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Research Article

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ASSOCIATION ASSESSMENT OF THROMBOPHILIC GENE MUTATIONS IN KASHMIRI WOMEN WITH RECURRENT MISCARRIAGES

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ABSTRACT

Aim: To access the prevalence of factor V Leiden and prothrombin G20210A mutations in Kashmiri women with recurrent miscarriage in comparison to control group of parous women. **Method:** FVL and prothrombin G20210A mutations were analysed in 100 women with RM in comparison to 100 parous women. The mutations were analysed using ARMS- PCR approach. **Result:** All the recurrent miscarriage cases and controls were found to have FVL *GG* genotype and prothrombin 20210 *GG* genotype. There was no association of these mutations with the development of recurrent miscarriage. **Conclusion:** We conclude that FVL and prothrombin G20210A mutations do not contribute to the pathophysiology of recurrent miscarriage in Kashmiri

women. Therefore, its screening is not indicated as an initial approach in Kashmiri women with RM.

KEYWORDS: Factor V Leiden; Prothrombin G20210A; Recurrent miscarriage; Kashmir.

INTRODUCTION

Recurrent miscarriage (RM) is defined as the loss of pregnancy as three early consecutive losses (<12 weeks gestation) or two late pregnancy losses (>12 weeks gestation). RM is recognised as one of the most disturbing health issues in women with a worldwide incidence of one per cent of women in the reproductive age group (Stirrat GM, 2009; Clifford K et al, 1997; Maconochie N et al 2004). Several causative factors have been implicated in RM including chromosomal abnormalities of parent or fetus, anatomical malformations, autoimmune disorders, endocrine disorders, infections and other metabolic disorders

(Makino T et al 1990; Stephenson MD 1996; Rai R et al 1997; Lashen H et al 2004; Salim R et al 2003). However, the cause of 50% of RM cases still remain unexplained (Clifford K et al, 1997; Jauniaux E et al 2006; Yang C et al, 2006).

Thrombophilia has been indicated an the main risk factor for maternal thromboembolism thereby contributing to adverse pregnancy outcome especially recurrent pregnancy losses (RPL), pre-eclampsia, abruption placentae and intrauterine growth restriction (Brenner B et al 1999; Visser J et al 2011; Colman-Brochu S, 2004; Abbate R et al 2003). A number of studies have focused on a higher prevalence of certain inherited thrombophilias, such as factor V Leiden (FVL) and prothrombin G20210A mutations in women with unexplained recurrent pregnancy losses (Kupferminc MJ et al 1999; Rey E et al 2003; Roque H et al 2004; Jivraj S et al 2004; Robertson L et al 2006; Glueck CJ et al 2008). However, these reports have found conflicting results (Coppens M et al 2007; Dilley A et al 2002; Finan RR et al 2002; Foka ZJ et al 2002). This heterogeneity has been also reflected in many existing meta-analyses (Rey E et al 2003; Robertson L et al 2006; Kist WJ et al 2008; Kovalevsky G et al 2004). The purpose of this study was to study the prevalence of FVL and prothrombin G20210A mutation in Kashmiri women with a history of more than two miscarriages as compared to ethinicity matched healthy parous women controls.

MATERIALS AND METHODS

Participants

Our study comprised 100 consecutive women with unexplained recurrent miscarriages referred to our department by TRUST Maternity Hospital and Lal Ded Maternity Hospital, Kashmir. The inclusion criteria for the patients were those having at least two recurrent pregnancy losses and those women between 18-40 yrs of age. The women with anatomical defects of uterus, primary or secondary anti-phospholipid syndrome, previous history of thromboembolism, hormonal imbalance, hypertension, diabetes and thyroid problems, overt malignancy and abnormal self or parental karyotype were excluded from our study. The control group consisted of 100 healthy age matched parous women with at least one live child and without any history of miscarriages or gestational complications. The inclusion and exclusion criteria of patients and controls is given in **Table 1**.

Ethics

The study was approved by the ethical committee of the institute. Informed consent was obtained from both patients and controls to participate in the study.

Sample Collection and processing

About 2-3 ml of blood was collected in ethylenediamine tetraacetic acid EDTA containing vials from both patients and controls. DNA was isolated from the blood using a Zymogen (Irvine, CA, USA) DNA extraction kit. The quality of the extracted DNA was accessed by agarose gel electrophoresis. The extracted DNA was stored at 4°C.

Molecular Analysis

Factor V Leiden G1691A and Prothrombin G20210A mutation detection

Molecular diagnosis of FVL and prothrombin G20210A mutation was performed by using amplification refractory mutation system (ARMS). The PCR primers were designed based to amplify the sequences containing the position 1691 of the factor V gene and position 20210 of the prothrombin gene. The primers used in the PCR amplification reactions include a common primer and normal allele specific primer or a mutation specific primer. The primer sequences are given in **Table 2**. A 150bp fragment form exon 10 of Factor V gene surrounding nucleotide 1691 and 350bp fragment from PT gene surrounding nucleotide 20210 was amplified.

PCR programme

The PCR was carried out in a final volume of 30ul mixture containing 1X PCR buffer (Biotools), 0.2 mM dNTP mixture (biotools), 150 ng each primer (Sigma), 1U Taq DNA polymerase (Biotools 5U/μl) and 200 ng genomic DNA (0.2 μg/μl). Amplification was done at at 94 for 7 min, 30 cycles of 94°C for 30 s, X°C for 30 sec min, 72°C for 30sec min followed by extension at 72°C for 7 min (**Table 2**). The PCR products were visualized on a 2-3% agarose gel containing 0.5μg/ml ethidium Bromide and photographed.

Statistical analysis

Statistical analysis was performed by using Chi-square-testing and Fisher's exact test. A value of p < 0.05 was considered significant.

RESULTS

A total of 100 women with > 2 consecutive miscarriages were evaluated for FVL G1691A-and prothrombin G20210A mutations in an age controlled study with 100 healthy parous women who had >1 live birth without pregnancy loss. Those RM patients who were <18 or >40 yrs of age or had other gestational complications like anatomical defects of uterus, anti-

phospholipid syndrome, abnormal karyotype, malignancy, diabetes, or thyroid problem were excluded from this study.

In our study all the 100 RM cases and 100 controls were found to be GG homozygous. No association was found between the FVL mutation and prothrombin G20210A mutation and RM was found (**Table 3**).

Table 1: Inclusion and exclusion criteria of patients and controls.

	RM Cases	Controls		
Inclusion criteria	> 2 consecutive miscarriages	>1 live birth without		
inclusion criteria	> 2 consecutive iniscarriages	pregnancy loss		
Exclusion Criteria	<18 > 40 yrs of age. Anatomical defects of uterus. Anti-phospholipid syndrome. Abnormal Karyotype. Malignancy. Diabetes, Thyroid, Hypertension	Previous miscarriages or gestational complications		

Table 2: Primer sequences for FVL and prothrombin gene amplification.

Mutation	Primer Sequence	Annealing temperature	Amplicon Size
FVL 1691G/A	(C): 5'-GGA CTA CTT GAC AAT TAC TGT TCT CTT G-3'	56 ⁰	150bp
	(N): 5'-GCA GAT CCC TGG ACA GAC G-3' (M): 5'-GCA GAT CCC TGG ACA GAC A-3'	30	
PT 20210 G/A	(C): 5'-TCTAGAAACAGTTGCCTG GCAG-3' (N): 5'-GCACTG GGA GCA TTG AGG ATC-3' (M): 5'-GCA CTG GGA GCA TTG AGG ATT-3'	60 ⁰	340bp

Table 3: Allelic and genotypic frequency of FVL G1691A and prothrombin G20210A mutation in RM cases and healthy controls.

Group	No	Factor V Leiden G1691A G > A genotype			P value	Allele Frequency		P value
		GG	GA	AA		G	Α	
Control	100	100	0	0	1	200	0	1
RM	100	100	0	0	1	200	0	1
		Prothrom	bin G202					
Controls	100	100	0	0	1	200	0	1
RM	100	100	0	0		200	0	

DISCUSSION

In current study we analysed the frequency of FVL and prothrombin gene mutations in 100 RM cases compared to 100 parous women. Our data shows both the mutations to be absent in

both RM cases as well as healthy controls. Our results do not establish an association between recurrent miscarriage and the thrombophilia-related variants. Previously we have found FVL and prothrombin mutations to be present in arterial stroke patients from Kashmiri population (Mahrukh et al 2017), but their frequency in healthy controls including parous women and those with RM is zero. It is well documented that FVL and prothrombin mutation is completely absent or extremely rare in patients of African, Japanese and South East Asian descent (Seligsohn U et al 2001). Recurrent miscarriages associated with FVL and prothrombin mutations are thought to be as a result of thrombosis in utero-placental vasculature. Several studies have reported an association between inherited thrombophilias and increased pregnancy complications, such as severe preeclampsia, fetal growth restriction, stillbirth and abruptio placentae (Kutteh WH et al 2006; Onderoglu L et al 2006, Coulam CB et al 2006; v et al 2006). The heparin thromboprophylaxis during pregnancy has been observed to lead to higher live birth in women with previous history of adverse pregnancy outcomes and thrombophilic defects (Brenner et al 2000: Younis et al 2000).

The FVL mutation (G to A substitution at position 1691 of the FV gene) prevents the inactivation of coagulation factor V by activated protein C, thereby leading to a state of hypercoagulability. FVL which is a significant increased risk factor for systemic thrombosis has also been associated with placental thrombosis (Rai et al 1996; Dizon et al 1997). The placental infarction was more often seen in fetus with FVL allele as compared to normal allele (Dizon et al 1997). A lower live birth rate (38%) was reported in women who had the FVL mutation compared to those who have a normal factor V genotype (69%) (Rai et al., **2002).** A histologically proven placental infarction has been reported in association with fetal carriage of FVL (Dizon-Townson et al., 1997). The impact of thrombophilic mutations on RM across different populations remains a controversial issue. While some studies have shown the association between recurrent pregnancy loss and FVL mutation from Brazil (Souza SS et al 1999), Israel (Brenner B et al 1999), Sweden (Wramsby ML et al 2000), Austria (Hopmeier P et al., 2008), USA (Glueck CJ et al. 2008), UK (Dawood F et al. 2007), Lebanon (Finan et al. 2002) and Tunisia (Mahjoub T et al 2005; Zammiti W et al 2006) while other studies failed to demonstrate an association between Factor V leiden mutation and recurrent miscarriage (Dizon-Townson DS, et al 1997; Pauer HU et al 1998), Alfirevie et al. 2001, Altintas A et al 2007; Sotiriadis A et al 2007; Zahed LF et al 2006). In a study in Asian region including 85 women with RPL and 31 controls, both FVL and prothrombin gene mutation were not found to be associated with recurrent miscarriage (Biswas A et al Mahrukh et al.

2008). This heterogenity in the incidence of the mutation in different populations may be due to racial and ethnic differences in the gene frequency among different populations (Burchard

EG et L 2003).

The prothrombin gene mutation (G to A substitution at position 20210 in the 3'-UTR of the prothrombin gene) is the second most common known inherited risk factor for thrombosis. The prothrombin A20210G has been identified as a risk factor for pregnancy loss in several studies and has been associated mostly to early recurrent pregnancy loss (Martinelli I et al, 2000). Most of the evidence that associates PT G20210A mutation to pregnancy failure originates in case-control studies (Finan RR et al 2002; Foka ZJ et al 2002; Reznikoff-Etievan MF et al 2001). Nevertheless, in most investigations, this relationship is consistent and, as described for FLV, is weaker with 1st trimester compared with 2nd trimester RM (Rev E et al 2003; Robertson L et al 2002;, Kovalevsky G et al 2004)]. Earlier in a study the prevalence of prothrombin G20210A was higher in women with both embryonic (17%) and fetal (16.9%) losses compared with controls (3%) (**Ivanov PD et al 2009**).

CONCLUSION

This is the first study conducted in Kashmiri population that has analyzed the role of thrombophilic markers on the pathogenesis of RM. The results obtained in this study are in accordance with the results of most of the previous research across the globe and indicate that the FVL and prothrombin G20210A mutations are not associated with recurrent miscarriage in Kashmiri women population. We suggest the thrombophilic gene screening should not be an initial approach in Kashmiri women with RM. Also large-scale studies should be performed to clarify the association between thrombophilic gene variants and RM.

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