Maternal inherited thrombophilia and pregnancy outcomes

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Abstract. Thrombophilia is a group of genetical disorders that cause blood to clot abnormally. Thrombophilia is linked to recurrent pregnancy loss, foetal growth restriction, late miscarriages, stillbirth and preeclampsia. Clinicians usually apply the term thrombophilia only to patients with atypical thrombosis. A successful outcome of pregnancy requires an efficient uteroplacental circulation. Since this system may be compromised by disorders associated with a prothrombotic state, it was postulated that maternal thrombophilia might be a risk factor for preeclampsia and intrauterine growth retardation. The study included 459 pregnant women with gestational ages ranging from 14 weeks to 28 weeks and the patients in the study were tested for hereditary thrombophilia. The type of thrombophilic mutation most common found was the MTHFR mutation (25.7%), followed by the prothrombin gene mutation (20.9%) and the Leiden factor V mutation (15.7%). Also 15.03% patients had been diagnosed with preeclampsia and 6.75% of the pregnant women had IUGR fetuses.

Introduction

Thrombophilia is defined as a predisposition to thrombosis, being a pathology associated with increased tendency of venous thromboembolism. Thrombophilia contributes to more than half of thromboembolic events during pregnancy (1).

Thrombophilia is not a disease itself, but is an important risk factor for thrombosis. Thrombophilic defects can be

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accompanied by the following manifestations (2): Repeated miscarriages, intrauterine growth restriction, preeclampsia, HELLP syndrome and neonatal fulminant purple.

Fetal growth restriction may be linked to preeclampsia but is also associated with inherited thrombophilia. Multiple studies have reported an association between thrombophilia defects and adverse pregnancy outcomes, including both preeclampsia and intrauterine growth restriction. It is hypothesized a maternal predisposition to clotting would lead to thrombosis in the placental vasculature, thereby restricting oxygen and nutrient exchange, resulting in fetal growth restriction (3).

Preeclampsia plays an important role in maternal and fetal morbidity and mortality. Although the etiology of preeclampsia remains unknown, it has been suggested that abnormal placentation and endothelial cell dysfunction are key features in the pathogenesis of preeclampsia (4).

The most common maternal indication for early termination of pregnancy is preeclampsia, the most common fetal indication for early termination of pregnancy is intrauterine growth restriction accompanied by acute fetal distress (5).

Scientists are investigating thrombophilias at a molecular level, over the past 40 years many investigations have been conducted in relation to this condition. Multiple studies have been done in different types of populations to understand the inheritance patterns and risks for individuals diagnosed with an inherited thrombophilia (6). Familial thrombosis was initially considered an autosomal dominant disorder with varying penetrance. Nonetheless, recent studies suggest that congenital thrombophilia may be the result of the combination of two or more gene defects in a family (7).

American College of Obstetricians and Gynecologists (ACOG) recommends that screening for thrombophilia should be performed on patients with a personal history of venous thromboembolism that occurred during a transient risk factor (i.e., pregnancy, surgery, prolonged immobility); patients presenting with thrombosis at a young age (<40 years), patients from families with a history of thrombosis (>2 members), thrombosis at unusual site, more than 3 miscarriages, late miscarriage and foetal death (8).

Inherited thrombophilias include factor V Leiden mutation, prothrombin G20210A mutation (also referred to as factor II

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mutation), protein C deficiency, protein S deficiency, antithrombin deficiency and methylenetetrahydrofolate reductase mutations (9). Thrombophilia can lead to pregnancy complications, including miscarriage, IUGR and stillbirth (10,11).

Patients and methods

A retrospective 6-month cohort study was conducted in University Emergency Hospital of Bucharest (Bucharest, Romania) between June-December 2018. We included in the study 459 pregnant women with gestational ages ranging from 14 weeks to 28 weeks. All the patients included in the study were tested for hereditary thrombophilia and laboratory samples, protein C, protein S, antithrombin III and homocysteine. Genetic analysis collected included mutations of factor V Leiden, gene MTHFR, mutation of factor XIII and prothrombin G20210A gene mutation.

This study was approved by the Ethics Committee of the University Emergency Hospital of Bucharest and informed consent was obtained from all the patients.

Statistical calculation of value. Statistical data collection and processing of data were performed in SPSS version 21. Depending on the typology of the data, coding, analysis of the data and the application of the statistical tests were performed. Also Cramer V (Cramér's phi) test and cluster analysis were performed. Cramér's V varies from 0 (corresponding to no association between the variables) to 1 (complete association) and can reach 1 only when each variable is completely determined by the other.

Based on the literature, the interpretation of Cramer V values is described in Table I.

The objective of cluster analysis is to find similar groups of subjects, where 'similarity' between each pair of subjects means some global measure over the whole set of characteristics (Table I) (12).

Results

The presence of thrombophilic mutations are the result of laboratory analyzes and the normality interval recorded depends on the laboratory reagents used so we opted to investigate the actual values of protein S and protein C by cluster analysis. Thus, using cluster analysis, we classified thrombophilic mutation levels into 3 categories, deficiency, normal and excess.

The average age of patients included in the study was 33 years (\pm 5.20), the average weight was 68 kg (\pm 12.42), the average height was 165 cm (\pm 10.20) and the average body mass index was 25.20 (\pm 4.62).

The average value of protein S was 50.57% (±13.68), with a variation between 20.1 and 111.8. According to the cluster analysis, the number of patients with protein S deficiency was 68, the number of patients with normal level of protein S was 230, and the number of patients with protein S excess was 161 (Table II).

The average value of protein C was 114.37% (±69.52), with a variation between 7.5 and 1078.0. According to the cluster analysis, the number of patients with protein C deficiency was 32, the number of patients with the normal level of protein C was 380, and the number of patients with protein C excess was 47 (Table III).

Table I. The interpretation of Cramer V values.

Value	Interpretation	
0	No association	
1	Perfect association	
<0.10	Poor association	
0.10-0.30	Weak to moderate association	
0.30	Moderate association	
0.30-0.50	Moderate to strong association	
0.50	Strong association	

Table II. Classification of the protein S level according to the cluster analysis.

Category	Frequency	Percentage (%)
Valid data		
Deficit	68	14.8
Normal	230	50.1
Excess	161	35.1
Total	459	100.0

Table III. Classification of the protein C level according to the cluster analysis.

Category	Frequency	Percentage (%)
Valid data		
Deficit	32	7.0
Normal	380	82.8
Excess	47	10.2
Total	459	100.0

In the study group, 15.03% patients had been diagnosed with preeclampsia and 6.75% of the fetuses had intrauterine growth restriction (Table IV).

Fetal growth restriction is a condition in which fetus is smaller than expected for gestational age. It is described as an estimated weight lower than the 10th percentile.

The type of thrombophilic mutation most common in the study group was the MTHFR mutation (25.7%), followed by the prothrombin gene mutation (20.9%) and the Leiden factor V mutation (15.7%) (Table V).

Determination of the correlation and risk between high risk thrombophilia and IUGR. In this study only data with statistical significance (P<0.05) were used, and Chi-square test was employed to ascertain the association.

The results showed that the value of the Cramer V coefficient reveals an association of moderate to strong intensity between the presence of prothrombin mutation and the presence of intrauterine growth restriction (Cramer's V=0.33, P<0.001). Patients who had prothrombin mutation had a 11.69

Table IV. Distribution of obstetric pathology.

Obstetric pathology	Frequency	Percentage (%)
No medical history	359	78.21
Preeclampsia	69	15.03
Intrauterine growth restriction	31	6.75

Table V. Thrombophilic mutations in this study group.

Thrombophilic mutations	Frequency	Percentage (%)
Protein S deficiency	68	14.8
Protein C deficiency	32	7.0
Factor V Leiden	72	15.7
Prothrombin G20210A	96	20.9
MTHFR mutations	118	25.7
Hyperhomocysteinemia	48	10.5
Antithrombin deficiency	47	10.2
Factor XIII deficiency	59	12.9

Table VI. Association between the presence of prothrombin mutation and the presence of intrauterine growth restriction.

		95% confidence level	
Items	Value	Minimum	Maximum
Odds ratio for IUGR (yes/no)	11.694	5.176	26.418
Total patients	459		

Table VII. Association between antithrombin deficiency and the presence of intrauterine growth restriction.

		95% confidence level	
Items	Value	Minimum	Maximum
Odds ratio for IUGR (yes/no)	60.373	23.561	154.696
Total patients	459		

higher risk of having a fetus with intrauterine growth restriction, compared with pregnant women who did not have this thrombophilic mutation (Table VI).

The results showed that the value of the Cramer V coefficient reveals a strong association between antithrombin deficiency and the presence of intrauterine growth restriction (Cramer's V=0.59, P<0.001). Patients with antithrombin deficiency have a 60.37-fold increased risk of having intrauterine growth restriction compared to pregnant women without thrombophilic mutation (Table VII).

Discussion

Proteins C and S are two vitamin K-dependent plasma proteins and are part of the natural anticoagulant system. Many pregnant women deficient in proteins C and S have been described and have an associated thrombotic tendency, but not all of them will experience thrombotic complications (13). In this study, we obtained mostly normal values of proteins C and S. However, 14.8% (68 pregnant women) were diagnosed with protein C deficiency, and 7% (32 patients) were diagnosed with protein S deficiency.

Intrauterine growth restriction (IUGR) and preeclampsia are an important cause of fetal and neonatal morbidity and mortality. Some studies showed association between inherited thrombophilia and complications, such as interauterine fetal death, preeclampsia and placental abruption but association between IUGR and thrombophilia is still controversial (14).

Hypertensive disorders of pregnancy, including preeclam psia, consist of a spectrum of conditions which are associated with an important maternal and feta morbidity and mortality (15). The incidence in general population is estimated to be between 3 and 10% of all pregnancies (16,17). In this study 15.03% of the patients were diagnosed with preeclampsia. The results from this study indicate that 15.03% patients had been diagnosed with preeclampsia and 6.75% of the fetuses had intrauterine growth restriction. The latest studies have suggested an important role of maternal thrombotic disorders in complications such as preeclampsia, intrauterine fetal death and IUGR.

In conclusion, we consider that thrombophilia is not a disease itself because a disease manifests when the symptoms of the disease begin, not when a diagnosis occurs.

Thrombophilia remains a pathological condition caused by a combination of risk factors. Thrombophilia refers to disorders which are associated with a persistent hypercoagulable state and a tendency towards thrombosis. Severe pregnancy complications such as preeclampsia and intrauterine growth retardation has been shown to be associated with thrombophilia.

Unfortunately, there is no curative treatment for thrombophilia, only prophylactic treatment with anticoagulants so pregnant women diagnosed with hereditary thrombophilia can be considered patients with high-risk pregnancies.

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Availability of data and materials

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

DIV, LVA and DMG collected, analyzed and interpreted the patient data regarding pregnancy and thrombophilia. OM, FG, REB and MMC contributed substantially to the conception of the study and had a major role in writing the manuscript. All authors read and approved the final version of the manuscript.

Ethics approval and consent to participate

The study was approved by the Ethics Committee of the University Emergency Hospital of Bucharest (Bucharest, Romania) and informed consent was obtained from all the patients.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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