Serum Nesfatin-1 in patients with type 2 diabetes mellitus: A cross sectional study

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Abstract

Background and objective: Nesfatin-1 is a newly described peptide, derived from nucleobindin2. Nesfatin-1 suppresses food intake and it is involved in regulating insulin secretion. The aim of this study was to compare serum levels of Nesfatin-1 in patients having type 2 diabetes and non-diabetic subjects.

Methods: This cross-sectional study included 90 participants; 64 patients with type 2 diabetes mellitus (32 males and 32 females) and 26 control subjects (13 males and 13 female). Body mass index, fasting serum level of glucose, fasting serum level of insulin, and glycated hemoglobin were estimated. Homeostasis model assessment of insulin resistance was calculated. Nesfatin-1 level was measured using enzyme-linked immune-sorbent assay kit. The data was analyzed using Graphpad prism 7.04 for windows.

Results: Type 2 diabetes patients aged from 33-78 years and the control group aged from 32-75 years. Nesfatin-1 level in the diabetic group was significantly lower than controls. The median interquartile range (IQR) of Nesfatin-1 was 0.765 (0.4-1.173) in diabetes and 1.02 (0.775-1.458) in controls. The diabetes group has significantly higher homeostasis model assessment of insulin resistance compared with non-diabetics. Serum Nesfatin-1 was correlated negatively with body mass index, fasting serum glucose, fasting serum insulin, glycatedhemoglobin, and homeostasis assessment of insulin resistance.

Conclusion: Serum Nesfatin-1 level is negatively correlated with fasting serum glucose, fasting serum insulin, and glycated hemoglobin. This association supports the role of Nesfatin-1 in increased insulin resistance in patients with type 2 diabetes.

Keywords: Type 2 diabetes mellitus; Nesfatin-1; Homeostasis model assessment of insulin resistance; Glycated hemoglobin.

Introduction

Nesfatin-1 (NES-1) is a newly discovered multi-functional peptide hormone with an approximate molecular weight of 9.8 kilo Dalton and a half-life of 23.5 minutes. It was first discovered in 2006 by Oh-I and his coworkers. 1-3 NES-1 is derived from nucleobindin2 (NUCB2) precursor, which is DNA and calcium binding protein that is found in the plasma membrane and neuroplasma. NUCB2 is highly conserved in humans, rats, and mice, that shares more than 85% of homology between humans and the other mammal species and even demonstrates similarities with lower organisms.3-5 The NUCB2

possibly precursor protein is translationally cleaved by the enzyme prohormone-convertase into the N-terminal NES-1amino acids 1-82 (AA 1-82), NES-2 (AA 85-163) and the C-terminal NES-3 (AA 166-396). Several biological actions been NES-1 have identified. specifically of the middle part of it, which corresponds to AA 24-53, it has a key role in physiological effects of NES-1, particularly for an anorexic effect, whereas no biological action has been described for NES-2 and NES-3.6,7 NUCB2 and NES-1 are expressed by the central nervous system (CNS) and peripheral tissues. In CNS, expressed prominently in the

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Hypothalamus, particularly in the supraoptic and paraventricular nuclei, spinal cord autonomic nuclei, pituitary gland, brain stem nuclei, nucleus tractus-solitarus, and in peripheral tissues including gastric mucosa, adipocytes, pancreatic endocrine beta cells (β-cells), and testis tissue. It has been found that its expression level is 20-fold higher in the gastric oxyntic mucosa than in the brain.^{8,9} It has been reported that NES-1 suppresses food intake and reduces weight gain when injected into the third ventricle of rodents or it administered peripherally. So, the major function of NES-1 is inhibition of food intake in time-dependent, dose-dependent, insulin-dependent manners. 10,11 Detection of NES-1 localizes with insulin in the beta islets pancreatic indicates involvement of NES-1 in regulating insulin secretion from pancreatic beta-cells and thus regulation of blood glucose. NES-1 affects glucose metabolism, the suggested mechanism for that is by increasing and insulin sensitivity decreasing insulin resistance. For measuring insulin resistance, homeostatic model assessment resistance (HOMA-IR) used.^{5,9,15-17} Type 2 diabetes mellitus (T2DM) is an important expanding global health problem. It is associated with alteration in glucose metabolism, insulin resistance, abnormality in fasting serum glucose (FSG), and also impaired glucose tolerance. 18,19 Factors such as obesity, unhealthy diet, low physical activity, and genetic factors contribute to some pathophysiological disturbances that are responsible for impaired glucose homeostasis in T2DM. Impaired insulin secretion and insulin resistance are still the core defects in T2DM.20 This study aimed to find out the level of serum NES-1 in patients having T2DM and to compare it with a serum level of NES-1 in normal subjects. It also aimed to find out the relation between levels of NES-1 and T2DM and detect the relation between NES-1, insulin level, and insulin resistance.

Methods

Subjects

This cross-sectional study was conducted at Layla Qasim Diabetic center in Erbil city in January and February 2018. The study included 90 participants: 64 patients with T2DM and 26 subjects free from diabetes mellitus comprised the control group. The diagnoses of Type 2 DM were made at least before one year by a physician according to WHO criteria. After verbal obtained from consent was participants, information was obtained from them using a questionnaire designed for the study. Body weight and height were measured in all subjects using a scale (Seca-USA) and a wall-mounted stadiometer to the nearest 0.5 kg and 0.5 cm, respectively. Body mass index (BMI) was calculated as the body weight in kilograms divided by the height in meters squared. The subjects in the control group had normal glucose levels (no diabetes history). Clinical exclusion criteria for cases and controls were Type 1 DM, pregnancy, hormone replacement therapy, cancer, severe acute or chronic infectious disease, systemic diseases (heart failure, kidney, liver, or lung diseases), thyroid, and adrenal disorders. All procedures were conducted in accordance with The Code of Ethics of the World Medical Association (Declaration of Helsinki) for experiments involving humans.²¹ The Local Ethics Committee at the College of Medicine, Hawler Medical University approved the study protocol.

Biochemical parameters and hormone analyses

Blood samples (5 ml) were drawn from a forearm vein in the morning after overnight fasting for hormone and other biochemical analyses. About 2.5 ml of the blood sample was placed into vacuum blood collection tube and then centrifuged at 4500 rpm for 5 minutes to obtain serum, some of the separated serum was used to estimate FSG and some for fasting serum insulin (FSI) estimation by using an automated immunoassay analyzer

(Cobas e411, Roche Diagnostics GmbH, D-68298 Mannheim, Germany), while the rest of the serum was immediately placed in Eppendorf tubes and frozen at -40 $^{\circ}$ C to be used later for the analysis of NES-1. The other 2.5 ml of the blood sample was placed into tubes containing Ethyle Diamine Tetra Acetic Acid (EDTA) for direct analysis of glycated hemoglobin (HbA_{1C)} using (GesanChem 200, Campobello di Mazara- Italy). The HOMA-IR calculated using the following equation:

 $HOMA- IR = FSI (mIU/L) \times FSG (mg/dI)/405.$

Serum NES-1 assay

Fasting serum NES-1 level was measured using a commercial enzyme-linked immune -sorbent assay (ELISA) kit (Bioassay Laboratory, Technology YangpuDist, Shanghai, China), by Absorbance Microplate Reader (ELx800™, Operator's Manual , Biokit, Wefen com. Spain) The standard curve range was 0.3-90 ng/ml. The coefficients of variation for inter-assay and the intra-assay were 10% and 8%, respectively.

Statistical analysis

data was analyzed using Graphpad Prism 7.04 for windows. All data were expressed as mean±standard error of the mean (SE) or median IQR. Statistical analysis was used according to the (t-test for two independent samples), and the Mann-Whitney test applied for data, which was not normally distributed. Data accuracy for diagnosis of T2DM was presented in terms of sensitivity and specificity by the Receiver Operating Characteristic (ROC) curve, which is a graphical display of sensitivity on the y-axis and (1-specificity) on the x-axis for varying cut-off points of test values. The area under the curve (AUC) (normal range 0.5-1) is used for measuring the quantitative accuracy. An area of 0.5 represents a worthless test, while an area of 1 represents a perfect test. Rough guide for classifying the accuracy of a diagnostic test is the traditional academic point system: 0.90-1 = excellent, 0.80-0.90 =

good, 0.70-0.80 = fair, 0.60-0.70 = poor, 0.50-0.60= fail. Correlations between data variables showed by Spearman correlation coefficient (8). A P value of less than 0.05 was considered as statistical significant.

Results

The study included 90 participants: 64 patients with T2DM (32 males; 32 females) aged from 33 to 78 years and 26 subjects free from DM comprised the control group (13 males; 13 female) aged from 32 to 75. The mean BMI of T2DM cases and controls were 30.21±0.692 and 27.76±0.952, respectively.

Comparison of biochemical parameters between T2DM and non-diabetic groups As shown in Table 1 and Figure 1, the fasting serum NES-1 level was significantly (P = 0.042) lower in patients with T2DM in comparison with the control group and the median IQR was 0.765 (0.4-1.173) ng/ml and 1.02 (0.775-1.458) ng/ml, respectively. The FSG, FSI, HbA_{1C} and the calculated HOMA-IR levels were significantly higher in patients with T2DM compared to the control group. At the same time, the calculated BMI was also higher in the T2DM group, but the difference was not significant statistically. The results were expressed as Mean±SEM, except for serum NES-1, FSI, HOMA-IR levels in which they were expressed as median IQR because their data was not normally distributed (Table 1).

Table 1: Comparison between the groups studied according to laboratory investigations fasting serum levels of NES-1, glucose and insulin, and calculated HOMA-IR, calculated BMI and HbA $_{1C}$ between the study groups.

Parameters	Control (n=26)	T2DM (n=64)	P values
NES-1(ng/ml): Median IQR	1.02 (0.775-1.458)	0.765(0.4-1.173)	0.042
FSI (mIU/L): Median IQR	11.21(7.768-14.44)	15.44(9.42-24.77)	0.033
HOMA-IR: Median IQR	2.39(1.823-3.2)	5.15(2.35-9.875)	<0.001
FSG (mg/dl)	88.96±1.659	165.9 ± 10.42	<0.001
BMI (kg/m²)	27.76±0.952	30.21±0.692	0.051
HbA1C (%)	5.038 <u>±</u> 0.151	8.029±0.218	<0.001

(IQR)= Interquartile range

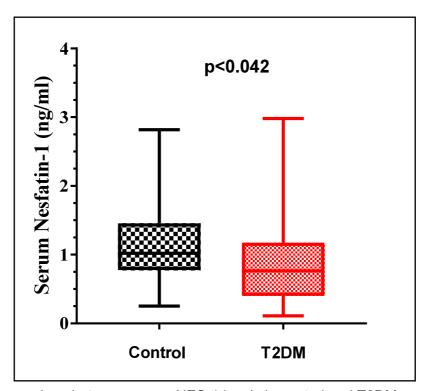


Figure 1: Comparison between serum NES-1 levels in control and T2DM groups.

Correlation coefficients (r) between data variables

Table 2 shows the correlations between data variables by correlation coefficient (r) and *P* values. NES-1 showed no significant negligible negative correlation with BMI, FSG, HbA1C, and FSI. However, it has a significant negligible negative correlation with HOMA-IR (Figure 2). On the other

hand, our results showed that BMI had a significant low positive correlation with insulin and HOMA-IR and a significant negligible positive correlation with HbA_{1C}. Moreover, the ROC curve of NES-1 shows discrimination between T2DM and control subjects, with an AUC of 0.64 and P = 0.042, as seen in Figure 3.

Table 2: Correlations between data variables in all subjects represented by correlation coefficient (r) and *P* values (p):

		NES-1	ВМІ	FBG	HbA1C	Insulin	HOMA-IR
NES-1	R		- 0.011	- 0.216	- 0.07	-0.05	-0.221
	р		0.919	0.057	0.511	0.665	0.042
BMI	R	- 0.011		0.188	0.214	0.42	0.416
	р	0.919		0.099	0.043	0.004	0.005
FSG	R	-0.216	0.188		0.867	0.206	0.628
	р	0.057	0.099		<0.001	0.071	0.009
HbA1C	R	-0.07	0.214	0.867		0.094	0.405
	р	0.511	0.043	<0.001		0.409	0.004
Insulin	R	-0.05	0.42	0.206	0.094		0.841
	р	0.665	0.004	0.071	0.409		<0.001
HOMA-IR	R	-0.221	0.416	0.628	0.405	0.841	
	р	0.042	0.005	0.009	0.004	<0.001	

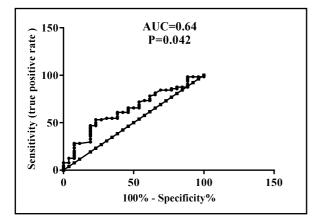


Figure 2: Correlation coefficient between NES-1 and HOMA-IR.

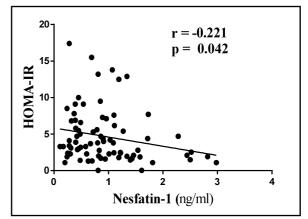


Figure 3: The ROC curve shows the sensitivity and specificity of NES-1 concentration during T2DM. AUC: Area under the curve

Discussion

In the present study, we compared fasting serum NES-1 levels in patients having T2DM with non-diabetic persons. We found NES-1 was significantly that level (P = 0.0425) lower in patients with T2DM compared to non-diabetic subjects. This result is in agreement with documented results in some previous studies. 19,24 This finding suggests that low level of serum NES-1 in T2BM patients might be one of the factors causing diabetic hyperphagia because it has been reported that NES-1 has a reducing effect on food intake. Thus, it acts as an anorexigenic agent, but the regulatory mechanism remains not clear. On the other hand, the ROC curve analysis showed that NES-1 represents as a biomarker for T2DM, with an AUC of 0.64 and P = 0.0042. The exact mechanism by which NES-1affect T2DM is not fully understood yet, however, some researches support our finding. 19, 24 The results of our study show increased FSG, HbA_{1C}, FSI, and HOMA-IR in T2DM as compared with the non-diabetic group. Calculating of data according to HOMA-IR formula shows that there is a significant difference (P < 0.001) in insulin resistance between T2DM and control groups, so; lower NES-1 levels in T2DM group could explain the significantly high insulin resistance in this group because as mentioned before NES-1 suggested to affect glucose metabolism increasing insulin sensitivity and resistance.5,9,17 insulin decreasing Furthermore, our results demonstrated that serum NES-1 level is negatively correlated with all the FSG, HbA_{1C}, FSI, HOMA-IR, and BMI. These negative correlations can be explained by the role of NES-1 in glucose metabolism and its anorexigenic effect, and this finding agreed with the results of a study done by Tsuchiya and his coworkers.²⁵ An important point to be mentioned is that in our study, the least duration of diagnosis was one year, and newly diagnosed T2DM patients were not included, in which most of them were on oral hypoglycemic drugs. This made the

difference with another study done by Zhang and his colleagues (2012) that included newly diagnosed T2DM patients in which the level of NES-1 was higher in T2DM group compared with the control group and the author explained this by the possibility that β -cells secrete normal amounts of insulin and NES-1 during the early stages of diabetes only. Hence, they designed their study based on excluding effects of microvascular and macrovascular complications and multiple medications on plasma NES-1.4 Recent studies reported anti-inflammatory effects for NES-1, and the explained mechanism is that there are elevated circulating NES-1 levels in newly diagnosed T2DM patients, as discovered by Zhang and his coworkers, and this elevation of NES-1 can reduce blood glucose, decrease food intake, increasing lipid metabolism, and countering inflammation.4,26 Accordingly, we hypothesized that in newly diagnosed T2DM there will be a compensatory mechanism that leads to the elevation of NES-1 level. In patients with T2DM who are receiving anti-diabetic treatment, circulating NES-1 levels decreased, by the of action of anti-diabetic treatments, as these drugs causing the reduction of blood glucose, increased insulin sensitivity, and additionally controlled food intake.²⁷ For the future, we recommend more studies concerning the NES-1 levels in response to diabetic therapy, including the effects of insulin and oral anti-diabetic treatment is needed. To elucidate the physiological role of NES-1, further studies are required. Establishing the physiological and biochemical effects of NES-1 on the regulatory mechanisms of energy systems in the body will contribute to clarify the factors that could play a role in the impairment of glucose metabolism.

Conclusion

Serum Nesfatin-1 levels are negatively correlated with fasting serum glucose, fasting serum insulin, and glycated hemoglobin. This association supports the

role of Nesfatin-1 in increased insulin resistance in patients with type 2 diabetes mellitus.

Competing interests

The authors declare no competing interests.

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