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Association of vitamin D deficiency and vitamin D receptor (VDR) gene single-nucleotide polymorphism (rs7975232) with risk of preeclampsia

Asma Aziz^{a,†}, Mohsin Shah^{a,†} (), Sami Siraj^b, Waheed Iqbal^b, Amin Jan^a, Imran Khan^b, Sajjad Ahmed^{a,b}, Salvatore Giovanni Vitale^c and Stefano Angioni^c

^aDepartment of Physiology, Institute of Basic Medical Sciences, Khyber Medical University, Peshawar, Pakistan; ^bDepartment of Pharmacology, Institute of Pharmaceutical Sciences, Khyber Medical University, Peshawar, Pakistan; ^cDivision of Gynecology and Obstetrics, Department of Surgical Sciences, University of Cagliari, Cagliari, Italy

ABSTRACT

Background: Preeclampsia has a multifactorial-yet-elusive etiology. Recent reports suggest a link between preeclampsia and vitamin D (VD) metabolic axis. Genetic variations like single-nucleotide polymorphisms (SNPs) of vitamin D receptor (VDR) gene can alter the metabolic role of VD, which have been shown by several genetic association studies. However, there is discordance among these studies.

Objective: The current study aimed to investigate the association of VDR gene polymorphism (Apal) and VD deficiency with risk of developing preeclampsia.

Patients and Method: In this case–control study, 40 preeclamptic and 40 normotensive pregnant women were compared for VD status and VDR gene polymorphism. Serum 25-hydroxyvitamin-D [25(OH) D] level was determined by enzyme-linked immunosorbent assay (ELISA) and VDR gene polymorphism Apa1 was analyzed by Allele specific polymerase chain reaction (AS-PCR) using sequence specific primers.

Results: Serum levels of 25(OH) D were very low but comparable in both preeclamptic and normotensive pregnant women. The difference between the two groups were not statistically significant (p=.423). VDR gene polymorphism Apal (rs7975232) was found not to have significant association with the risk of developing preeclampsia. The frequencies of wild genotype (GG) in preeclamptic and normotensive women were 27.5% and 22.5% respectively. A total of 25% of preeclamptic women had mutant homozygous genotype (TT) and 17.5% of normotensive women had mutant homozygous genotype. The frequency of mutant heterozygous genotype (GT) in preeclamptic patients was 47.5% and in normotensive women was 60%. The variation of wild and mutant genotypes between the two groups was not statistically significant (p>.05). **Conclusion:** This study showed that VDR gene polymorphism (Apal) and VD deficiency are not associated with the risk of preeclampsia.

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KEYWORDS

Preeclampsia; VDR gene; vitamin D; SNP; Apal

Introduction

Preeclampsia (PE) is a pregnancy complication of second half of pregnancy, labor, or the early puerperium with serious consequences for mother and infant [1]. American college of obstetricians and gynecologists defined PE as a new onset hypertension with blood pressure \geq 140/90 mmHg and proteinuria (\geq 0.3 g per 24 h) after 20 weeks of gestation [2]. PE affects about 3%-7% of women in their 1st pregnancies with a morbidity and mortality rate of 10%-15% [3,4]. However, in developing countries like Pakistan, reported prevalence of PE and eclampsia is around 19% and about 1.12% women dies (1 in 89) due to maternal causes in which major cause of death is PE or eclampsia [5]. Abnormal placentation, intolerance of maternal immune system to placental and fetal antigens, and endothelial cells injury are some of the proposed causes of preeclampsia [6-8]. However, PE is known as the 'disease of hypotheses' regarding its causes and pathogenesis [6-8]. Recent studies have linked low levels of vitamin D (VD) with the risk of PE. Being known for calcium homeostasis, it has also been suggested that VD plays important role in modulating immune and cardiovascular systems during pregnancy that have an impact on blood pressure [9]. Optimum levels of VD promote healthy gestation and fetal growth [10], whereas low maternal VD levels have been reported in PE by various researchers [3,11–13]. Furthermore, studies also reported that VD supplementation lowers the risk of PE [14,15]. However, the recent studies based on VD and PE are conflicting and inconsistent [16,17].

VD exerts its biological action by binding to a high affinity receptor, which is known as VD receptor (VDR). VDR is a nuclear receptor that belongs to the super family of steroid hormone receptors, it functions as a nuclear transcription factor [18]. Since, the discovery of VDR gene, genetic variations in its DNA sequence have been extensively studied and linked to multiple disorders [19] including PE. However, where some researchers have found strong association of single-nucleotide polymorphisms (SNPs) in VDR gene complex with PE [20,21],

CONTACT Mohsin Shah 🖾 mohsin.ibms@kmu.edu.pk 🗊 Department of Physiology, Institute of Basic Medical Sciences, Khyber Medical University, Peshawar, Pakistan and Sami Siraj 🖾 samisiraj.ibms@kmu.edu.pk 🗊 Department of Pharmacology, Institute of Pharmaceutical Sciences, Khyber Medical University, Peshawar, Pakistan 🚯 Supplemental data for this article can be accessed online at https://doi.org/10.1080/09513590.2022.2146089

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others have shown no association [21]. In our previous work, we described the role of neurokinin-B in the pathophysiology of preeclampsia [22]. Here, we intend to find the role of VD and its receptor gene polymorphism in PE in our population. Thus, a case-control study was designed to investigate the level of VD and VDR gene polymorphism in patients with PE compared with healthy individuals and to find out the possible association of VD levels and VDR gene polymorphism with the risk of PE.

Material and methods

Study design and setting

This was a case-control study. A total of 80 subjects (40 preeclamptic and 40 normal pregnant women) were enrolled in the study during their third trimester of pregnancy (gestational age = 34 ± 3 weeks) from obstetrics and gynecology wards of Hayatabad Medical Complex, Peshawar and Saidu Group of Teaching Hospital, Swat. The patients were diagnosed according to American college of obstetrician and gynecology (ACOG) 2002, criteria for preeclampsia [2]. According to this criteria, a women is considered as preeclamptic if she has hypertension, systolic blood pressure (SBP) ≥140 mm Hg or diastolic blood pressure (DBP) \geq 90 mm Hg on 2 occasions >4 h apart after 20 weeks' gestation in a previously normotensive woman and proteinuria ≥300 mg protein per 24-h urine. Informed written consent was taken from all the participants. Subject's demographics were recorded on predesigned proforma and 5 mL of blood was collected from antecubital vein. About 2.5 mL was transferred to gel tube from which serum was extracted for biochemical analysis and remaining 2.5 mL to EDTA tube which was stored for genetic analysis. The experimentation was carried out in the physiology laboratory, Institute of Basic Medical Sciences and the study was approved by the Advanced Studies and Research Board of Khyber Medical University, Peshawar (Study approval No. DIR/KMU-AS&RB/CV/000037).

Inclusion-exclusion criteria

Women suffering from chronic diseases like hypertension, cardiac pathologies, bad obstetrical history, or any fetal anomalies and those who do not want to participate in the study were excluded from the study. The term bad obstetrical history was defined as female who experience any loss in her pregnancy either abortion, still birth, early neonatal death, or demise of the fetus due to any other cause related to pregnancy. Subjects were examined for height, weight, pulse, edema, fundal height (height of the uterus), fetal heart sounds, and SBP and DBP. Obstetrical ultrasound of each patient was also done to exclude any fetal anomalies. Subjects taking VD supplements were excluded from the study. The subjects were also investigated for hemoglobin, blood grouping, and urinary proteins.

Determination of vitamin D

Serum level of 25-hydroxycholecalciferol was measured by enzyme-linked immunosorbent assay (ELISA) kit (Kit Catalogue No. EQ 6411-9601) purchased from Euroimmune Medizinischelabordiagnostika AG. The ELISA kit used solid-phase competitive protein binding assay to determine the level of 25-hydroxycholecalciferol in human serum samples. All the steps of the assay were performed according to the manufacturer's instructions. The results were analyzed by three independent observers apart from the researchers of the study who were blind to the subject status. VD levels >25 ng/mL is the optimal level, severe VD deficiency <10 ng/mL, and insufficiency 10–19 ng/mL [23].

DNA extraction and VDR genotyping

Genomic DNA was extracted from peripheral blood using commercially available PureLink Genomic DNA Mini Kit (Kit Catalog No. K1820-02). The *ApaI* (rs7975232) VDR gene polymorphism was genotyped using allele-specific polymerase chain reaction (AS-PCR). Specific primers (Table 1) were designed for VDR rs7975232 using online tools [24].

The reaction was performed in total volume of 20 µL including 2 µL PCR (10×) buffer, 0.8 µL MgCl₂ (50 mM), 1 µL dNTPs (10 mM), $0.4 \mu \text{L}$ DNA Tag Polymerase $(5 \text{ u/}\mu\text{L})$, $0.6 \mu\text{L}$ $(10 \mu\text{m})$ of each forward, reverse, and allele specific primers, 2 µL (50 ng/ µL) of DNA template and sufficient molecular grade water to make total volume of 20 µL. The reaction mixtures were amplified in automatic thermal cycler with the following conditions; initial denaturation at 95 °C for 4 min followed by 30 cycles of 94°C for 1 min, 64°C for 40s, 72°C for 1 min, and final extension at 72°C for 7 min. The amplicon was electrophoresed on 1.5% agarose gel stained with 2µL ethidium bromide (10 mg/ mL) at 90 mA for 45 min and visualized under ultraviolet (UV) trans-illuminator. Depending upon the presence of G or T specific band along with the control band, genotypes were assigned as homozygous wild (GG), homozygous mutant (TT), and heterozygous mutant (GT) for ApaI polymorphism.

Statistical analysis

All the continuous variables were presented as mean \pm standard deviation whereas categorical variables were expressed in terms of frequencies and percentages. To compare continuous and categorical variables, independent sample *t* test and Pearson's chi square test of independence was applied, wherever applicable. SNP data was calculated in terms of Hardy–Weinberg equilibrium (HWE), where chi-square statistics was applied for genotype and allele frequency analysis. Regression analysis was employed to determine the impact of different variables on disease progression. The odds ratio (OR) with 95% CI was calculated to determine the risk. *p* value <.05 was considered significant.

Results

The mean age of the study population preeclamptic group was 33.05 ± 5.26 years while mean age of control group was 27.55 ± 4.91 years (Table 2). Statistically significant difference was observed in the BMI of disease group compared to control group (p < .001). Similarly, a statistical significant difference was observed in the pulse rate of preeclamptic and normotensive pregnant women (p = .01). Both SBP and DBP were statistically significant between the groups (both p < .001). Similarly, a statistically significant difference was obtained in the fundal height

 Table 1. Allele flanking and specific primer sequences and their respective PCR product sizes.

Control primers	Forward Reverse	CCAAACACTTCGAGCACAAGG AGAGCAGAGTTCCAAGCAGAGG	592 bp
Allele-specific	T-Reverse	GGTGGGATT GAGCAGTGAAGT	312 bp
primers	G-Reverse	GGTGGGATTGAGCAGTGATGG	

 Table 2. Demographic and clinical variables between normotensive (control) and preeclamptic (cases) patients.

Variables	Disease	$Mean \pm SD$	95% CI	p value
Age (years)	Yes	33.05 ± 5.267	3.2-7.7	<0.001
	No	27.55 ± 4.914		
BMI (m/kg ²)	Yes	35.742 ± 4.6465	8.0-11.8	<.001
	No	25.792 ± 3.9651		
Pulse	Yes	80.08 ± 4.196	0.4-3.8	.01
	No	77.95 ± 3.449		
Systolic (mmHg)	Yes	163.98±16.671	42.6-54.3	<.001
	No	115.50 ± 8.149		
Diastolic (mmHg)	Yes	106.25 ± 8.605	31.6-39.3	<.001
	No	70.75 ± 8.883		
Fundal height (weeks)	Yes	34.78 ± 2.587	0.1-2.6	.03
	No	33.40 ± 3.112		
Gravida	Yes	5.50 ± 3.063	1.1–3.4	<.001
	No	3.20 ± 2.151		
Para	Yes	4.025 ± 2.828	1.1-3.2	<.001
	No	1.800 ± 1.771		
Period of gestation	Yes	35.21 ± 2.803	0.5-3.1	.007
	No	33.38 ± 3.094		
Vitamin D (ng/mL)	Yes	6.790 ± 9.151	-5.8 to 2.3	.44
	No	8.535 ± 9.431		
Hb (g/dL)	Yes	11.615 ± 1.387	-0.8 to 0.08	.1
-	No	12.003 ± .600		

Table 3. Frequency and percentage distribution of different study parameters.

Variables		Frequency	Percent
Disease	Yes	40	50.0
	No	40	50.0
Edema status	No pitting edema	36	45.0
	mild pitting edema	17	21.3
	moderate pitting edema	27	33.8
Genotypes	GG	20	25.0
	GT	43	53.8
	TT	17	21.3
Urine albumin	30-100	24	30.0
	100-300	15	18.8
	300-1000	1	1.3
	Nil	40	50.0
BMI categories	Normal	19	23.8
	Over weight	19	23.8
	Obese	42	52.5
Vitamin D levels (ng/mL)	Normal	2	2.5
	Insufficient	8	10.0
	Deficient	70	87.5
Urine puss cells	Normal	62	77.5
	Infected	18	22.5

of the diseased group compared to control group (p=.03). The difference of both gravida and para of the preeclamptic women compared to normotensive control was statistically significant (both p < .001). A similar statistically significant difference was obtained in the period of gestation of preeclamptic when compared to normotensive women (p=.007). The level of VD (p=.4) and Hb (p=.1) between the two groups in this report did not attain statistical significance. All the demographics of the study population are summarized in Table 2.

The frequencies and percentages of different variables in the study population are shown in Table 3. To rule out the edema status, 45% of the total study population were presented with non-pitting edema, 21.3% had mild while 33.8% had moderate pitting edema. Urine albumin was absent in 50% of study cases, 30% had 30–100 mg/g urine albumin, 18.8% had 100–300 while 1.3% have above 300 mg/g urine albumin. Urinary tract infection (UTI) was evaluated based on pus cells in urine, 77.5% had normal urine pus cells while 22.5% were presented with UTI. The BMI of 23.8% individuals were in normal range, while 23.8% and 52.5% cases were overweight and obese respectively.



Figure 1. Genotype distribution among the two study groups.

 Table 4. VDR Apal (rs7975232) genotypes distribution based on different study variables.

	Genotypes Apal (rs7975232)			p values	
Variables	TT (n = 17)	GG (n=43)	TG (n=20)	TT vs TG	TT vs. GG
BMI	32.01±4.9	29.7±6.8	30.7 ± 7.0	.78	.54
Pulse	80.1 ± 4.2	78.4 ± 3.6	78.8 ± 3.9	.46	.36
Systolic	143.2 ± 25.1	140.6 ± 28.4	137.9 ± 28.6	.78	.95
Diastolic	90.2 ± 16.9	90.2 ± 20.2	86.9 ± 20.9	.83	1.0
Fundal height	33.8 ± 2.8	34.3 ± 2.5	34.07 ± 3.1	.97	.90
Gravida	3.8 ± 2.3	4.3 ± 3.4	4.5 ± 2.7	.71	.87
Para	2.6 ± 2.2	2.6 ± 3.0	3.1 ± 2.5	.78	1.0
POG	34.2 ± 3.1	34.4 ± 2.1	34.2 ± 3.2	.99	.99
Vitamin D	3.9 ± 1.8	7.2 ± 6.5	9.3±11.5	.10	.51
Hb	11.7 ± 1.4	12.0 ± 0.9	11.7 ± 0.9	1.0	.69

The levels of VD were normal in 2.5%, 10% were presented in VD insufficiency while 87.5% were VD deficient (Supplementary Table S1). The VDR-GG genotype was 25%, GT was 53.8%, and TT was 21.3% in study population. The overall genotype distribution between disease group and control group is also shown in Figure 1. Similarly, no statistical difference was found between VDR-APA-I genotypes and study variables (preeclamptic vs. control) (p > .05). Levels of VD were low in control group but even lower in disease group though the difference was not statistically significant (Supplementary Table S2). Table 4 summarizes the difference in study variables based on genotypes distribution. To find the impact of genotypes, levels of VD, and BMI on disease, logistic regression analysis was performed as shown in Table 5. The HWE was calculated using web-based online tool. Both the disease group and control group were not consistent with HWE as shown in Table 6. No statistically significant association was found between genotypes and VD levels among disease and control group. However, individual with TG genotypes were 1.8 times more likely to develop PE rather than TT genotypes. Similarly, individuals with insufficient and deficient VD levels were 3.5 times and 2.4 times more likely to develop PE rather than those with normal VD levels. Obesity was strongly associated with the development of PE with p < .001. To ascertain the effect of modifying effect of obesity on genotype association with the disease, multivariate logistic regression was performed using two independent variables, i.e. genotypes and obesity. However, our data suggest that there was no significant influence of obesity on association of VDR genetics with the disease (Supplementary Table S3 and Figure 1).

Discussion

In the present study, we compared VD receptor gene polymorphism (Apa1) and VD status in total of 80 preeclamptic and

 Table 5. Regression analysis of study variables between disease and control group.

Variables	Phenotype	OR (95% CI)	p value
Genotypes	Т	_	_
<i>,</i> ,	G	1.05 (0.56–1.95)	1.0
	TT	_	_
	GG	1.16 (0.31-4.32)	.92
	TG	1.8 (0.57–5.63)	.46
Vitamin D levels	Normal	_	_
	Insufficient	3.5 (0.01–723)	.63
	Deficient	2.4 (0.01-349)	.72
BMI	Normal	_	_
	Overweight	0.2 (0.01-2.2)	.19
	Obese	0.01 (0.001-0.09)	<.001

 Table 6. Hardy–Weinberg equilibrium between disease and control group.

Genotype	De Observe value Predictive		χ2	p value
Cases				
TT	10	9.5	0.09	.75
TG	19	20.0		
GG	11	10.5		
Controls				
TT	07	9.0	1.64	.19
TG	24	20.0		
GG	09	11.0		

normotensive pregnant women of the province of KPK Pakistan. In previous studies, VD has been shown to affect incidence of preeclampsia and VD supplementation also has shown some promising results in various interventional studies [25]. Therefore, a case-control study was designed to address the objective of the present study. However, we found that neither VD level nor VDR SNP ApaI has any statistically significant association with the incidence of preeclampsia.

According to the present study, Apa 1 genotype distribution between preeclamptic and normotensive subjects was homozygous wild genotype (GG), present in 11 preeclamptic (27.5%) and 9 normotensive (22.5%) subjects. Heterozygous mutant genotype (GT) was present in 19 preeclamptic (47.5%) and 24 normotensive (60%) subjects. Homozygous mutant (TT) genotype was present in 10 preeclamptic (25%) and 7 normotensive (17.5%) subjects (Figure 1), showing no statistically significant variation between preeclamptic and normotensive women regarding wild and mutant genotypes (p > .05). Apa-I (rs7975232) is an intron variant and doesn't cause a change in the linear code. However, it does have a documented effect on mRNA stability and, therefore, expression of VDR gene. The phenotypic manifestation of VDR-rs7975232 is not consistent across different descents or ancestries [26]. The present study is in accordance with a previous study conducted by Rezende et al., where they found no correlation between different variants of VD receptor gene (Fok1, Apa1, and Bsm1) and preeclampsia [27]. Similarly, Rezavand et al. found no association of preeclampsia with VD receptor SNP Fok1 [21]. Association between preeclampsia and allelic variant of VD receptor gene rs12831006 was confirmed by Baca et al., while they found no statistically significant association between SNP rs1544410 (Bsm1) and preeclampsia [28]. In the current study, in preeclamptic women minor (GT, TT), genotypes were more prevalent in women with high BMI as shown in Table 4. There was no statistically significant difference between genotypes in terms of anthropometric and various clinical parameters (Table 4).

In both normotensive and preeclamptic groups, the VD levels were lower however, the level of VD was comparable in both preeclamptic and normotensive women (p=.44). As shown in

Table 2, the mean value of VD in preeclamptic patients was 6.79 ng/mL and mean VD level in control group was 8.53 ng/mL. Thus, according to the results of this study 1.3% of the control group was adequate regarding VD status and also 1.3% of the preeclamptic women had adequate VD levels. In control group, 2.5% of the women were deficient and 23.7% were severely deficient regarding VD level, while 5% of preeclamptic patients were deficient and 8.8% were severely deficient. Among these 35% of preeclampsia patients and 18.8% of normotensive women had very severely deficient VD levels as shown in Supplementary Table S1. Thus 93.8% of the study group had VD deficiency. The findings of this study are consistent with a cross-sectional study conducted in Lahore Pakistan [5]. They compared serum VD levels in preeclamptic and normotensive pregnant women and concluded that 95% of women were VD deficient with VD levels comparable in both groups. Another study done on a small group of pregnant women in Karachi Pakistan [29] showed similar results with 90% of pregnant women having VD deficiency.

The high prevalence of VD deficiency in the study group could rise because the pregnant women were not taking VD or calcium supplements and were of low socioeconomic status having poor diet not fortified with VD and calcium supplements. Furthermore, most of these women were housewives who spent most of their time indoor at home and wearing veil and folk weave clothing and thus less exposed to sun. The findings of the current report do not associate low levels of VD with preeclampsia unlike the work done by previous researchers [3,12,30]. Bodnar et al. conducted a nested case-control study and found that 55 nulliparous women who developed preeclampsia had VD deficiency [3]. Similarly, Baker et al. also conducted a nested case-control study and reported that women having VD levels <20 ng/mL had four times greater risk of developing severe preeclampsia [12]. Concomitantly, the result of our study is in agreement with previous studies conducted on vitamin D status and preeclampsia, where they found no association of VD level with preeclampsia [5,31]. Powe et al. conducted a nested casecontrol study and found no association of first trimester total 25-hydroxyvitamin-D (25 (OH) D) levels and subsequent development of preeclampsia [31]. Similarly, Shand et al. conducted a casecontrol study on 221 pregnant women who were VD deficient however, VD deficiency was not correlated with preeclampsia or gestational hypertension [32]. The discrepancy between their and the current study may be due to the different populations that were studied. Another reason of discrepancy might be the measurement of serum 25 (OH) D in third trimester of pregnancy in the present study, while Bodnar et al. assessed vitamin D level at 10 weeks of gestation. The possibility may be that third trimester VD status has no significant influence on placentation and development of preeclampsia as compared to first trimester VD concentrations. The difference between our results and Bodnar et al. may also be due to difference in weight and height measurements [3]. Bodnar et al. calculated BMI from self-reported pre-pregnancy weights and heights while we measured weights and heights at time of sampling. 25 (OH) D levels are more likely associated with BMI at time of measurement than pre-pregnancy BMI. Lower circulating 25 (OH) D levels in obese subjects have been attributed to the deposition of VD in adipose tissue [33]. As the risk of preeclampsia increases in obese pregnant women [34], therefore it is inferred that pre-pregnancy BMI may not be a potential confounding factor as compared to BMI measured at time of 25 (OH) D measurement. The chances of preeclampsia increase with the increase in BMI as shown in our study where 97.5% of the preeclamptic patients had high BMI and only 2.5% had normal BMI. While in control group 42.5% were of normal BMI and 57.5% were of high BMI. According to the

statement of American College Of Obstetricians and Gynecologists (AGOG) Technical Bulletin 219 Washington DC 1996, the risk of preeclampsia increases with high BMI by the ratio of 3:1. Association of two predictor variables, i.e. obesity and genotypes, with preeclampsia was also assessed. However, our data suggest that there was no significant influence of obesity on association of VDR genetics with the disease.

This study has certain limitations. The sample size of our study was not adequate. Therefore, including larger cohort would have represented the population more accurately. The second limitation of this study is that we did not include a control group of non-pregnant women. Inclusion of this third group of normotensive non-pregnant women would have improved our study results. Also, we only considered Apa1 polymorphism while results of the study would have improved if other vitamin D receptor gene polymorphisms (Taq1, Bsm1, and Cdx-2) were considered and their possible interaction with Apa1 evaluated.

Conclusion

This study shows that pregnant women of Pakistani population both normotensive and preeclamptic are very deficient regarding 25 (OH) D levels. Similarly, we observed no significant association of VD deficiency and VD receptor gene polymorphism (Apa1) with preeclampsia. The levels of 25 (OH) D were similar in both cases and controls and also mutant genotype frequency was similar in both groups.

Compliance with ethics guidelines

All procedures involving human participants performed in this study were in accordance to the ethical standards of Khyber Medical University's ethical committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. The study was approved by the institution board of study and informed consents were obtained from all individual participants included in the study.

Author contributions

AA and MS were involved in study design, execution, writing of the manuscript and analysis of the data. AJ helped in recruiting control subjects. SGV and SA helped in the organization of the dataset and in data analysis. WI and SA helped in data analysis as well as experimentation. SS helped in execution of the data and analysis of the results. IK, SGV and SA provided critical discussion. All authors reviewed and approved the final version of the manuscript.

Disclosure statement

The authors report no conflicts of interest.

Data availability

Available from the corresponding author on reasonable request.

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ORCID

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