

Study of Serum Spexin Level in Obese and Non-Obese type 2 Diabetic Patient

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Abstract

Background: Spexin is a peptide, which is discovered as an important regulatory adipokine in obesity, insulin resistance (IR), and diabetes. **Aim:** To evaluate serum Spexin levels in type-2 diabetic obese patients in comparison to its levels in non-obese diabetic (NOD) and to assess its relationship to IR and glycemic control. **Patients and methods:** A total of 20 NOD (group 1), 22 obese non-diabetic (OND) (group 2), and 24 obese diabetic patients (OD) (group 3), with an average age of 43.8 ± 8.9 , 49 ± 8.8 , 45.1 ± 11.6 years respectively, and 20 age and sex-matched healthy participants (control group) were included. Height and weight were measured using standard techniques. Glucose levels, triglycerides (TG), total cholesterol (TC), high-density lipoprotein cholesterol (HDL-C), low-density lipoprotein cholesterol (LDL-C), fasting insulin (FI), calculation of Homeostasis model assessment insulin resistance (HOMA-IR), and Spexin levels were measured in each patient. **Results:** The median serum Spexin levels were significantly lower in the OD than in the other groups and lower in the NOD and OND than in the non-obese non-diabetic (NOND) ($p < 0.001$). Spexin level at a cutoff value ≤ 0.70 can significantly discriminate OD from NOND. Spexin levels were negatively correlated with age, body mass index (BMI), LDL-C, TG, FI, and HOMA-IR. **Conclusion:** Spexin plays an important role in glucose homeostasis and lipid metabolism. The presence of diabetes is associated with lower Spexin levels. Further investigations and additional studies in larger populations are required to understand the exact function of this peptide and to validate the observations in the current study.

Keywords: Obesity, diabetes, adipokine, insulin resistance

Introduction

Diabetes Mellitus (DM) is a metabolic chronic disorder, which is characterized by chronic elevation of the blood glucose level. In 2015, nearly 415 million adult subjects with ages ranging from 20 to 79 years suffered from DM⁽¹⁾. It is a global public health burden as this number is expected

to rise to another 200 million by 2040. Approximately 1 in 11 adults all over the world suffer from DM, 90% of them are Type II DM (T2DM)⁽²⁾. The major factors contributing to the increasing rates of T2DM are the increase in a sedentary lifestyle, obesity, and aging of the population⁽³⁾. High levels of adiposity, evaluated by increased BMI, are the single main risk factor for the

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development of T2DM⁽⁴⁾ and are related to several metabolic abnormal defects which lead to IR⁽⁵⁾. Obesity management by using an operation for weight loss has been demonstrated to be successful to prevent and treat cases suffering from T2DM⁽⁶⁾. Spexin is a neuropeptide generated in the white fatty tissue, brain, heart, thyroid, lung, ovary, liver, adrenal, testis, pancreas, muscle, and stomach⁽⁷⁾. It was recognized as a significant regulator of obesity and its associated metabolic disorders including IR and DM⁽⁶⁾. Preliminary studies in mice with diet-induced obesity, T2DM, and hepatic steatosis suggest that the administration of Spexin may be an effective treatment for these three conditions⁽⁸⁾. Spexin role is not well recognized; and the strict mechanisms throughout which Spexin applies its effects are unclear even now, due to insufficient information available on the receptors of Spexin⁽⁹⁾. Walewski et al. found a considerably lower concentration of Spexin in obese than in lean adults and suggested a possible satiety-inducing function for it in humans⁽¹⁰⁾. Decreased levels of Spexin have been detected in T2DM patients compared to the non-diabetic people, which recommends that Spexin may affect the metabolism of lipid and glucose⁽¹¹⁾. Insulin and Spexin localization in β cells were established in pigs and humans, but not in mice⁽¹²⁾. This denotes that Spexin possibly could affect the function of the pancreas. The present study aimed to: evaluate serum Spexin levels in type-2 diabetic obese patients in comparison to its levels in NOD and to assess its relationship to IR and glycemic control

Subjects and Methods

This case-control study included 44 patients with T2DM (24 OD, 20 NOD) and 22 OND, with a mean age of 45.1 ± 11.6 ,

43.8 ± 8.9 , and 49 ± 8.8 years respectively, selected from the Outpatient Clinics of Mansoura University Hospital, Faculty of Medicine, Egypt, from March 2018 to February 2019. Moreover, 20 healthy controls matched for sex and age were also selected from the same locality. Informed consent was obtained from all participants, and approval was given by the ethics committee of our institution. DM was diagnosed according to the American Diabetes Association (ADA) criteria⁽¹³⁾. We excluded patients who had Type 1 DM, lean patients, patients with intrinsic renal or hepatic diseases, patients with organ failure, and patients who had an endocrinal cause of obesity or diabetes. Full history and clinical examination were taken from all participants. The patient's height and weight were measured using standard techniques. BMI was calculated as weight in kilograms divided by height in meters squared (kg/m^2). Blood pressure (BP) was taken in the sitting position using a random-zero sphygmomanometer. Venous blood samples after overnight fasting were collected. Creatinine level in the blood serum was determined. Serum TG levels were estimated according to the Allain et al. method⁽¹⁴⁾. TC was estimated according to the Fossati and Prencipe⁽¹⁵⁾ method. Serum HDL-C was estimated according to the Williams et al. method⁽¹⁶⁾. LDL-C levels in blood serum were calculated according to the equation, which was described by Trinder⁽¹⁷⁾: $\text{LDL-C} = \text{TC} - \text{HDL-C} - \text{TG} / 5$. Estimation of blood glucose was performed according to the Trivedi et al. method⁽¹⁸⁾. Serum insulin was assayed by ELISA supplied by Eagle Biosciences, Inc. Catalog Number: INS31-K0120A (Northwest Blvd., Suite 112, Nashua, NH 03063). Calculation of HOMA-IR. $\text{*HOMA} = \text{Fasting glucose (mg/dl)} \times \text{FI (ml u/ml)} / 405$ (19). Spexin was assayed by ELISA kits supplied by Mybiosource (USA). This kit

utilizes ELISA depending on the Avidin-Biotin complex technology. Addition of Spexin (C12orf39) to the wells that are coated with monoclonal antibody of Spexin (C12orf39) and after that undergo incubation for a certain period. Then, putting anti C12orf39 antibodies that are labeled by biotin for uniting with streptavidin-HRP leads to the formation of the immunocomplex. Washing to get rid of the excess unbound enzymes after the termination of the incubation period. In addition to substrate A and B, then the solution will become blue and turn into yellow color with the acid effect.

Statistical analysis

Data were fed to the computer and ana-

lyzed using IBM SPSS software package version 22.0. Qualitative data were described using numbers and percentages. Quantitative data were described using the mean. The standard deviation for parametric data after testing normality using Kolmogorov-Smirnov test, and median (min-max) for non-parametric data. The significance of the obtained results was judged at the (0.05) level.

Results

Table 1 shows an insignificant difference between groups regarding age ($P=0.3656$) Female sex was higher in OD ($n=23$) than in the NOND ($n=12$) and the NOD ($n=13$) ($p=0.011$). TC was significantly higher in the OD than in the control group,

Table 1: Clinical and biochemical characteristics of the study groups

Variable	NOND (n= 20)	NOD (n= 20)	OND (n= 22)	OD (n= 24)	P value
Sex (n%)					
Males	8(40%)	7 (35%)	3 (13.6%)	1 (4.2%)	0.011
Females	12(60%)	13 (65%)	19 (86.4%)	23 (95.8%)	
Age (years)*	45.63±8.6	43.8±8.9	4.9±8.8	45.1±11.6	0.3656
SBP (mmHg)*	118.5±13.9	126.5±15.3	125.9±12.6	134.6±18.9	0.011
DBP (mmHg)*	73±8.6	76±13.1	77.7±10.2	77.5±11.1	0.479
TC*	232.8±63.7	254.1±79.0	267.8±95.7	311.6±124.2	0.049
LDL-C*	130±58.2	135±79.1	167±94.1	188.7±113.7	0.109
HDL-C*	65.5±15.9	64.0±17.5	60.5±16.1	62.7±19.3	0.818
TG*	187.9±98.6	277.7±112.2	202.7±90.5	303.3±107.4	<0.001
Fasting insulin@	2.4 (1.4-3.9)	4.3 (3.5-5.1)	4.9 (3.9-6.0)	7.15 (5.90-11.75)	<0.001
HOMA-IR@	0.54 (0.28-0.82)	2.49 (1.37-3.16)	1.13 (0.95-1.63)	3.43 (1.69-4.71)	<0.001

*Data are presented as (Mean ± SD), @ Data are presented as median (IQR), P-value: One-Way ANOVA. NOND: Non-obese non-diabetic, NOD; non obese diabetic, OND; obese non-diabetic, OD; obese diabetic. SBP: systolic blood pressure, DBP: diastolic blood pressure, HOMA-IR; Homeostasis model assessment insulin resistance, TC; Total cholesterol, LDL-C: low-density lipoprotein cholesterol, TG: triglyceride, HDL-C: high-density lipoprotein cholesterol.

difference between the four groups ($p=0.049$). TG was significantly higher in the OD than in the OND ($p<0.001$) and higher in the NOD than in the control group ($p<0.001$). The other tested parameters such as FI was significantly higher in the OD than in all three other groups. HOMA-IR was higher in the OD than in the OND and

NOND ($p<0.001$). It was also higher in the OD than in the NOD, but this difference was not statistically significant. Tables 2 and 3 show that the serum levels of Spexin were significantly lower in OD than in all other three groups and were lower in the NOD and OND than in the NOND ($p<0.001$). Spexin levels at cutoff value ≤ 0.70 can

Table 2: Comparison of Spexin levels in the study groups

Statistic	NOND (n= 20)	NOD (n= 20)	OND (n= 22)	OD (n= 24)	KW value	P-value
Median (IQR)	1.85 (0.83 – 2.27)	0.7 (0.7 – 0.7)	0.7 (0.6 – 0.7)	0.6 (0.52 – 0.6)	72.579	<0.001
Range	0.8 – 3.8	0.7 – 0.8	0.6 – 0.7	0.17 – 0.6		
Pairwise comparisons	A	B	B	C		

P value: Kruskal -Wallis H test. IQR; inter quartile range. Similar letters = Insignificant difference, Different letters = Significant difference. NOND; Non-obese non diabetic, NOD; non obese diabetic, OND; obese non diabetic, OD; obese diabetic.

Table 3: Spexin cutoff values in different groups

Discrimination of	Cutoff value	P-value	AUC (95% CI)	SN	SP	PPV	NPV
OD from NOND	≤ 0.70	<0.001	1.0	100%	100%	100%	100%
OND from NOND	≤ 0.75	<0.001	1.0	100%	100%	100%	100%
NOD from NOND	≤ 0.75	<0.001	0.975 (0.938–1.000)	80%	100%	100%	83.3%
MS from Non-MS	≤0.75	<0.001	0.783 (0.677–0.889)	94%	60%	77.4%	87.5%

AUC=Area under the Receiver operating characteristic curve, PPV=Positive predictive value. NPV=Negative predictive value, SN =sensitivity, SP=specificity. NOND; non-obese non-diabetic, NOD; non-obese diabetic, OND; obese non-diabetic, OD; obese diabetic, MS; Metabolic Syndrome.

significantly discriminate OD from NOND. Spexin at a cutoff value of ≤0.75 can significantly discriminate OND as well as NOD from NOND and discriminate metabolic syndrome (MS) from Non-MS ($p<0.001$). Tables 4 shows the associations between Spexin and other parameters. Spexin levels were negatively correlated with age, BMI, FI, LDL-C, TC, TG, and HOMA-IR. Tables 5 shows a significantly higher BMI, WC, HC, WHR, FI, and HOMA-IR in the MS than in the NO-MS ($p<0.001$).

Discussion

We have evaluated the serum Spexin levels in T2DM obese patients and assessed its relationship to IR and glycemic control, as the Spexin role is not well recognized, and its mechanism of action is still unclear. We found a non-significant difference between the four groups in age, as the age was matched. TC and TG levels were

significantly higher in the OD than in the control group. Also, TG levels were significantly higher in the OD than in the OND. Moreover, TG levels were significantly higher in NOD than in the control group. The effects of obesity on lipid metabolism include elevated LDL-C, VLDL-C, TG, and low levels of protective HDL-C⁽²⁰⁾. These results coincide with a previous study that reported elevated TG, VLDL, Apo B, and non-HDL-C levels in obese patients⁽²¹⁾. SBP was significantly higher in the OD than in the control group. There is a nearly linear relationship between BMI and BP, and weight loss reduces BP in most hypertensive individuals⁽²²⁾. This result was matched with the results of Blüher⁽²³⁾, who stated that obesity considerably augments the possibility of the development of diseases like T2DM, hypertension, fatty liver disease, stroke, myocardial infarction, osteoarthritis, and dementia.

Table 4: Correlation of Spexin to other study parameters		
Parameter	Correlation coefficient	P-value
Age	-0.413	<0.001
BMI	-0.824	<0.001
Fasting insulin	-0.600	<0.001
HOMA-IR	-0.520	<0.001
TC	-0.264	0.014
LDL-C	-0.216	0.046
HDL-C	-0.010	0.924
TG	-0.317	0.003

P-value: Spearman's correlation test. BMI; body mass index, HOMA-IR; Homeostasis model assessment insulin resistance, TC; total cholesterol, LDL-C: low-density lipoprotein cholesterol, HDL-C: high-density lipoprotein cholesterol, TG: triglyceride.

Our study showed that the Spexin levels were significantly lower in all groups than in the control group and lower in the NOD and OND than in the NOND. Spexin is a neuropeptide. The gene that encoded Spexin (Ch12:orf39) was the most down-regulated gene in obese omental and sc human fat (24). Spexin not only decreased bodyweight but also improved glucose tolerance by reducing IR along with HbA1c⁽²⁵⁾. These results coincide with the results of Karaca et al., and Al-Daghriet al.⁽²⁶⁻²⁷⁾ who found that the median fasting serum Spexin levels were significantly lower in diabetic than in the control subjects that recommends that Spexin may affect the metabolism of glucose. Also, Walewski et al. and Behrooz et al.^(10,28) found considerably lower concentrations of Spexin in obese than in the lean adults and children respectively and suggested a possible satiety-inducing function for it in humans. On the other hand, our results were opposing to the results of Hodges et al.⁽⁸⁾ who evaluated the glucose ingestion effect on the concentration of Spexin in the blood in adolescents. The authors stated that the me

dian on centration of the fasting Spexin in the serum was not different between the study groups and it did not show considerable correlation with any of the body fitness, composition, or biochemical measurements in the blood. Our study showed that only 5 cases (5.8%) had IR; 4 of them had MS and only 1 with no-MS. However, this difference didn't achieve a statistical significance, probably due to the small sample size. Also, it showed that BMI, WC, HC, WHR, and HOMA-IR were significantly higher in patients with MS than in those with NO-MS. In accordance with the 3rd Report of The National Cholesterol Education Program (NCEP) Expert Panel⁽²⁹⁾ which defined MS as: 1) abdominal obesity in which WC>102cm (men), and >88cm (women); 2) TG \geq 150mg/dL; 3) HDL-C<40mg/dL in men and <50 mg/dL in women; 4) blood pressure \geq 130/ \geq 85 mm/Hg; and 5) FBG \geq 110mg/dL. Obesity results in IR that together put stress on β -cells, resulting in a β -cell failure and a progressive decrease in the secretion of insulin⁽³⁰⁾. Because of IR, more amounts of insulin are released at any serum glucose level.

Table 5: anthropometric parameters and homeostasis model assessment of insulin resistance as regard metabolic syndrome

Statistic	MS	No-MS	Z value	P-value
BMI (kg/m²)				
Median (IQR)	34.2 (29 – 38.7)	28.0 (23.7 – 35.6)	-3.894	<0.001
Range	26.5 – 50.2	20.3 – 45.2		
Mean Rank	52.19	30.84		
WC (cm)				
Median (IQR)	109 (100 - 117)	86 (80 - 102)	-5.670	<0.001
Range	85 – 134	53 – 125		
Mean Rank	56.14	25.09		
HC (cm)				
Median (IQR)	120 (112 - 130)	103 (96 - 114)	-5.013	<0.001
Range	102 – 150	80 – 144		
Mean Rank	54.68	27.21		
WHR				
Median (IQR)	0.89 (0.86 – 0.93)	0.86 (0.81 – 0.90)	-2.809	0.005
Range	0.73 – 1.03	0.38 – 0.98		
Mean Rank	49.46	34.37		
Fasting insulin				
Median (IQR)	5.6 (4.1 – 8.4)	3.7 (1.9 – 5.1)	-4.428	<0.001
Minimum-Maximum	1.4 – 20	1.2 – 8		
HOMA-IR				
Median (IQR)	2.48 (1.38 – 3.90)	0.83 (0.47 – 1.27)	-5.929	<0.001
Minimum-Maximum	0.29 – 14.62	0.19 – 3.32		

P-value: Mann-Whitney U test. BMI: body mass index, MS; Metabolic Syndrome, WC: waist circumference, HC: hip circumference, WHR: waist to hip ratio, HOMA-IR; Homeostasis model assessment insulin resistance

The present study showed also that FI was significantly higher in the OD than in the three all other groups. HOMA-IR was significantly higher in OD than in the OND and NOD. It was also higher in OD than in the NOD but this difference was statistically not significant. This study revealed that there was a significantly negative correlation between Spexin and age, FI, HOMA-IR, BMI, TC, LDL-C, and TG. These results were in accordance with the previous studies by Walewski et al., and Kumar et al.^(10,31) who reported that expression of Spexin in the adipose tissue and its levels in the blood were majorly reduced in the subjects suffering from obesity. Chen et al.⁽³²⁾ detected that the levels of serum Spexin were

decreased in IR obese children in comparison with the non-IR obese children and serum Spexin levels correlated significantly and inversely with FI level and HOMA-IR in obese children. Also, Karaca et al. and Al-Daghriet al.^(26,27) reported that Spexin inversely correlated with the blood glucose level, HbA1c, TG, and LDL-C. The limitations in our study were small sample size, factors that might affect Spexin such as physical activity and dietary intake were not taken into consideration.

Conclusion

We concluded that Spexin plays an important role in glucose homeostasis and li

lipid metabolism. The presence of diabetes is associated with lower Spexin levels. Spexin levels were found to be negatively correlated with age, BMI, FI, TC, LDL-C, TG, and HOMA-IR. To understand the exact function of this peptide and to validate the observations in the current study, further investigations and additional studies on larger populations are required.

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