

ORIGINAL RESEARCH

Role of MMP9 and TIMP2 Gene Polymorphisms in Varicose Veins Formation in the Population of North India

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ABSTRACT

Aim and Objective: To find out Role of MMP9 and TIMP2 gene polymorphisms in the varicose veins formation in North Indian population. **Methods:** A total of 600 (300 cases and 300 controls) individuals were selected for the study, who had diagnosed for varicose veins complication from January 2020 to July 2020 at Government Medical College, Jalaun (Orai) UP, (India). Cases and controls age was between 10-60 years. Fresh 5 ml of EDTA blood was drawn for genetic analyses from both patients and controls as per standardized protocols. For the identification of MMP-9 and TIMP-2 genes polymorphism, DNA were isolated from the fresh blood by using Qiagen kit protocol. PCR-RFLP was performed to determine genotype variants. **Results:** The 50% males and 50% females were selected for this study. 59 (19.66%) cases were reported those had family history of varicose veins formation while it was found quite less 8 (0.02%) in the controls with OR=44.22, 95% CI=14.78, p-value <0.0001. In the PCR-RFLP the distribution of genotypes frequencies in the cases and controls groups it has found that gene frequencies of MMP-9 genotypes have not found statistical significance, whereas the gene frequency of TIMP-2, GG (homozygous) genotype was found statistically significant which was higher in the controls than in the cases. **Conclusion:** To our knowledge, this is the first study to report on MMP-9 and TIMP-2 gene polymorphisms in relation to varicose veins formation in north Indian population. In our sense, such type of studies are required to know the strength and exact nature of the genetic variation to know which of the variant within a haplotype cluster could be functionally related to varicose veins formation in the populations. This study may have demonstrated a statistically significant result in the studied population in large group.

Key Words: Varicose vein formation, Matrix Metallo-proteinase-9, Tissue Inhibitor of Matrix Metallo-proteinase-2, Gene polymorphism, Genotypes

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INTRODUCTION

Development of varicose veins and persistent venous shortage are the most common disorders of the lower limbs. It is more common in females than males of human being.¹ There are several factors are associated for this disease such as old age, ligamentous laxity, prolonged standing, pregnancy, obesity, leg trauma and venous thromboses.²⁻⁷ There are few more risk factors involved to cause this disease such as family history and genetic abnormalities for varicose vein formation. The risk of developing varicose veins has been proven for an individual is around 90% when parents suffered from varicose disease. The probability of this disease is around 20% when parents are normal.⁸ The inner wall of blood vein is

known as tunica intima which is made up of smooth endothelial lining surrounded by an extracellular matrix (ECM) consisting of elastin, collagen and proteoglycans. The elastin and collagen are important structures and responsible for maintaining the integrity of vein wall. This layer has also contains valves to keep the blood flowing in a single upward direction towards the heart.^{9,10} There are four mechanisms are known which may be accountable for the patho-physiology of varicose vein formation. First one is due to loss of elasticity of the vein wall, mainly due to increasing age. The second one is due to absence or non functioning of the venous valves. In the third case it may be due to dysfunctioning of muscular pumping system. In the fourth cases it may

be due to the difficulties of the venous system. All these obstructions may cause venous hypertension and dilated veins in the lower extremities.^{10,11} Matrix Metallo-proteinases (MMPs) is a group of more than 24 endopeptidases (enzymes). All the enzymes are neutral and catalyze the degradation of the proteins in the ECM. The degradation of proteins in ECM is a well in order and performs many physiological functions such as angiogenesis, tissue remodeling and wound healing. However, due to over-expression of MMP gene several mul-functioning may be arises *e.g.* varicose veins formation, cancer, autoimmune diseases, cardiovascular diseases etc.¹² One of the most important Matrix metalloproteinases-9 (MMP9) is a member of the gelatinase family called gelatinase B which is responsible to degrades collagen types IV, V, and XI, telopeptides of type I collagen along with elastin.¹³ In begin, MMP9 is produced in inactive zymogenic form and needs enzymatic activation. MMP activity is regulated by α_2 -macroglobulin and Tissue Inhibitors of Matrix Metallo-proteinases (TIMPs).^{14,15} TIMPs are responsible for the control of the activation of the proenzyme and substrate degradation.¹⁶ The TIMP family has four members synthesized in vascular smooth muscle cells, endothelial cells and macrophages.¹⁷ The expression of MMP1, 2, 3, 9 and 13 is increased in patients of varicose veins formation. The key reason of this increase is may be due to an imbalance between the activities of MMPs and TIMPs, by which in ECM accumulated in the vein wall.^{18,19} The genetic polymorphisms of MMPs and TIMPs are may be important determining factors for this imbalance. The aim of this study was to identify and evaluate the genetic polymorphisms of MMP9 and TIMP2 genes in order to varicose vein formation and examine these polymorphisms. The study also focuses to relate MMP-9 and TIMP-2 gene polymorphism with the relation of varicose vein formation of North Indian population.

MATERIALS AND METHODS

This study was performed at Government Medical College, Jalaun (Orai) UP. (India). This medical college has a tertiary health care association. All the reagents and chemicals were ordered from different suppliers such as Sigma Aldrich, Fermentas, Bio-Rad and Qaigen. For the study total 600 samples (300 cases and 300 controls) were considered. Before the collection of samples a written signed consent was taken from all the cases and controls along with a filled questionnaire form. This study was performed between January 2020 to July 2020 and entire protocol was approval by the ethics committee of the Government Medical College, Jalaun (Orai) UP. (India). Cases and controls age was between 10-60 years. Family history record of cases and controls was also collected. The patients were categorized based on clinical, etiological, anatomical and pathophysiological.²⁰ High ligation with stripping

with or without additional individual ligation and excision were performed for symptoms, complications or cosmetic needs. 2.0 ml of EDTA blood was collected for genetic analyses from each sample.

Sample Size: Sample size was calculated by using below mentioned formula wherein the calculated minimum sample size was found for our study is 150.²¹

$$N = \frac{(Z_{1-\alpha/2} + Z_{1-\beta}) (\sigma_1^2 + \sigma_2^2)}{(\mu_1 - \mu_2)^2}$$

$$\mu_1 = 64.27$$

$$\mu_2 = 70.615$$

$$\sigma_1 = 13.505$$

$$\sigma_2 = 16.36$$

$$Z_{1-\alpha/2} = 1.96 \text{ for } 95\% \text{ confidence,}$$

$$Z_{1-\beta} = 1.84 \text{ for } 80\% \text{ power}$$

Inclusion Criteria: Normal persons above the age of 10 years were recruited as controls. The persons suffering from varicose veins disease with no confounding factors will be recruited as subjects for the study. Individual who needed ligation and excision symptoms were also included as subjects for this study.

Exclusion Criteria: The following exclusion criteria's were needed for the cases and controls as mentioned below.

1. Individuals who had previous history of diabetes mellitus.
2. Individuals known risk of thyroid disease *e.g.* hypothyroidism or hyperthyroidism and taking thyroid medication.
3. Individuals with hormonal replacement therapy, taking vitamin D supplements.
4. Individuals with viral infections, tuberculosis, cancer, mental illness, epilepsy and acute inflammation.

Biochemical Measurement: Control persons were selected in the hospital according standard protocol of WHO *i.e.* fasting plasma glucose <7.0 mmol/l, and history of non management of hypoglycemic record. High ligation with stripping with or without additional individual ligation and excision were performed for symptoms, complications or cosmetic needs.²² The varicose veins disease was confirmed and examined by senior surgeons of surgery department. There were several clinical examinations tested *e.g.* edema and skin trophic modifications on the legs as regular examination. This disease was also confirmed by the department of radiology with the use of duplex ultrasonography scan. A questionnaire was collected from patients who were related to venous complications *e.g.* skin trophic modifications, family history of thromboembolic disease, history of varicose veins, exercise activities and occupations etc. Fresh 5 ml of EDTA blood was drawn for genetic analyses from both patients and controls as per standardized protocols. For the identification of MMP-9 and

TIMP-2 genes polymorphism, first DNA were isolated from the fresh collected blood by using Qiagen kit with following user manual. Purity of DNA was analyzed by running 1% agarose gel and by estimating the 260/280 nm absorbance. Primers were synthesized from Chromous Biotech Pvt. Ltd, Bangaluru. Isolated DNA was amplified with the use of PCR (Bio-RAD, T-100). The PCR conditions were as; initial denaturation for 5 min at 95°C, for 35 cycles, denaturation at 94°C for 30 sec, 56°C for 30 sec, 72°C for 1 min and for final extension at 72°C for 7 min for MMP-9 gene. The only difference in the annealing temperature (55°C for 30 seconds) for the amplification of TIMP-2 gene and rest conditions were similar as MMP-9 gene. Amplified DNA was resolved with 1.0% agarose gel containing ethidium bromide and bromophenol blue. Gel photographs were analyzed with software connected in gel

documentation system (Bio-RAD).²³ For the analysis of restriction fragment length polymorphism (RFLP) of MMP-9 and TIMP-2 genes PAE-I and HGA-I restriction endonuclease enzymes (Fermentas Pvt. Ltd.) were used respectively with following standard digestion protocol.²⁴ Digested fragments were electrophoresis in a 2% agarose gel for allele identification. The statistical analyze was performed by using SPSS software version 20 (USA). Statistical power of the study was analyzed under log-additive model, assuming 10% population possibility by Quanto. Means and standard deviations values were calculated by inverse normal units of the parameters. Allele frequencies of cases and controls of the study population were analyzed by equal opportunity of proportions Z-test. The odds ratio (OR) and 95% confidence interval (CI) were calculated with respect to minor alleles.²⁵

Table 1: Primers oligoes for the Amplification of MMP-9 and TIMP-2 Genes

S.N.	Primer sequences	Tm (°C)
1.	MMP-9 F (5'- GCCTGGCACATAGTAGGCC-3')	59.0
2.	MMP-9 R (5'-CTTCCTAGCCAGCCGGCATC-3')	60.0
3.	TIMP-2 F (5'- GATCCTGTCAGTTTCTCAA-3')	58.5
4.	TIMP-2 R (5'-TTTCCCCTTAGCTCGACTCT-3')	61.0

RESULTS

Clinical and Biochemical Measurement

The study was conducted with the 300 cases and 300 controls of randomly selected varicose veins patients and healthy persons. Demographic data of the patients and controls are mentioned in Table 2. Assessments of age, BMI, systolic and diastolic blood pressures and smoking have showed no statistically significant differences between the groups. The BMI was estimated according to the WHO Asia pacific guidelines as non-obese (BMI <25 kg/m²) and obese (BMI >25 kg/m²). Though, evaluations of gender and family history record have demonstrated a significant difference between groups with significant P value <0.0001. The most of the data analyzed values were shown significantly greater in cases with respect to controls. The cases and controls were 150 (50%) males and 150 (50%). The family history of varicose

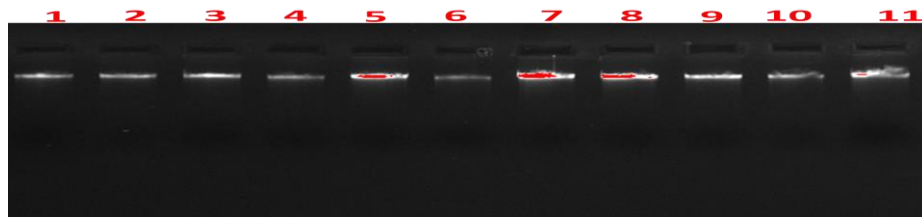
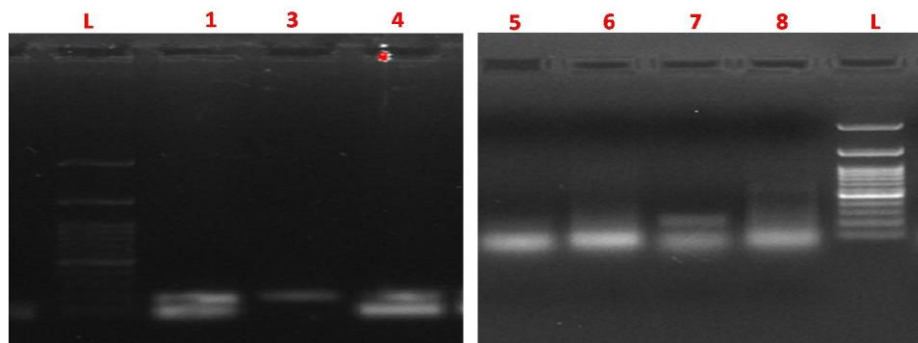
veins were found positive in 59 (19.66%) cases, whereas it was found positive only in 8 (0.02%) in the controls with OR=44.22, 95% CI=14.78, p-value <0.0001). In the Fig. 1, isolated DNA from the cases and controls are presented. In the Fig. 2 and 3 demonstrated the data of PCR-RFLP of the 1562C/T polymorphism of MMP-9 and 418G/C polymorphism of TIMP-2 gene. The genotypic frequency distributions of MMP-9 and TIMP-2 had not illustrated any significant deviation from the predictable Hardy-Weinberg equilibrium in the any studied groups. Table 3 showed the distribution of allelic genotypic frequencies in the cases and controls groups. Frequencies of MMP-9 genotypes have not found statistical significance results between patients and controls, whereas the frequency of TIMP-2, GG (homozygous) genotype was found statistically significant higher in the controls than in the cases.

Table 2: Demographic Data of the Studied Cases and Controls

S.N.	Clinical Characteristics	Cases	Controls	P-value
1.	No. of person involved	300	300	-
2.	No. of male gender	150	150	-
3.	Mean age ±SD (years)	52.36	54.18	0.001
4.	BMI (Kg/m2) ±SD	24.13±1.4	22.42±1.4	0.110
5.	No. of Smoking persons	102	95	0.001
6.	Systolic BP (mmHg) ±SD	120±1.5	119±1.4	<0.005
7.	Diastolic BP (mmHg) ±SD	80±4.0	80±1.7	<0.005
8.	Positive family record	59	08	<0.0001

Table 3: Genotypes frequencies distribution of MMP-9 and TIMP-2 alleles in patients of varicose veins and the control group

Gene	Genotype	Cases (n=300), n (%)	Controls (n=300), n (%)	Adjusted OR	95% CI	p-value
MMP-9 genotype	CC	0.00	0.00	-	-	-
	CT	108 (36%)	60 (20%)	1.326	0.74-1.27	0.756
	TT	192 (64%)	240 (80%)	0.945	0.83-1.13	0.367
TIMP-2 genotype	CC	0.00	0.00	-	-	-
	CG	252 (84%)	196 (64%)	1.201	0.75-1.81	0.401
	GG	48 (16%)	104 (36%)	0.391	0.18-0.84	0.010

**Figure 1: Photograph of Isolated DNA (MMP 9 and TIMP2 gene)****Figure 2 and 3: PCR-RFLP analysis of 1562C/T polymorphism of MMP9 promoter and 418G/C polymorphism of TIMP2 gene promoters. Lane 1-3 corresponded to MMP 9 gene while lane 5-8 corresponded to the TIMP gene, L represented 100bp ladder**

DISCUSSION

In the available literature there are few genetic studies have been mentioned about the MMP-9 and TIMP-2 gene polymorphisms in relation to varicose veins formation. But detailed study with genotypes variation for this region is not studied so far. Varicose veins formation on lower extremity is mainly a cosmetic problem due to poor quality of life as a result of chronic venous shortage. In the present study, genotypes of MMP-9 have not statistically significantly in the patients and controls. MMPs are the main proteinases which humiliating the extracellular matrix.²⁶ MMP-9 plays a crucial role in the digestion of denatured collagens. It has mentioned in the literature that MMPs have been played a big role in many tissues including cardiovascular system. The MMP expression had been increased mechanical pressure in tissues and responsible in varicose veins formation due to amplified venous hydrostatic pressure²⁷ resulting increased hydrostatic pressure in the veins and cell permeability, primarily to leucocyte infiltration, smooth muscle proliferation and finally involved in venous valvular degeneration.²⁸ It has been reported that polymorphisms in the promoter region of MMP-9 is related to the varicose veins in

the Chinese population.²⁹ In the present study identical results have been obtained in relation between polymorphism of MMP-9 and varicose veins formation. There are similar finding had been obtained in Turkish study population.³⁰ The Chinese study explained that those with CC genotype had a greater risk of varicose veins formation, though, in this study none of the cases and controls were carriers the CC genotype. This may be due to the race differences.³¹ TIMPs are small proteins and regulate the action of MMPs and inhibit MMPs activities by binding to their active zinc-binding area.³² In this study, CG (heterozygous) genotypes is responsible for higher risk of varicose veins formation while individuals with the GG genotype may have a lower risk for varicose vein formation. The anti-angiogenic properties of the endogenous TIMPs are well recorded. TIMP-2 plays a key role in the inhibition of endothelial cell proliferation.³³

CONCLUSION

To our knowledge, this is the first study to report on MMP-9 and TIMP-2 gene polymorphisms in relation to varicose veins formation in north Indian population. In our sense, such type of studies are

required to know the strength and exact nature of the genetic variation to know which of the variant within a haplotype cluster could be functionally related to varicose veins formation in the populations. This study may have demonstrated a statistically significant result in the studied population in large group.

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DECLARATIONS

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Conflicts of interest: The authors are thankful to the Government Medical College, Jalaun (Orai) UP (India) for the financial support.

Ethics approval: This study was approved by the ethical committee of Government Medical College, Jalaun (Orai) UP (India).

Consent to participate: We have consent to participate.

Consent for publication: We have consent for the publication of this paper.

Availability of data and material: The entire data and materials were self generated.

Code availability: Not applicable.

Authors' contributions: Sushil Kumar Bundela and Ramesh Kumar Yadav both the authors were equally contributed to the work at Government Medical College, Jalaun (Orai) UP (India).

Conflict of interest: There is no any conflict of interest regarding the publication of this work.

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