggested that this low-grade inflammation plays a role in the pathogenetic processes that cause T2DM.

The current study was found higher IL-6 level in MetS patients than healthy controls. Several studies have suggested that TNF- α , and IL-6, mediate insulin resistance, and the inflammatory response.^{19,20} The previous study reported that high levels of IL-6 have been observed in adipose tissues from patients with diabetes mellitus or obesity, particularly those with features of MetS.²¹ According to the obtained results in this study, the authors have thought that changes in TNF- α and IL-6 levels are a result of diabetes formation stages rather than a causal factor in the transformation of prediabetes stages to T2DM. Insulin resis-

tance (IR) and b-cell dysfunction include the underlying pathophysiology of prediabetes and its subsequent progression to overt diabetes. Recent studies show that leptin, resistin, IL-6 and TNF- α play an important role in the emergence of insulin resistance.²¹ In this study, it was found that TNF- α and IL-6 levels were statistically significantly higher in all stages of prediabetes (BAG, IGT and MetS), compared to the control group. It is seen in the literature that chronically elevated levels of inflammatory biomarkers such as TNF- α and IL-6 directly cause insulin resistance and impairment of insulin sensitivity and increase the risk of T2DM.¹⁹⁻²¹ The present findings suggest that these cytokines could be used for early diagnosis of diabetes in the future. Because IL-6 levels were found to be statistically significantly increased in IFG, IGT and MetS patients.

The current study found that elevated betatrophin levels were associated with T2DM and higher BMI. This relationship of betatrophin and T2DM has been previously reported by Fu et al.¹⁶ Recent reports have shown that a betatrophin transcript variant and its expression levels correlate with clinical or pathological symptoms. The genetic R59W variant of betatrophine has been shown to be associated with the reduction of HDL and LDL in African-American and Spanish participants, and higher levels of circulating betatrophin protein were found in overweight / obese groups in type 2 diabetes patients.²² A previous study reported that serum betatrophin levels increased significantly in prediabetic stages (IFG-IGT) and T2DM.²³ These findings were thought that serum betatrophin levels are associated with glucose intolerance. Similar findings by Liu et al.,²⁴ reported that serum betatrophin levels were higher in T2DM and IGT patients than healthy controls, and it was found to be closely related to insulin resistance. The present findings of Liu et al. also support the above-mentioned hypothesis.

Chen *et al.*²³ found that betatrophin levels increased in patients with T2DM. It has also been observed that betatrophine affects the lipid profile and insulin resistance states. There is a significant positive relationship between HbA1c and triglyceride levels in type 2 diabetic patients.²⁵ Proinflammatory cytokine levels and betatrophin and HbA1c levels are found to be related in a similar way with MetS in the T2DM group, and dyslipidemic values with increased circulating levels of IL-6 and TNF- α play a role in a cellular balance in prediabetic stages.

The major limitations of this study included its single-centre design, low number of patients, so the current study may face us to can not generalize to other settings in the rest of the country or across the world. Here, the lack of VLDL, total cholesterol, and insulin values in the patient groups in this study was also seen as one of the reasons limiting the results of this study.

CONCLUSION

The circulating levels of betatrophin and TNF- α were increased in MetS, IFG and IGT. IL-6 was decreased in MetS compared to the healty controls. Further studies are needed to elucidate the role of betatrophin and inflammatory parameters in the development of T2DM and prediabetic syndromes whether betatrophin could have clinical applications in the development of new antidiabetic drugs.

ETHICAL APPROVAL:

The study protocol was approved by the Ethics Committee of Firat University (No. 05-16, Date: 16/03/2017).

PATIENTS' CONSENT:

Written informed consents were obtained from all patients. The study was conducted in accordance with the principles of the Declaration of Helsinki.

CONFLICT OF INTEREST:

The authors declared no conflict of interest.

AUTHORS' CONTRIBUTION:

EO, AB: Data collection, design, compiling, and discussion. Led and conceived the project, and authored the manuscript. BY: Collecting and analysing data, discussion.

ED, MFG: Collecting statistics and analysing data.

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