



Biochemical Significance of Soluble Endoglin as a Possible Marker Predicting Cardiovascular Diseases in obese and obese Type 2 Diabetic Patients

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ABSTRACT

Obesity-associated insulin resistance is a major risk factor for Type 2 diabetes and cardiovascular diseases. Endoglin is co-receptor for members of the transforming growth factor (TGF)-superfamily and is highly expressed on vascular endothelial cells. The objective is to examine endoglin as a possible marker predicting cardiovascular complications in obese subjects with or without type 2 diabetes mellitus (type 2 DM). Four groups (22 subjects each) were included: normal control, obese without type 2 DM, non-obese type 2 DM and obese with type 2 DM. Plasma glucose, insulin, glycosylated hemoglobin (HbA1c) levels and homeostasis model assessment of insulin resistance (HOMA-IR) were significantly elevated ($P < 0.05$) in both groups of type 2 DM as compared to the control group indicating poor glycemic control. Unfavorable serum lipids patterns were dramatically found in the obese and obese DM patients than the control and type 2 diabetic groups respectively. The significant increase in serum malondialdehyde (MDA) and the significant decrease ($P < 0.05$) in the total antioxidant capacity (TAC) levels were observed in all groups compared to the control group, however, with higher % change values in the obese type 2 DM group (48.22 and -48.71 respectively) showing that oxidative stress is severely affected by obesity. Soluble endoglin levels were significantly higher ($P < 0.05$) in sera of obese subjects than those of control ones and in obese DM patients than in non-obese DM patients and these elevations were parallel to that of serum VCAM-1 suggesting different degrees of endothelial dysfunctions. Soluble endoglin was significantly correlated in the non-obese and obese type 2 DM with plasma glucose ($r = 0.542, 0.652$ respectively), insulin ($r = 0.589, 0.682$ respectively) and HOMA-IR ($r = 0.517, 0.551$ respectively); also with total cholesterol in the obese non type 2 DM and obese type 2 DM groups ($r = 0.446, 0.459$ respectively) and with MDA ($r = 0.921$) in the obese diabetic group. This indicates the association between sol. endoglin with poor glycemic control, impaired lipid metabolism and oxidative stress in these groups. Therefore, soluble endoglin may suggest to be used a useful marker in the early detection of the risk of cardiovascular complications in obese and obese type 2 diabetic patients.

Introduction

Rates of diabetes have increased markedly over the last 50 years in parallel with obesity. Vascular disease is the main cause for disability and death in patients with diabetes mellitus ^[1].

Endothelial dysfunction indicated by impaired endothelium dependent vasodilatation is common in early and otherwise uncomplicated type 2 diabetes ^[2] and has also been shown to be predictive of future adverse cardiovascular events which are partly due to the frequent association of the disease with other cardiovascular risk factors, including hypertension, obesity and dyslipidemia ^[3].

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Circulating biomarkers have been important in predicting development of diabetes and its complications as well as providing targets for therapy^[4]. Endoglin (also known as CD105) is a type I transmembrane glycoprotein that may have roles in hematopoiesis, cardiovascular development and angiogenesis. The glycoprotein consists of a homodimer of 180 kDA with disulfide links^[5]. There are two isoforms of endoglin created by alternative splicing: the long isoform (L-endoglin) and the short isoform (S-endoglin)^[6]. A soluble form of endoglin can be produced by the proteolytic cleaving action of metalloproteinase MMP-14 in the extracellular domain near the membrane^[7].

Endoglin has been shown to interact with TGF- β receptor 3 and TGF- β receptor 1. It is highly expressed on vascular endothelial cells. Endoglin and activin like kinase-1 (ALK-1) proteins are specific endothelial receptors of the transforming growth factor- β (TGF- β) superfamily that are essential for vascular integrity^[8]. Through binding to the TGF- β type II receptor, TGF- β can activate two distinct type I receptors (ALK1 and ALK5) in endothelial cells, each one leading to opposite effects on endothelial cell proliferation and migration^[9]. Endoglin plays a pivotal role in the balance of ALK1 and ALK5 signaling to regulate endothelial cell proliferation in response to TGF- β ^[10]. Endoglin expression was demonstrated in atherosclerotic vessels predominantly in endothelial cells and smooth muscle cells in various types of blood vessels in mice and humans, suggesting its participation on atherogenesis^[11].

Supporting the link between endoglin and metabolism is the relationship between plasma levels of endoglin and glycemia that was recently found in diabetic hypertensive patients^[12]. In addition soluble endoglin (sol. endoglin) are released under hypoxic stress and is an indicator of hypertension- and diabetes-associated vascular pathologies in humans and animals^[13, 14] and there is a clear association between endothelial dysfunction and alterations in glucose metabolism or metabolic syndrome^[15].

Accordingly, the present study aimed to assist the relationship between sol. endoglin and hyperglycemia, oxidative stress and endothelial dysfunction in obese and/or Type 2 diabetic patients and to evaluate whether sol. endoglin can be used as a possible biomarker predicting future adverse cardiovascular events in those patients.

Subjects and Methods

Subjects:

This study included 88 volunteers of both sexes (closely sex-matched by ratio in each subject group) and aged from 45 to 65 years. The participants presented as 44 non type 2 DM subjects and 44 age matched subjects diagnosed as type 2 DM patients (recent onset <5 years) according to Report of the Expert Committee on the Diagnosis and classification of Diabetes Mellitus (2006)

The participants were selected from the outpatient's clinic of National Institute of Diabetes and Endocrinology (NIDE), Cairo, Egypt. The participants in the diabetic groups were treated with oral hypoglycemic agents only. Hypoglycemic medications were withheld on the morning of the study.

Design and study populations: Subjects included in this study were further classified into the followings: 22 normal control non obese subjects (BMI < 30 kg/m²), 22 obese subjects without type 2 DM (BMI \geq 30 kg/m²), 22 non obese type 2 DM patients and 22 obese type 2 DM patients (BMI \geq 30 kg/m²).

Exclusion Criteria: Patients with type 2 DM complications such as diabetic nephropathy, neuropathy, retinopathy, hypertension, and heart disease based on clinical and laboratory investigations were excluded from the study. The patients were instructed not to engage in any vigorous exercise for at least 3 days before the study. Each subject included in the study signed an informed consent form to participate in the study after full explanation of the purpose and nature of all procedures used. Approval was taken from the research committee of General Organization of teaching Hospitals and Institutions.

Methods:

Blood sampling: A total of 9 ml of venous blood obtained were collected by venous puncture from each subject after 12 hours overnight fasting processed as follows: 5ml were subdivided into tubes with or without anticoagulant for separation of plasma and serum samples respectively, 2 ml were added in tube with EDTA without centrifugation (whole blood sample) for assaying HbA1c % and another 2 ml of blood samples were added in tubes with potassium oxalate and sodium fluoride for assaying of glucose.

Biochemical analysis

Diabetic biomarkers: Plasma glucose concentration was assayed by glucose oxidase method according to **Trinder**^[16]. Blood HbA1c % was measured according to the method of **Grey et al.**^[17] using an immunoturbidimetric assay on Dimension RxL Max (Dade Behring). Plasma insulin was determined using human commercially available enzyme-linked immunosorbent assay kits (Bio Source Europe S.A., Nivelles, Belgium) according to **Flier et al.**^[18]. Insulin resistance was measured using homeostasis model assessment of insulin resistance (HOMA-IR) by multiplying fasting plasma glucose (mmol/L) and fasting plasma insulin (mU/L) divided by 22.5^[19].

Lipid profile and oxidative status parameters: Commercial kits based on different techniques, purchased from Bio Med, Egy-Chem were used for the determinations of serum total cholesterol^[20], HDL-C^[21] and triglycerides^[22]. Serum LDL-C was calculated as follows: LDL-C = TC - HDL-C / TG/5^[23] while, atherogenic index was calculated from ratio of TC/HDL-C^[24].

Oxidative status markers: Malondialdehyde (MDA) was determined in serum as an index of lipid peroxidation according to the method described by **Uchiyama and Mihara** [25] while total antioxidant capacity (TAC) was determined according to **Koracevic et al.** [26] method.

Endothelial function parameters: Serum VCAM-1 was determined using quantitative sandwich enzyme immunoassay technique (Human) designed by Cloud-Clone Corp. (USA) Catalogue: SEA547Hu. Soluble endoglin was assayed by ELISA technique provided by Human Endoglin (ENG) Kit CATALOG #: 95702. Glory Science Co., Ltd USA, according to the method of **Abdalla et al.** [27].

Statistical analysis: All results were expressed as the mean \pm SD. Statistical analysis was performed with Statistical Package for the Social Science for Windows (SPSS, version 16.0, Chicago, IL, USA). The data were analyzed by one-way analysis of variance (ANOVA). Pearson's correlation analysis was used to determine the correlation among the parameters assessed. The *P*-value less than 0.05 were considered statistically significant.

Results

Demographic characteristics of the patients and controls are presented in Table (1): No statistical differences were found between all groups with respect to age and blood pressure. However, BMI showed significant increase ($P<0.05$) in obese group against the control group and in obese type 2 DM group against the non-obese type 2 DM group.

Diabetic biomarkers of all groups presented in Table (2): Obese subjects exhibited significant elevations ($P<0.05$) in plasma insulin and HOMA-IR values compared to the normal non obese control group (% change: 44.07 and 54.09% respectively). Group of patients diagnosed as type 2 DM showed significant elevations ($P<0.05$) in plasma glucose, HbA1c %, plasma insulin and HOMA-IR levels as compared to the control group with % changes: 133.40, 68.53, 52.54 and 252.27% respectively. Obese type 2 DM patients as compared to the non-obese diabetic ones, exhibited significant increases ($P<0.05$) in plasma glucose levels (41.73%), HbA1c (22.54%) and HOMA-IR levels (56.13 %).

Table (3) demonstrated the lipid profile pattern in all groups: In obese group of patients there were significant elevations ($P<0.05$) in serum TC, LDL-C, TG levels and TC/HDL ratio while HDL-C showed significant reduction ($P<0.05$) compared to normal non obese controls (% changes: 29.81, 21.95, 149.58, 41.35 and -15.19 respectively). Non obese type 2 DM patients exhibited significant increases ($P<0.05$) in the levels of serum TC, LDL-C, TG, TC/HDL ratio and a significant decrease ($P<0.05$) in HDL-C levels as compared to the normal control ones with % changes: 26.31, 38.03, 77.95, 48.65 and -15.42% respectively. Obese type 2 DM group of patients had significantly higher ($P<0.05$) levels of serum TC, LDL-C, TG and TC/HDL ratio (% changes: 28.06, 40.22, 40.81 and 58.18% respectively) while they had significantly lower ($P<0.05$) HDL-C levels (-20.14%) than the non-obese type 2 DM group.

Table 1: Age, BMI, systolic and diastolic blood pressure of participants included in the different studied groups.

Parameter Groups	Age (Years)	BMI (kg/m ²)	SBP (mm Hg)	DBP (mm Hg)
Normal Control	52.69 \pm 11.08	27.55 \pm 1.80	124.00 \pm 23.00	78.00 \pm 14.00
Obese without DM *% change	53.90 \pm 11.68 1.10	34.19 \pm 2.40 ^a 24.10	135.00 \pm 15.00 8.80	85.00 \pm 10.00 8.97
Non obese T2 DM *% change	55.27 \pm 12.74 4.89	27.64 \pm 1.40 ^b 0.32	136.00 \pm 11.00 9.70	85.00 \pm 8.00 8.97
Obese with T2 DM *% change **% change	60.17 \pm 14.03 14.19 8.87	33.70 \pm 1.50 ^{ac} 22.32 21.92	139.00 \pm 17.00 12.09 2.20	90.00 \pm 9.00 15.38 5.88

Values are represented as mean \pm SD of 22 subjects /group. *P* value was significant at < 0.05 .

a: Significant difference from normal control group; b: Significant difference from obese group; c: Significant difference from type 2 DM group. * % change from control group. ** % change from non-obese type 2 DM group

Table 2: Fasting plasma glucose, glycosylated hemoglobin, plasma insulin levels and HOMA-IR scores of participants included in the different studied groups.

Parameters Groups	Plasma Glucose (mmol /L)	HbA1c %	Plasma insulin (mU/L)	HOMA-IR
Normal Control	5.00 ± 0.60	5.37± 0.60	10.03 ± 0.80	2.20 ± 0.50
Obese without DM *% change	5.29 ± 0.56 5.80	5.61 ± 0.54 4.47	14.45± 1.75 ^a 44.07	3.39 ± 0.50 ^a 54.09
Non obese T2 DM *% change	11.67±1.70 ^{ab} 133.40	9.05±1.17 ^{ab} 68.53	15.30±1.20 ^a 52.54	7.75± 1.00 ^{ab} 252.27
Obese with T2 DM *% change **% change	16.54 ± 4.1 ^{abc} 230.80 41.73	11.09±1.90 ^{abc} 106.52 22.54	16.40 ± 1.00 ^{ab} 63.51 7.19	12.10±1.4 ^{abc} 450.00 56.13

Values are represented as mean ± SD of 22 subjects /group. *P* value was significant at < 0.05.

a: Significant difference from normal control group; b: Significant difference from obese group; c: Significant difference from type 2 DM group. * % change from control group. ** % change from non-obese type 2 DM group

Table 3: Lipid profile of participants included in the different studied groups.

Parameters Groups	Serum TC (mg %)	Serum HDL (mg %)	Serum LDL (mg %)	Serum TG (mg %)	TC / HDL
Normal Control	186.52± 28.97	51.08± 5.80	112.40± 21.01	104.16± 27.18	3.70± 0.40
Obese without DM *% change	242.12± 24.46 ^a 29.81	43.32± 6.58 ^a -15.19	137.08± 22.47 ^a 21.95	259.96± 36.90 ^a 149.58	5.23± 0.49 ^a 41.35
Non obese T2 DM *% change	235.60± 30.13 ^a 26.31	43.20± 4.36 ^a -15.42	155.15± 26.06 ^a 38.03	185.35± 25.97 ^{ab} 77.95	5.50± 0.60 ^a 48.65
Obese with T2 DM *%change **%change	301.70± 40.24 ^{abc} 61.75 28.06	34.50± 3.76 ^{abc} -32.46 -20.14	217.55± 31.04 ^{abc} 93.55 40.22	261.00± 27.21 ^{ac} 150.58 40.81	8.70± 0.77 ^{abc} 135.14 58.18

Values are represented as mean ± SD of 22 subjects /group. *P* value was significant at < 0.05.

a: Significant difference from normal control group; b: Significant difference from obese group; c: Significant difference from type 2 DM group. * % change from control group. ** % change from non-obese type 2 DM group

Parameters of oxidative status and endothelial function were listed in Table (4): Results of oxidative status parameters revealed that in obese patients without type2 DM serum levels of MDA were significantly increased ($P<0.05$) by 18.76% while serum TAC levels were significantly decreased ($P<0.05$) by 12.80% as compared to the normal control subjects. Non obese type2 DM

patients comparing to their respective normal controls showed the same results but with different % change values (MDA 31.14% & TAC -36.54%). In addition, obese type 2 DM patients had significantly higher ($P<0.05$) MDA levels by 13.02% while they had significant lower ($P<0.05$) TAC levels by 19.19% than the non-obese type 2 DM group.

Concerning the endothelial function parameters, the obese group of patients without type 2 DM had significantly higher ($P<0.05$) levels of serum VCAM-1 and sol. endoglin (% change: 33.88 & 17.39% respectively) than the control group. The two parameters showed the same significant changes in the non-obese type 2 DM patients compared to the normal control ones but with higher values of % changes (50.94 and 26.09% respectively). Furthermore, comparing obese type 2 DM patients to the non-obese diabetic ones, serum levels of VCAM-1 and sol. endoglin were also significantly increased ($P<0.05$) with % changes 12.87 and 36.21% respectively.

Correlation studies (Table 5): In the non-obese type 2 DM and obese type 2 DM groups of patients, positive significant correlations were found between sol. endoglin and plasma glucose ($r=0.542$ & $P<0.001$ and $r=0.652$ & $P<0.001$ respectively); also between sol. endoglin and insulin ($r=0.589$ & $P<0.001$ and $r=0.682$ & $P<0.001$ respectively) and between sol. endoglin and HOMA-IR ($r=0.517$ & $P<0.001$ and $r=0.551$ & $P<0.001$ respectively). Sol. endoglin also showed significant positive correlations with TC in the obese with or without type 2 DM groups ($r=0.459$ & $P<0.01$ and $r=0.446$ & $P<0.026$ respectively) however, it showed a positive significant correlation with MDA ($r=0.921$ & $P<0.001$) in the obese with type 2 DM group only.

Discussion

Obesity plays a central role in the insulin resistance syndrome, which includes hyperinsulinemia, hypertension, hyperlipidemia, Type 2 diabetes mellitus, and an increased risk of atherosclerotic cardiovascular

disease^[28].

This study enrolled obesity as a risk factor for cardiovascular diseases in obese without type 2 DM patients and with type 2 DM patients and assessed the association between sol. endoglin and insulin resistance, hyperlipidemia, oxidative stress and endothelial dysfunction. The study also examined the possibility of using sol. endoglin as a predictor of future adverse cardiovascular events in obese and/or type 2 DM patients.

All participants in this study were age matched and normotensive (Table 1). Results collected from Table (2) indicated that obesity induced a degree of insulin resistance manifested from the elevated HOMA-IR scores (54.09%). Furthermore, both of the type 2 DM groups either non obese or obese exhibited poor glycemic control and insulin resistance however, the involvement of obesity induced dramatic changes. These findings were in agreement with **Rexrode et al.**^[29] who reported that obesity has been strongly associated with insulin resistance in normoglycemic persons and in individuals with Type 2-diabetes and that increases the risk of cardiovascular disease in adults. Also **Šindelka et al.**^[30] concluded that obesity may have even greater influence on the insulin action than diabetes mellitus itself. The positive significant correlations found between sol. endoglin and plasma glucose, insulin and HOMA-IR (Table 5) supported the link between endoglin and the glycemic status in the diabetic groups. **Blazques-Medela et al.**^[13] reported that there was a significant correlation between sol endoglin and glycemia however with hypertension as a risk factor for cardiovascular complications in patients with type 2 DM.

Table 4: Serum malondialdehyde, total antioxidant capacity, vascular cell adhesion molecule-1 and sol. endoglin levels of participants included in the different studied groups.

Parameters Groups	Serum MDA (nmol/L)	Serum TAC (mmol/L)	Serum VCAM-1 (ng/ml)	Serum Endoglin (pg/ml)
Normal Control	5.33±1.04	1.56±0.10	468.40±16.83	4.60±0.71
Obese without DM *% change	6.33±0.99 ^a 18.76	1.22±0.09 ^a -12.80	627.10±37.60 ^a 33.88	5.40±0.69 ^a 17.39
Non obese T2 DM *% change	6.99± 1.10 ^a 31.14	0.99±0.08 ^a -36.54	707.00±32.80 ^a 50.94	5.80±0.71 ^a 26.09
Obese with T2 DM *% change **% change	7.90±1.09 ^{abc} 48.22 13.02	0.80 ±0.08 ^{abc} -48.71 -19.19	798.00±47.40 ^{abc} 70.36 12.87	7.90±0.78 ^{abc} 71.74 36.21

Values are represented as mean ± SD of 22 subjects /group. P value was significant at <0.05 .

a: Significant difference from normal control group; b: Significant difference from obese group; c: Significant difference from type 2 DM group. * % change from control group. ** % change from non-obese type 2 DM group

Table 5: Pearson's correlation coefficient (r) between sol. endoglin and some biochemical parameters in the different studied groups.

		Glucose	Insulin	HOMA-IR	Total cholesterol	MDA
Obese without DM	Pearson's correlation(r)	N.S	N.S	N.S	0.446	N.S
	P value <				0.026	
T2DM	Pearson's correlation(r)	0.542	0.589	0.517	N.S	N.S
	P value <	0.001	0.001	0.001		
Obese with T2DM	Pearson's correlation(r)	0.652	0.682	0.551	0.459	0.921
	P value <	0.001	0.001	0.001	0.01	0.001

Impaired lipid metabolism resulting from uncontrolled hyperglycemia has been implicated in cardiovascular complications in diabetes patients^[31]. Unfavorable lipids patterns were found in the obese patients against the control subjects and in the obese type 2 DM patients against the non-obese ones (Table 3). These findings were manifested from the significant differences observed when comparing the various parameters of blood lipid profile in relation to TC, LDL-C, TG HDL-C levels and TC/HDL-C ratio in these groups. It was reported that dyslipidemia of obesity is commonly manifested as high plasma triglyceride levels, low HDL-C and normal LDL-C with preponderance of small dense LDL- particles^[32] and there is a linear correlation between the degree of obesity and plasma level of LDL cholesterol and triglycerides^[33]. Current results indicated that hypercholesterolemia were associated with increased levels of sol. endoglin in the obese groups as indicated from the positive significant correlations between sol. endoglin and TC in the obese groups with or without type 2 DM groups (Table 5). Recently, **Beiroa et al.**^[14] reported for first time that heterozygous endoglin deficiency in mice decreases high fat diet-induced hepatic triglyceride content and insulin levels. The overall findings of those authors indicated that endoglin is a potentially important physiological mediator of insulin levels and hepatic lipid metabolism. Accumulating evidence suggested a close relationship between hyperlipidemia and oxidative stress in obese patients^[34]. In the present study (Table 4) all the studied groups exhibited certain degrees of oxidative stress compared to the normal control group manifested from the elevation in levels MDA and the reduction in the levels of TAC. However, from the % change values it was severely presented in the obese diabetic group of patients compared to the control group (MDA 48.22% and TAC -48.71% respectively). This could be explained

on the bases that obesity is associated with increased oxygen consumption, cell injury/inflammation, increased fat deposition, and compromised antioxidant defense^[35] and that the greater the obesity, the greater demand on antioxidant enzymes to combat free-radical damage, hence antioxidant enzymes become depleted^[36]. Additionally, the increased superoxide production from fatty acids in mitochondria of the endothelial cells directly inactivates 2 critical anti-atherosclerotic enzymes, endothelial nitric oxide synthase and prostacyclin synthase causing defective angiogenesis^[37]. Furthermore, in the diabetic condition, oxidative stress impairs glucose uptake in muscle and adipocytes^[38] and affected insulin secretion from pancreatic β cells thus causing abnormalities in the secretion and action of insulin^[39]. In the present results the positive significant correlations between sol. endoglin and MDA in the obese type 2 DM group (Table 5) indicating possible association between sol. endoglin and oxidative stress in this group.

Endothelial dysfunction represents an early phase of vascular changes that eventually lead to atherosclerosis with all its unfavorable complications and the damaged endothelium presents a range of adhesion molecules to the arterial lumen^[40]. In the current study the levels of VCAM-1 was elevated ascending from the obese group without type 2 DM (33.88%) to the obese group with type 2 DM (70.36 %). Previous study of **Bořanská et al.**^[41] declared that circulating adhesion molecules like VCAM-1 in patients with obesity were elevated and that this effect may play an important role in the development of endothelial dysfunction-/atherosclerosis. Further studies reported that poor glycemic control together with increased glucose levels may be responsible for significant higher levels of VCAM-1 in diabetic patients^[42,43].

From the current results the elevations in the levels of

sol. endoglin were parallel to that of VCAM-1 levels in the sera of obese patients without type 2 DM against the controls and in the sera of obese type 2 DM patients against the non-obese ones (Table 4). This indicated an association between sol. endoglin and endothelial dysfunction which was found to be increased when obesity enrolled as a risk factor of cardiovascular alterations in these groups. The results of Kurki *et al.* [44] in mice indicated that obesity was associated with increased expressions of angiogenesis-related proteins among which were sol. endoglin. Recently, it was reported that in the normal state, L-endoglin modulated the TGF- β response but upon senescence of endothelial cells, S-endoglin is up-regulated, interacting with the TGF- β receptor complex containing ALK1 and ALK5 [45,46]. As a consequence of this interaction, S-endoglin up-regulate plasminogen activator inhibitor type1 (PAI-1)/extra-cellular matrix (ECM) synthesis which may lead to increased fibrosis; down-regulate inhibitor of DNA binding 1 (Id1) which associated with decreased angiogenesis and down-regulate endothelia nitric oxide synthase and up-regulate cyclooxygenase-2 which are involved in endothelial dysfunction and impaired vascular relaxation [47,48]. Also, endoglin regulates the half-life and activity of eNOS thus S-endoglin allows a switch that triggers the cardiovascular pathology [49]. In addition sol. endoglin amplifies the vascular damage mediated by vascular endothelial growth factor-1 (VEGF-1) [50].

Conclusion: It can be concluded that sol. endoglin is associated with poor glycemic control, impairment in lipid metabolism and oxidative stress. Additionally, sol. endoglin seems to be associated with impaired endothelial function which is a major characteristic of patients with obesity and /or diabetes. The results were relevant enough to establish the relative strength of prediction of cardiovascular risk according to the endoglin level presented by the patient.

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