Seroprevalence of IgG Antibodies for Dengue among Blood Donors

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ABSTRACT

BACKGROUND

Dengue fever is endemic India and the prevalence of dengue is on the rise owing to various social and economic factors. Prevalence of asymptomatic dengue infection varies widely from less than 1 % to 80 % in India. Transfusion transmissible dengue has been reported in different parts of the word. Prevalence of subclinical dengue among blood donor poses a threat to the blood supply leading to transfusion transmissible dengue. We wanted to estimate the prevalence of IgG antibodies for Dengue in the blood donor population.

METHODS

Six hundred and eight whole blood donors were included in the study during the period January 2017 to October 2018. Donor registration and education was done as per the national guidelines. Donors who gave a history of previous dengue or symptoms consistent with dengue were excluded from the study. Serum samples from whole blood donors were tested for IgG antibodies using ELISA technology.

RESULTS

Of the total of 608 donors, 602 were male donors and 55 % of the donors were in the age group 21 - 30 years. Majority (69 %) of the donors were from urban locations. Anti IgG antibodies for dengue were present in 4.14 % of donors of which 38 % of donors were in the age group 31 - 40 years. Three hundred and eighty-four donors revealed history of fever, myalgia and headache in the past one year. No statistical significance was found between fever, myalgia and the presence of IgG anti-dengue antibodies.

CONCLUSIONS

The subclinical or asymptomatic prevalence of dengue infection is low when compared to other studies in other parts of the country. Enquiring into donor history for history of dengue or symptoms of dengue and deferring such donors for a recommended period will prevent transfusion transmissible dengue.

KEY WORDS

Seroprevalence, Dengue IgG Antibodies, Subclinical Dengue, Blood Donors

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DOI: 10.14260/jemds/2020/679

How to Cite This Article: Bandi K, Thokala RP, Anandan A, et al. Seroprevalence of IgG antibodies for dengue among blood donors. J Evolution Med Dent Sci 2020; 9(42):3092-3096, DOI: 10.14260/jemds/2020/679

Submission 05-03-2020, Peer Review 11-09-2020, Acceptance 17-09-2020, Published 19-10-2020.

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BACKGROUND

Dengue is a vector borne disease that is a global threat with endemicity in more than 100 countries and 50 million new infections are being added every year. Disease burden of dengue is on rise in India from 2001, a disease that was initially confined to northern and southern states has spread to many states now. Rise in temperature, urbanisation, changes in environmental factors together with inadequate vector control had contributed to the existing disease burden.¹ The rate of asymptomatic or subclinical infection of dengue varied from less than 1 % to 80 % depending on the population studied. With increasing prevalence of dengue among the population, the blood recipient population is also at risk of acquiring dengue infection that can be subclinical in a majority of donor population. Dengue infection after solid organ transplantation, bone marrow transplantation and blood transfusion has been documented in a few studies.^{2,3}

Arboviral disease transmission through blood products is a well-recognised fact. Spread of West Nile virus and chikungunya virus through blood transfusion has already been documented. The high prevalence of dengue in various regions, asymptomatic infection rate, the period of viremia, the high viral load and wide spread prevalence of the vector had placed dengue in the red category for emerging or re-emerging agents that could pose threat to blood supply in North America by AABB (American Association of Blood Bank).⁴ The potential of dengue to emerge as a threat to blood safety is recognised even in European countries that has not documented autochthonous dengue infection since 1930.5 In India the prevalence of dengue infection varied with population studied with some parts having a prevalence rate of over 50 %.6 And in some parts of the country where the prevalence is close to zero among donor population.⁷ Prevalence of subclinical infection among donor population in middle east has also been observed.8

The disease burden of dengue is higher in India owing to the favourable conditions existing like the tropical climate, changing climate, increased temperature, and abundance of vectors, migrating population and inadequate vector control measures.^{9,10} Dengue infection can remain subclinical and viremia in the subclinical phase can transmit dengue infection through blood collected during this phase.¹¹ Studies on prevalence of dengue in blood donor population are limited from this part of the country, hence this study was done to determine the subclinical prevalence of dengue infection among healthy voluntary blood donors.

METHODS

This cross-sectional study was carried out after Institutional Ethics Committee approval. Serum samples from 608 whole (the sample size included was based on the number of donors with inclusion criteria who walked in to donate blood during the study period and the number of assay kits procured to analyse the samples) blood donors collected during the period January 2017 to October 2018 were included in the study, whole blood donors representing all parts of the state were tested for dengue specific IgG antibodies using ELISA after obtaining informed consent. Voluntary donors walking in to the blood bank were screened using a donor screening questionnaire. Additional information regarding dengue infections was enquired and recorded by the examining medical officer. Donors who revealed history of dengue infection in the past were excluded from the study. Donors who revealed symptoms consistent with symptoms of dengue in the past were also excluded from the study. Donors who revealed history of fever with symptoms not specifically attributable to dengue infection were included in the study.

Serum samples collected in pilot tubes for mandatory transfusion transmissible infection (TTI) screening were used for this study after the completion of mandatory TTI screening. These serum samples were tested for dengue specific IgG antibodies in a fully automated Elisa platform (EVOLISTM, Bio-Rad, using GAC-Capture ELISA kit). All samples were processed within 72 hours of collection and the results were recorded after verifying the validation criteria of a particular run. Results were interpreted according to the manufacturer instructions. If the dengue IgG units is < 9, then the result was interpreted as negative for dengue IgG antibodies, if dengue IgG units is > 11, then the result is interpreted as a positive for dengue IgG antibodies.

Statistical Analysis

Data was analysed using descriptive statistics. The unpaired ttest was employed to analyse continuous variables. Chi-Square Test and Fisher Exact Test was employed for categorical variables. Statistical significance was taken as p < 0.05. The data was analysed using SPSS version 16.

RESULTS

Six hundred and eight donor samples were tested for dengue IgG antibodies. Of the total 608 donors included in the study there were only 6 female donors and 602 male donors. Majority of the donors (55 %) belonged to age group of 21 - 30 years of age (Figure 1). Donors representing urban population were 69 % and rural side were 31 %. Among the total sample population, Dengue IgG antibodies were present in 4.14 % of blood donors and 95.86 % of blood donors were negative for anti-dengue IgG antibodies.



It is evident from Table 1 that most of the IgG positive blood donors were in 31 - 40 years age group (38.46 %) with

a mean age of 29.31 years. In IgG negative group most were in 21 - 30 years age group (56.36 %) with a mean age of 27.45 years. (p = 0.387). The data subjected to unpaired t-test reveals the existence of statistically non-significant association between age distribution and anti-dengue IgG prevalence (p > 0.05).

Age Distributi on – Groups		IgG Positive	IgG Positive %	IgG Negative	IgG Negative %	% of Age Group among Total Donors	
≤ 20 years	98	6	23.08	92	15.81	16.12	
21 - 30 years	336	8	30.77	328	56.36	55.26	
31 - 40 years	134	10	38.46	124	21.31	22.04	
41 - 50 years	32	2	7.69	30	5.15	5.26	
51 - 60 years	8	0	0.00	8	1.37	1.32	
Total	608	26	100.00	582	100.00	100.00	
p value 0.99							
Table 1. Distribution of Study Population According to IgG Positivity and Negativity							

As the number of female's donors were less than 1 %, statistically non-significant association was observed between gender status and anti-dengue IgG prevalence (p > 0.05). It is evident from the area of living status table (Table 2) that most of the IgG positive group subjects (n = 20) belonged to corporation area (76.92 %) and in IgG negative group too majority belonged to corporation area (68.73 %) (p = 0.761). This data subjected to Fishers Exact test reveals the existence of statistically non-significant association between area of living status and IgG study groups (p > 0.05).

Area of Living - Groups	IgG Positive	IgG Positive %	IgG Negative	IgG Negative %	Total	%	
Corporation Area	20	76.92	400	68.73	420	69.08	
Non Corporation Area	6	23.08	182	31.27	188	30.92	
Total	26	100.00	582	100.00	304	100.00	
1	p value			0.761			
Table 2. Distribution of Whole Blood Donors According to Type of Area							

Three hundred and eighty four Blood donors revealed symptoms like fever, myalgia, headache and vomiting in the past one year (Figure 2). These donors did not have serologically confirmed diagnosis of dengue or any other viral or bacterial illness. These donors did not require hospital admission and were treated either by their family physician on outpatient basis or took self-medication and recovered.



Among the 26 donors reactive for anti-dengue IgG antibodies (Table 3), 18 donors revealed history of fever, 16

donors revealed history of myalgia and 2 revealed history of vomiting and none of them reported history of rashes and headache. Of the donors who were negative for IgG dengue n = 582, 172 donors revealed history of fever, 140 donors revealed history of myalgia, 8 donors revealed history of vomiting, 28 donor revealed history of headache.

Symptoms - Groups	IgG Positive	%	IgG Negativ e	%	Total	%	P Value Fishers Exact Test
Fever	18	69.23	172	29.55	190	31.25	0.003
Myalgia	16	61.54	140	24.05	156	25.66	0.006
Vomiting	2	3,84	8	1.3	20	3.29	0.062
Headache	0	0.00	28	4.81	28	4.61	> 0.999
Table 3. Donor Population with History of Symptoms of Viral Illness in the Preceding Year							

The data in Table 3, subjected to Fishers Exact test reveals the existence of statistically significant association between fever, and myalgia status and IgG study groups (p < 0.05). Similarly when data was subjected to fishers exact test it reveals the existence of statistically non-significant association between vomiting and headache status and IgG study groups (p > 0.05)

Logistic regression model for statistically significant independent predictors of having IgG antibodies against dengue virus risk factors when performed, found no significant association between the age, gender, area of living and symptoms suggestive of dengue in past one year.

DISCUSSION

In a study from New Delhi, in the northern part of the country, among a population of 200 donors tested, 58 % of the donors had a reactive test for IgG antibodies, when these donors were tested for the presence of virus using RT-PCR, none of the donors had detectable viremia.⁶

A study from northern India in 2013, where 1709 whole blood donations were tested for presence of NS1 antigen, the prevalence of NS1 was found to be zero. In this study donors were representative of the patient population suffering from dengue.⁷ As this study was aimed at detecting NS1 that is present in the acute phase of the dengue illness, it is only logical that anyone with fever or with the symptoms of viral illness would not have turned up for blood donation.

In a study conducted across six distinct geographic location across India during the period 2011 - 2012, samples from 2609 healthy children aged 5 - 10 years were tested for dengue IgG antibodies. A high prevalence rate of 56 % was recorded in this study. This study revealed that the disease burden of dengue in India is comparable to disease burden of Southeast Asian and Latin American countires.¹² Similarly another study from Andhra Pradesh revealed a high prevalence (89.5 %) of dengue IgG antibodies among apparently healthy people who visited hospital for Master Health Check-up.¹³ A prevalence study done in 2017 post dengue surge in 2016 in Pune among 1434 individuals covering men, women and children revealed a prevalence of 81 % for dengue specific IgG antibodies.¹⁴

A study from Chennai with a sample size of 1010 general population the prevalence of IgG specific dengue antibodies was found to be 93 % indicating that 93 % of the sample

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population had a previous dengue infection whereas only 1 % revealed a history of dengue infection indicating a high percentage subclinical infection.¹⁵ In another study from the country's capital New Delhi, 2125 individual households who were living around 50 confirmed dengue cases were tested for dengue specific antibodies which revealed a prevalence of 36 % for dengue specific IgG antibodies.¹⁶

On comparison with other studies, ¹²⁻¹⁶ the prevalence of IgG antibodies in the blood donors is as low as 4.14 %. In this present study. In the other studies mentioned above the sample population included in the study were general population without apparent disease, whereas in our study the sample population included are healthy donors in whom history of dengue and other history of symptoms consistent with viral illness has been elicited. Donors who gave history of dengue in past and donors who revealed symptoms consistent with dengue were excluded from this study. Such an exclusion criteria of this study would reveal the true subclinical infection burden of dengue among blood donor population.

In this study, the prevalence of dengue IgG antibodies is low, (4.14 %) as compared to a similar study from north India where a high prevalence of 58 % has been recorded. A study on the prevalence of asymptomatic dengue infection among blood donors from Australia, Hondorus and Brazil revealed a prevalence of 0.3 % in Honduras to 0.04 in Brazil.¹⁷ A study from Taiwan in 2015 tested eight thousand samples for dengue viremia NS1 and anti-dengue IgM and IgG antibodies revealed very low prevalence rate of 0 %, 0.21 % and 0.16 % respectively.¹⁸ A study from Brazil on transfusion transmitted dengue in year 2012 revealed a dengue viremia prevalence of less than 1 % among the blood donors.¹⁹

The prevalence of viremia as demonstrated by detecting dengue viral RNA has revealed prevalence of less than 1 % among blood donors.^[17-19] The period of viremia had been observed until day 8 on average for all the four serotypes. The period of NS1 antigenemia, on average lasts until day 9 in most of the reported cases.²⁰ There are only a few cases reported for transfusion transmissible dengue³ as the period of viremia and antigenemia resolve after the period of acute illness.

Studies Conducted in Malaysia²¹ and Singapore²² during the period 2009 - 2010, both places were endemic for dengue revealed a high prevalence of dengue IgG antibodies of 46 % and 50 % respectively. When compared to similar studies on blood donors^{5,6} the prevalence of anti-dengue IgG were less in this study as donors with previous dengue infection, donors who revealed history of symptoms consistent with dengue were excluded from the study. Such an exclusion would reveal the true subclinical infection prevalence among blood donors.

Limitation

IgG positive donors were not evaluated for the presence of antigen or for presence of the virus and the infectivity of donor blood unit at this point cannot be commented upon.

CONCLUSIONS

Prevalence of dengue infection is increasing in India with the prevalence increasing with age. Social, climatic and economic conditions combined with lack of vaccine are factors largely favouring the spread of the disease. Transfusion transmitted dengue though reported in clusters in few parts of the world, its incidence is very less. Prevalence of subclinical dengue infection is less when compared to similar studies from other parts of the country. When prospective blood donors are examined for history of dengue infection and for symptoms consistent with dengue, deferring such donors for the recommended deferral period of 6 months would largely prevent transfusion transmissible dengue.

Data sharing statement provided by the authors is available with the full text of this article at jemds.com.

Financial or other competing interests: None.

Disclosure forms provided by the authors are available with the full text of this article at jemds.com.

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