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

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 females). 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. Nesfatin-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. The NUCB2 precursor protein is possibly post-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 for NES-1 have been 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. NUCB2 and NES-1 are expressed by the central nervous system (CNS) and peripheral tissues. In CNS, expressed prominently in the...
Hypothalamus, particularly in the supra-optic and paraventricular nuclei, spinal cord autonomic nuclei, pituitary gland, brain stem nuclei, nucleus tractus-solitarius, 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, and insulin-dependent manners.\(^10,11\)

Detection of NES-1 localizes with insulin in the pancreatic beta islets indicates the 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 insulin sensitivity and decreasing insulin resistance. For measuring insulin resistance, homeostatic model assessment of insulin resistance (HOMA-IR) is 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 consent was obtained from the 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.
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(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 °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 (HbA1C) using (GesanChem 200, Campobello di Mazara- Italy). The HOMA-IR calculated using the following equation:

\[ \text{HOMA-IR} = \frac{\text{FSI} \times \text{FSG}}{405} \]

Serum NES-1 assay

Fasting serum NES-1 level was measured using a commercial enzyme-linked immune-sorbent assay (ELISA) kit (Bioassay Technology Laboratory, 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

The data was analyzed using the 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. 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, HbA1C 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\textsubscript{1C} between the study groups.

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Control (n=26)</th>
<th>T2DM (n=64)</th>
<th>P values</th>
</tr>
</thead>
<tbody>
<tr>
<td>NES-1 (ng/ml): Median IQR</td>
<td>1.02 (0.775-1.458)</td>
<td>0.765(0.4-1.173)</td>
<td>0.042</td>
</tr>
<tr>
<td>FSI (mIU/L): Median IQR</td>
<td>11.21(7.768-14.44)</td>
<td>15.44(9.42-24.77)</td>
<td>0.033</td>
</tr>
<tr>
<td>HOMA-IR: Median IQR</td>
<td>2.39(1.823-3.2)</td>
<td>5.15(2.35-9.875)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>FSG (mg/dl)</td>
<td>88.96±1.659</td>
<td>165.9 ± 10.42</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>BMI (kg/m\textsuperscript{2})</td>
<td>27.76±0.952</td>
<td>30.21±0.692</td>
<td>0.051</td>
</tr>
<tr>
<td>HbA1C (%)</td>
<td>5.038±0.151</td>
<td>8.029±0.218</td>
<td>&lt;0.001</td>
</tr>
</tbody>
</table>

(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 HbA1C. 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):

<table>
<thead>
<tr>
<th></th>
<th>NES-1</th>
<th>BMI</th>
<th>FBG</th>
<th>HbA1C</th>
<th>Insulin</th>
<th>HOMA-IR</th>
</tr>
</thead>
<tbody>
<tr>
<td>NES-1</td>
<td>R</td>
<td></td>
<td></td>
<td>-0.216</td>
<td>-0.07</td>
<td>-0.05</td>
</tr>
<tr>
<td>p</td>
<td></td>
<td>0.919</td>
<td>0.057</td>
<td>0.511</td>
<td>0.665</td>
<td>0.042</td>
</tr>
<tr>
<td>BMI</td>
<td>-0.011</td>
<td>R</td>
<td></td>
<td>0.188</td>
<td>0.214</td>
<td>0.42</td>
</tr>
<tr>
<td>p</td>
<td>0.919</td>
<td></td>
<td></td>
<td>0.099</td>
<td>0.043</td>
<td>0.004</td>
</tr>
<tr>
<td>FSG</td>
<td>-0.216</td>
<td>0.188</td>
<td>R</td>
<td></td>
<td>0.867</td>
<td>0.206</td>
</tr>
<tr>
<td>p</td>
<td>0.057</td>
<td>0.099</td>
<td></td>
<td></td>
<td>&lt;0.001</td>
<td>0.071</td>
</tr>
<tr>
<td>HbA1C</td>
<td>-0.07</td>
<td>0.214</td>
<td>0.867</td>
<td>R</td>
<td></td>
<td>0.094</td>
</tr>
<tr>
<td>p</td>
<td>0.511</td>
<td>0.043</td>
<td>&lt;0.001</td>
<td>R</td>
<td></td>
<td>0.409</td>
</tr>
<tr>
<td>Insulin</td>
<td>-0.05</td>
<td>0.42</td>
<td>0.206</td>
<td>0.094</td>
<td>R</td>
<td></td>
</tr>
<tr>
<td>p</td>
<td>0.665</td>
<td>0.004</td>
<td>0.071</td>
<td>0.409</td>
<td>&lt;0.001</td>
<td>R</td>
</tr>
<tr>
<td>HOMA-IR</td>
<td>-0.221</td>
<td>0.416</td>
<td>0.628</td>
<td>0.405</td>
<td>0.841</td>
<td>R</td>
</tr>
<tr>
<td>p</td>
<td>0.042</td>
<td>0.005</td>
<td>0.009</td>
<td>0.004</td>
<td>&lt;0.001</td>
<td></td>
</tr>
</tbody>
</table>

**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.
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.

**References**