

C-peptide as a biomarker in preeclampsia

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SUMMARY

Preeclampsia (PE) is a syndrome exclusive to human pregnancy and responsible for high perinatal morbidity and mortality, whose manifestations include hypertension, proteinuria, and edema. There is evidence that PE incidence is four times higher in diabetic type 1 women than in non-diabetic women; and increases in women with metabolic syndrome and insulin resistance. C-peptide is a 31 amino acid residue polypeptide part of the proinsulin, which is enzymatically cleaved into insulin and C-peptide molecule. Both are simultaneously secreted in equimolar concentrations into blood, thus serum C-peptide level might reflect real insulin production. Little is known about the relationship between circulating levels of C-peptide and the increase in blood pressure in PE. For this reason, we assessed serum levels of C-peptide, and insulin in women with normal pregnancy and with PE, in a population of 30 Venezuelan women. Serum samples were evaluated by multiplex microsphere analysis (Bio-Plex Pro Assays).

Our results show increased insulin and C-peptide serum levels in women with PE. Spearman correlation analysis in all subjects showed a positive association between SBP and C-peptide ($r= 0.4841$; $P<0.0251$) and insulin ($r= 0.4386$; $P<0.0221$), associated with a positive correlation between SBP and proteinuria and glucose. Our results suggest that an increase in C-peptide could be associated with the development of hypertension and insulin resistance in PE. Therefore, the quantification of this peptide could be a promising biomarker to predict the onset of pregnancy-induced hypertension.

Key words: Preeclampsia, C-peptide, Insulin resistance, hypertension.

RESUMEN

La preeclampsia (PE) es un síndrome exclusivo de la gestación humana y responsable de una alta morbilidad perinatal, cuyas manifestaciones incluyen la hipertensión arterial, proteinuria y edema. Existe evidencia que su incidencia es cuatro veces mayor en mujeres diabéticas tipo 1 que en las mujeres no diabéticas; y también incrementa en mujeres con síndrome metabólico y resistencia a la insulina. El péptido C es un polipéptido de 31 aminoácidos, un subproducto del proceso de la biosíntesis de la insulina, que resulta del clivaje de la proinsulina. Tanto la insulina como el péptido C son secretados en cantidades equimolares y liberados al torrente sanguíneo. Se conoce poco acerca de la relación que existe entre los niveles circulantes del péptido C y los incrementos de la presión arterial en la PE. Por ello, en el presente estudio se cuantificaron los niveles plasmáticos del péptido C e insulina en mujeres

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con embarazo normal y con PE, pertenecientes a una población de 30 mujeres venezolanas. Se evaluaron muestras de suero mediante el análisis multiplex de microesferas (Bio-Plex Pro Assays). Los resultados muestran incrementos significativos de los niveles plasmáticos del péptido C y la insulina en las mujeres con PE. Mediante el análisis de correlación de Spearman en todas las participantes, la PAS mostró una asociación positiva con el péptido C ($r=0,4841$; $P<0,0251$) y con la insulina ($r=0,4386$; $P<0,0221$), estos asociados a una correlación positiva entre PAS, proteinuria y glucosa. Los resultados sugieren que el incremento del péptido C podría estar asociado al desarrollo de hipertensión y resistencia a la insulina en pacientes con PE. Igualmente, indican que la cuantificación de este péptido circulante podría constituir un biomarcador prometedor para predecir la aparición de la hipertensión inducida por el embarazo.

Palabras clave: Preeclampsia, péptido C, resistencia a la insulina, hipertensión.

INTRODUCTION

Preeclampsia (PE) is a complex disease, exclusive to human pregnancy, being the main cause of fetal and maternal morbidity and mortality. For this reason, it has been called the disease of multiple theories, in which both environmental and genetic factors have been associated with its development. It is characterized by new-onset hypertension and proteinuria, which usually arises after 20 weeks of gestation, more frequently in the third trimester and reverses in the postpartum period (1-4). In women with pre-gestational diabetes mellitus (type 1 or type 2), the risks for preeclampsia and pregnant-induced hypertension (PIH) is elevated about four times compared to non-diabetic women (5). Likewise, women with metabolic syndrome are also at high risk of suffering PE (6). Metabolic syndrome and type 2 diabetes are known to be associated with insulin resistance (7). In effect, insulin sensitivity in late normal pregnancy is 45 to 70 percent lower than that of nonpregnant women, and pregnancy itself induces a state of peripheral insulin resistance, the purpose of which is likely to ensure a sustained postprandial supply of glucose to the fetus (8). Indeed, insulin sensitivity in late normal pregnancy is 45 to 70 percent lower than that of nonpregnant women, and it is restored postpartum (9). Several lines of evidence suggest

mid-trimester maternal insulin resistance is associated with a significantly increased risk of subsequent preeclampsia. Furthermore, the fact that PE usually begins late in pregnancy, when insulin resistance is highest (10), supports this possible association. However, the mediators responsible for the possible relationship between PE and insulin resistance have not been fully established. Recently, researchers have focused on several new potential mediators of insulin resistance in gestational diabetes mellitus (GDM), including C-peptide, which could be involved in the regulation of insulin resistance, kidney dysfunction, and the increase in blood pressure during pregnancy.

The connector peptide, or C-peptide, is a 31 amino acid residues that result from cleavage of proinsulin in its middle portion, between the segment A and B chains of insulin (11,12). Both C-peptide and insulin are stored within the secretory granules of pancreatic beta cells and are released into the bloodstream in equimolar amounts in response to glucose stimulation (13,14). Its deficiency, together with that of insulin, is the main characteristic of type 1 diabetes mellitus (T1DM), and of the late phase of type 2 diabetes mellitus (T2DM); therefore, quantification of circulating C-peptide in clinical practice is a useful and widely used method to monitor pancreatic beta-cell function, discrimination between T1DM and T2DM, detection of absolute insulin deficiency and the identification of patients with diabetes of adult-onset (15).

C-peptide has been considered for many years as an inactive peptide; however, numerous studies reveal that this molecule plays a physiological role in different cell types (16-18). There is evidence that indicates that C-peptide can suppress several molecular mechanisms involved in the pathophysiology of diabetic nephropathy and, therefore, prevents the onset and progression of diabetes-induced kidney failure (17,20). Likewise, studies in animal models of T1DM reveal that C-peptide has positive effects in the early phase of nephropathy, retinopathy, and neuropathy (21-23), also having a positive impact on the kidneys, retina, and nerves in patients with T1DM and T2DM (24-26). Yasuhi et al. (27) reported that during the second trimester of pregnancy there is a strong correlation

between circulating levels of C-peptide and glucose and the subsequent development of gestational hypertension. However, little is known about the relationship between blood pressure values and circulating levels of C-peptide in PE. Therefore, and to establish the possible correlation between the increase in blood pressure in PE and the alterations in the circulating levels of C-peptide, in this study serum levels of this peptide were determined in a group of Venezuelan healthy pregnant women and PE pregnant women.

MATERIALS AND METHODS

Subjects

A descriptive cross-sectional study was conducted in healthy pregnant women and in women with PE who meet the inclusion requirements. The sample consisted of 30 pregnant women (17 healthy pregnant women and 13 pregnant women with PE), aged between 17 and 40 years old and with gestational ages between 28 and 40 weeks, or in the last trimester of pregnancy, who attended the Gynecology Service of the Hospital Universitario de Caracas, Caracas-Venezuela, between the dates of June 2014 to July 2015 were selected. Pre-eclamptic pregnant women were defined as those who presented proteinuria (≥ 0.3 g/24-hour urine) and/or systolic blood pressure (SBP) values equal to or greater than 140 mmHg, or diastolic blood pressure (DBP) equal to or greater than 90 mmHg, measured twice, 4-6 hours apart.

Exclusion criteria were established: chronic arterial hypertension, autoimmune diseases, type 1 and type 2 diabetes mellitus, angiopathy, mental retardation, psychiatric diseases, neurological disorders, multiple pregnancies, chronic kidney disease, patients with a body mass index equal to or greater than 30 kg/m², maternal or fetal infection, and congenital anomalies of the fetus. The selected patients underwent a physical examination, blood pressure measurement with the use of a mercury sphygmomanometer, afterward biological samples were taken.

All the volunteers signed and dated the informed consent which was written simply, and

carefully read and orally explained by trained personnel, indicating what the study consisted of. Likewise, the participants clarified their doubts with the investigator in charge.

All the procedures used were reviewed and approved by the Bioethics Committee of the Hospital Universitario de Caracas, Caracas-Venezuela, and complied with the Declaration of Helsinki for experimentation studies with human beings (1975 and revised in 1983).

Biological samples

The pre-analytical conditions were the ones recommended worldwide for this type of assessment: Blood samples were obtained after 8-12 hours of fasting and without a regular diet. The blood sample was obtained by direct venipuncture in the antecubital region with multiple needles (Venojet[®]), using tubes with a coagulation accelerator (Vacutainer[®]). Immediately, they were centrifugated at 3000 rpm for 15 minutes. Serum was separated and concentrations of insulin and C-peptide, BUN, and creatinine were determined. Serum aliquots were stored at a temperature of -80 °C until further analysis.

Blood pressure determination

In the sitting position, the systolic and diastolic blood pressure was determined with the use of a mercury sphygmomanometer, which was placed on the left arm.

Biochemical methods

Blood glucose values were determined by an enzymatic method, using the commercial kit (Stanbio) with reference values being 70–105 mg/dL. Serum creatinine was determined by a colorimetric method using a commercial kit (Laboratorios Biogamma C.A., Caracas, Venezuela). The results were expressed in mg/dL. Urinary proteins were determined using a commercial kit and the results were expressed as g/24h (0.05-0.15 g/24h) (STAT FAX 3300 Chemistry Analyzer).

Serum diabetes markers determination

All serum samples were evaluated in duplicate by multiplex microsphere analysis (Bio-Plex Pro Assays, Life Science Group, BIORAD). Briefly, the Bio-Plex® system is based on three technological cores. The first uses up to 100 polystyrenes (5.6 µm) or magnetic (8 µm) microspheres, fluorescently stained encoded with a spectral code (xMAP Technology), which allows the simultaneous detection of 100 different molecules in one of the wells of the 96-well microplate. The second is a flow cytometer with two laser beams associated with an optical system that allows the different molecules attached to the surface of the microspheres to be quantified. The third is a high-speed digital signal processor that handles fluorescence data with high efficiency. With this technique concentrations of C-peptide and insulin were assessed and expressed as pg/mL.

Statistic analysis

The data were expressed as the mean ± standard error of the mean (S.E.M.). Data distribution was evaluated using the Shapiro-Wilk, Kolmogorov-Smirnov, Jarque-Bera normality tests. The data were analyzed using the GraphPad InStat program 5.0 version (GraphPad Software, Inc). The Mann Whitney U test was used to compare the values of the variables of the groups under study. The correlations between variables were made by the Spearman correlation test. A value of P<0.05 was considered statistically significant.

RESULTS

Characteristics of the patients

The clinical characteristics of the patients are described in Table 1. No significant differences were found in age, weight, height, gestational age and body mass index between the two groups studied. In contrast, SBP and DBP values were significantly higher in women with PE when compared with control pregnant women.

Table 1
Clinical characteristics of pregnant control and preeclamptic patients

Parameter	Healthy pregnant women	Preeclamptic patients	P
	N=30	N= 17	N= 13
Age (years)	26.47 ± 0.92	28.62 ± 2.70	NS
Weight (kg)	71.80 ± 2.97	74.24 ± 2.46	NS
Size (m)	1.60 ± 0.02	1.59 ± 0.01	NS
BMI (kg/m ²)	28.4 ± 0.91	29.33 ± 0.99	NS
Gestational age (weeks)	35.5 ± 1.08 (28 - 40)	35.38 ± 0.99 (29 - 41)	NS
SBP (mmHg)	112.90 ± 2.26	153.58 ± 2.74***	<0.0001
DBP (mmHg)	74.20 ± 1.81	103.16 ± 3.61****	<0.0001

NS: no significant

Laboratory parameters

Table 2 shows significant increases in blood glucose, blood urea nitrogen (BUN), proteinuria,

and creatinine in preeclampsia pregnant women when compared to pregnant controls of the same gestational age.

Table 2
Laboratory parameters of healthy and preeclamptic pregnant women

Parameter	Healthy pregnant women	Preeclamptic patients	P
	N=30	N=13	
Glycemia (mg/dL)	73.50 ± 2.46	86.23 ± 5.2*	0.0305
BUN (mg/dL)	11.40 ± 1.23	17.61 ± 2.10*	0.0123
Proteinuria (g/24h)	0.25 ± 0.10	4.0 ± 2.4 ***	0.0001
Serum Creatinine (mg/dL)	0.54 ± 0.02	0.66 ± 0.05*	0.042

Serum C-peptide and insulin levels in pregnant women with preeclampsia or controls

As seen in Figure 1, a significant increase in serum levels of C-peptide and insulin was observed in the group of PE compared to the control group.

Correlation between SBP and clinical laboratory parameters

Spearman correlation analysis in all subjects shows that there is a significant positive correlation between SBP vs. proteinuria, glycemia, C-peptide, and insulin (Table 4).

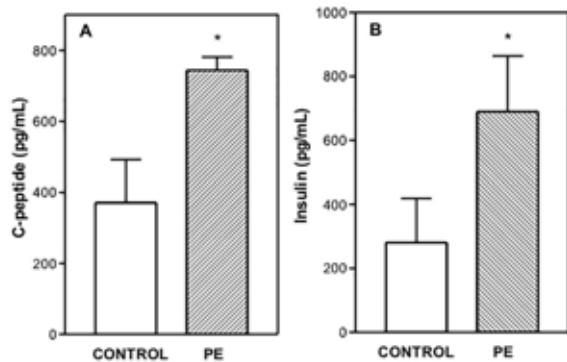


Figure 1. Serum levels of C-peptide C (A) and insulin (B) in pregnant control or preeclampsia (PE) women. *P<0.05.

Table 4
Spearman's correlation analysis

	r	P
SBP & Proteinuria	0.8192	0.0001
SBP & Glycemia	0.4150	0.0281
SBP & C-peptide	0.4082	0.0251
SBP & Insulin	0.4386	0.0221

DISCUSSION

In the present study, we demonstrate an increased serum level of C-peptide in Venezuelan women in their third trimester of pregnancy

with PE, when compared with those of pregnant normotensive control, which was associated with a significant increase in serum glucose and insulin. Furthermore, SBP values of all the population of the pregnant woman were positively correlated with serum levels of C-peptide, insulin, glucose, and proteinuria, suggesting the coexistence of PE with possible maternal insulin resistance in the last trimester of pregnancy. In this regard, it was shown that in the second trimester of pregnancy, there is a strong correlation between C-peptide levels and fasting and postprandial glucose, and the probability of subsequent development of gestational hypertension, being this association independent of maternal obesity and blood pressure during mid-pregnancy (27). These findings may reflect an amplified response of pancreatic beta-cells to glycemic stimulus, similar to those found in states of insulin resistance. Likewise, in a preliminary study in 301 patients from sub-Saharan Africa, dysregulation of C-peptide was evaluated in the presence of PE and HIV infection, and it was found that serum C-peptide was elevated in PE when compared to non-pregnant and normotensive pregnant women, while the alteration of C-peptide levels was not affected by HIV status (28). Contrary to our findings, El Sayed et al. (29) in a study carried out on 60 pregnant women in Alexandria, and in the Hyperglycemia and Adverse Pregnancy Outcome (HAPO) Study Cooperative Research Group (30), it was reported that serum C-peptide and insulin levels were significantly lower in women with severe preeclampsia than in women with mild preeclampsia and normotensive pregnant women. Similarly, Valensise et al. (31) reported higher insulin levels, but not higher C-peptide values, in pregnant women undergoing oral glucose tolerance tests who develop gestational hypertension. These disagreements could be due to the difference in the methods used to assess serum C-peptide levels and the involvement of different ethnic groups in the different studies.

Currently, the possible role of the C-peptide in PE has not yet been clarified. PE is a hypertensive disorder that occurs mainly due to placental maladjustment, which involves endothelial dysfunction, which consistently affects the kidney producing proteinuria (32), morphological evidence of glomerular endotheliosis (33),

decreased renal plasma flow (RPF) and glomerular filtration rate. Therefore, the increased C-peptide levels in PE could be associated with its renoprotective effect, as its beneficial actions have been described in T1DM complications since it is capable of improving glomerular hyperfiltration and hypertrophy, and proteinuria (34-36), and by improving erythrocytes deformability in T1DM patients by increasing Na⁺/K⁺ATPase activity (37,38). Indeed, acute or long-term infusion of C-peptide has been shown to reduce early disturbances of diabetic nephropathy such as hyperfiltration, glomerular hypertrophy, and microalbuminuria. Likewise, in a randomized study in 21 normotensive patients with T1DM and incipient nephropathy, who received combined treatment with C-peptide and insulin for 3 months, it was shown a reduction of albuminuria, indicating that C-peptide can improve renal function in patients with type 1 diabetes mellitus (39). In addition, it was shown in experimentally induced diabetic nephropathy that C-peptide administration reverted the higher fasting serum glucose and lower insulin levels and decreased renal injury markers such as serum urea, creatinine, tumor necrosis factor-alpha (TNF- α), angiotensin II, malonyl dialdehyde, total antioxidant, Bcl-2, and NO in renal tissue. Beneficial effects of C-peptide were partially antagonized by L-NAME, indicating that NO partially mediates C-peptide effects, and suggesting that C-peptide deficiency could contribute to renal and metabolic alterations in T1DM (40).

C-peptide exerts important physiological effects on different cells such as endothelial, beta-pancreatic, renal tubule, and fibroblast cells (16,25). There is evidence indicating that plasma C-peptide concentrations provide an indirect measure of insulin secretion reserve (41,42), and have been suggested as an indicator of pancreatic beta-cell function (43). Therefore it has increased the interest in this peptide due to its unique property as an independent marker of insulin biosynthesis and secretion, because unlike insulin, C-peptide is subjected to an insignificant liver first-pass metabolism, making it a special diagnostic tool in diabetes (44). Insulin resistance is known to be a common feature linking

hypertension and hyperinsulinemia (45). Indeed, in cross-sectional studies in fasting patients in the third trimester of pregnancy (46), with postprandial hyperinsulinemia (47), with insulin resistance (48), are conditions identified as being associated with high blood pressure in late pregnancy. Several lines of evidence suggest that PE may be associated with higher degrees of insulin resistance than those that characterize normal pregnancy. Postulated mechanisms, through which insulin resistance could increase blood pressure in pregnancy, as occurs in essential hypertensive patients, include activation of the sympathetic nervous system (49,50), renal sodium retention (51), increased cation transport (52) and association with endothelial dysfunction (53). Similarly, it has been reported that there are higher risks of PE with various conditions associated with insulin resistance including gestational diabetes, polycystic ovary syndrome, obesity, and weight gain (54).

Even though serum C-peptide has been established as a marker of insulin resistance and obesity, specifically in T2DM and basal levels of C-peptide are significantly elevated in patients with metabolic syndrome and diabetes (55), the most recent studies have shown that C-peptide serum levels are strongly and positively associated with stroke events, independent of serum insulin level in diabetics (56). In this sense, it has been postulated that C-peptide levels constitute a better predictor of total mortality or mortality related to cardiovascular events, than serum insulin itself and other measurements derived from insulin resistance in non-diabetic individuals; however, it is not clear what the mechanism of such association (57). Likewise, C-peptide has been postulated as a risk factor for cardiovascular disease (CVD) since its serum levels were significantly and negatively associated with serum levels of high-density lipoproteins (HDL-C) in individuals without diabetes, indicating that C-peptide serum levels are associated with CVD death whose cause, at least in part, may be due to low HDL-C levels (56,58). In a study by Cabrera et al. (59) in which they analyze the association between serum C-peptide and coronary heart disease, it was shown that elevated C-peptide is associated with the incidence of myocardial infarction and coronary heart disease in the general population, in consequence, it could be a

predictor of coronary events earlier than impaired fasting glucose. Furthermore, evidence-based clinical, epidemiological, and experimental studies have shown that hyperinsulinemia is associated with different cardiovascular risk factors, such as hypertension (60). Indeed, it is known that hyperinsulinemic and insulin-resistant individuals have increased sympathetic nervous system activity and salt sensitivity, factors which increase the probability that they will develop hypertension. Therefore, since elevated insulin and C-peptide levels coexist in PE, C-peptide may exhibit an insulin-like mechanism promoting hypertension during pregnancy (11). That is why pregnancy provides an opportunity to examine the association between increased beta-cell secretion and high blood pressure in subjects with no history of previous hypertension. It could then be inferred that the increase in C-peptide secretion and therefore its circulating levels described in the present study, may constitute a compensatory response to the increase in blood pressure during the last trimester of pregnancy, so its measurement could be an early marker of risk for pregnancy-induced hypertension.

In conclusion, we report that a population of Venezuelan pregnant women with 28-40 weeks of gestation presents higher serum C-peptide levels when compared with normotensive pregnant women. These findings support the concept that serum C-peptide levels could serve as an early predictor of preeclampsia and insulin resistance in pregnant women. However, large-scale and prospective studies are required to assess whether the association between increased blood pressure, proteinuria, and increased C-peptide levels are early markers to predict the development of PE.

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CONFLICT OF INTEREST

The authors have no conflict of interest.

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