Original Research Article

Seroprevalence of dengue in the tribal district of central India

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A R T I C L E   I N F O

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A B S T R A C T

Background: Dengue is a vector-borne disease that is a major public health threat globally affecting three billion people with approximately 200 million cases of morbidity and 50,000 cases of mortality annually. It is transmitted to humans by Aedes mosquitoes. It is caused by the dengue virus (DENV, 1–4 serotypes). The clinical manifestations of dengue vary from mild fever to severe hemorrhage, shock, and death. It is critical to make an early and accurate laboratory diagnosis of DENV infection for effective patient management. Dengue is now no more restricted to the urban population, it has become endemic in the rural population as well.

Aim and Objectives: 1. To find the seroprevalence of dengue virus infection in the tribal population of Gondia district. 2. To observe the seasonal variation of dengue cases.

Materials and Methods: This study was conducted under the NVBDC program at the Department of Microbiology, GMC, Gondia which is the sentinel center for Dengue and Chikungunya. The Study was carried out from October 2018 to September 2020. Serum samples of patients with dengue-like clinical illness were subjected to IgM antibody detection by dengue MAC ELISA.

Results: Seroprevalence of dengue in Gondia was found to be 12.37% (48/388). The most common age group affected was 21-30 years (39.59%). Males (54.17%) were affected more than females (45.83%). The peak of dengue cases was observed in September 2019. Less number of cases were reported in the year 2020.

Conclusion: Newer diagnostic techniques, public awareness programs, better education, and proper monitoring of vector control are required to prevent dengue outbreaks.

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1. Introduction

Dengue, also known as "tropical flu," is a viral disease transmitted to humans by Aedes mosquitoes.¹ Dengue virus is a positive-stranded RNA virus in the genus Flavivirus, family Flaviviridae.² There are four antigenically related but distinct serotypes of the dengue virus: DENV-1, DENV-2, DENV-3, and DENV-4.³

Dengue was reclassified by WHO IN 2009 according to levels of severity as dengue without warning signs, dengue with warning signs like abdominal pain, persistent vomiting, fluid accumulation, mucosal bleeding, lethargy, liver enlargement, increasing hematocrit with decreasing platelets, and severe dengue which included dengue with severe plasma leakage, severe bleeding, or organ failure.⁴ There is a need for early diagnosis of dengue infection since early diagnosis can prevent fatal cases. The most widely used serological method for dengue diagnosis is ELISA.³

Dengue virus (DENV) infection has created a serious global public health challenge to three billion people, resulting in approximately 200 million cases of morbidity and 50,000 cases of mortality annually. In 2019, the Ministry of Health and Family welfare reported 136422 dengue cases with 132 deaths all over India. A total number

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45
of 12374 dengue cases with 25 deaths were reported in Maharashtra.\(^5\)

Increasing incidence of the disease has been attributed to improper water storage, rampant unplanned urbanization, overcrowding, inadequate sanitation, and sewer system leading to the proliferation of vector breeding sites in urban, semi-urban, and rural areas.\(^6\) So, we conducted this study to know the seroprevalence of Dengue in Gondia, a tribal district of central India.

2. Materials and Methods

This cross-sectional study was conducted under National Vector Borne Disease Control Programme (NVBDCP) at the Department of Microbiology, GMC, Gondia, which is the sentinel center for Dengue and Chikungunya. The study was carried out from October 2018 to September 2020. Serum samples of patients with dengue-like clinical illness were subjected to IgM antibody detection by dengue MAC ELISA.

Blood samples were collected and serum was separated as per the standard guidelines.\(^7,8\) Samples were processed by NIV Dengue IgM Capture ELISA KIT. Serum was diluted 1:100 in tubes or deep well plate using sample diluents for DEN IgM. IgM coated strips were washed three times with 1X wash buffer. 50 \(\mu l\) of diluted samples were transferred from the deep well plate to respective wells as per the protocol on the ELISA sheet using a multichannel pipette. 50 \(\mu l\) of DEN IgM Positive and Negative controls were added to respective wells as per the protocol. The plate was covered with aluminum foil to prevent the evaporation of samples. The plate was kept in a closed humidified box inside the incubator at 37\(^0\)C for 1 hour. After washing five times with wash buffer, the plate was tapped after the last wash on a tissue paper to remove traces of wash buffer content. 50 \(\mu l\) of DEN antigen was added to each well of the plate. 50 \(\mu l\) of Anti DEN Monoclonal antibody HxB (biotin-labeled) was added to each well.

In each well, a 50 \(\mu l\) of Avidin-HRP was added. The plate was kept in a closed humidified box inside the incubator at 37\(^0\)C for 1 hour.100 \(\mu l\) of Liquid TMB substrate (TMB/H2O2) was added to each well. The plate was incubated at room temperature in dark for 10 minutes. The reaction was stopped exactly after 10 minutes with a 100 \(\mu l\) stop solution. Absorbance was measured at 450 nm within 10 minutes after the termination of the reaction. If the OD value of the sample tested was less than the OD value of negative control by factor 2.0, the sample was considered as negative. If the OD value of the sample tested exceeded the OD of negative control by factor 3.0, the sample was considered as positive. If the OD value for the sample tested exceeded OD of negative control by a factor of 2.0 but was less than OD of negative control by a factor of 3.0, the sample was considered as equivocal.

Data was analysed using statistical package for social sciences version 16 (SPSS V16). Significance value of less than 0.05 (\(p<0.05\)) was considered for statistical tests.

3. Results

A total number of 388 samples were tested, out of which 48 were positive. So, the seroprevalence of dengue in Gondia was found to be 12.37% as shown in Figure 1.

![Fig. 1: Seroprevalence of dengue in Gondia](image)

Table 1 shows that Males (54.17%) were affected more than females (45.83%). The difference was not statistically significant.

![Fig. 2: Age wise distribution of positive cases](image)

Figure 2 shows that maximum positive cases were in the age group of 21-30 years (39.59%). Figure 3 shows the peak of Dengue cases was observed in September 2019.

Table 2 shows year-wise distribution of dengue cases. Less number of cases were reported in 2020.

4. Discussion

At present, information about dengue infection is quite limited, and officially reported cases are inadequate. With an increase in morbidity, mortality rates, and economic burden, understanding of dengue infection is essential to assist policymakers and public health managers to prepare
Table 1: Gender wise distribution of positive cases

<table>
<thead>
<tr>
<th>Age in years</th>
<th>Male</th>
<th>Male (%)</th>
<th>Female</th>
<th>Female (%)</th>
<th>Total</th>
<th>Total (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>0-10</td>
<td>1 (2.08)</td>
<td></td>
<td>1 (2.08)</td>
<td></td>
<td>02 (4.16)</td>
<td></td>
</tr>
<tr>
<td>11-20</td>
<td>7 (14.58)</td>
<td></td>
<td>6 (12.5)</td>
<td></td>
<td>13 (27.08)</td>
<td></td>
</tr>
<tr>
<td>21-30</td>
<td>10 (20.84)</td>
<td></td>
<td>9 (18.75)</td>
<td></td>
<td>19 (39.59)</td>
<td></td>
</tr>
<tr>
<td>31-40</td>
<td>4 (8.33)</td>
<td></td>
<td>3 (6.25)</td>
<td></td>
<td>07 (14.58)</td>
<td></td>
</tr>
<tr>
<td>41-50</td>
<td>2 (4.17)</td>
<td></td>
<td>2 (4.17)</td>
<td></td>
<td>04 (8.34)</td>
<td></td>
</tr>
<tr>
<td>&gt;50</td>
<td>2 (4.17)</td>
<td></td>
<td>1 (2.08)</td>
<td></td>
<td>03 (6.25)</td>
<td></td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>26 (54.17%)</strong></td>
<td></td>
<td><strong>22 (45.83%)</strong></td>
<td></td>
<td><strong>48 (100%)</strong></td>
<td></td>
</tr>
</tbody>
</table>

Table 2: Year-wise distribution of dengue cases

<table>
<thead>
<tr>
<th>Year</th>
<th>No. of samples tested</th>
<th>Year</th>
<th>No. of positive cases</th>
</tr>
</thead>
<tbody>
<tr>
<td>2018</td>
<td>74</td>
<td>2019</td>
<td>227</td>
</tr>
<tr>
<td></td>
<td>9 (2.32%)</td>
<td></td>
<td>35 (9.02%)</td>
</tr>
<tr>
<td>2019</td>
<td>227</td>
<td>2020</td>
<td>87</td>
</tr>
<tr>
<td></td>
<td>4 (1.03%)</td>
<td></td>
<td>48 (12.37%)</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>388</strong></td>
<td><strong>No. of positive cases</strong></td>
<td><strong>9 (2.32%)</strong></td>
</tr>
</tbody>
</table>

Fig. 3: Month-wise distribution of Dengue cases

for and control outbreaks with limited resources for both diagnosis and treatment.

Dengue was earlier considered as an urban and semi-urban disease but in recent years, dengue has become endemic in rural areas of India as well due to water storage practices and large-scale development activities in rural areas. This has increased the scale of the dengue challenge in the country. Thus, the present study was carried out to study the seropositivity and the seasonal variations of dengue infection in our region.

4.1. Seroprevalence of dengue infection

A total number of 388 samples were tested, out of which 48 were positive. So the Seroprevalence of dengue in our study was found to be 12.37%. Seroprevalence in Study by Rao MS et al. from Andra Pradesh, South India was 17.7% and Sood S et al. from Rajasthan was 18.99%. A higher seroprevalence of 31.3% was reported in central India by Ukey PM et al. and 24.49% by Deshkar ST et al.  

4.2. Age-wise distribution

In Maharashtra, dengue is the second most leading cause of death after diarrhea in children below five years of age. In our study, 21-30 (39.59%) years were the most common age group involved. Kumar A et al. observed that a maximum number of cases, 57.3%, were in the age group of 15-44 years. In our study, 4.16% cases were in the age group of 0-10 years, whereas, Ukey PM et al., Deshkar ST et al., Rao MS et al., observed maximum cases in the age group of 0-10 years which was 43.90%, 40.50%, 35.84% respectively. In general, the prevalence of dengue infection is relatively seen more in children compared to adults but persons of all age groups are susceptible to dengue infections.

4.3. Gender-wise distribution

Our study showed Males (54.17%) were affected more than females (45.83%) which coincides with a study by, Karoli R et al. in which 58% male patients and 42% females. In a study by Kumar M., males being affected significantly more compared to females. Similarly, many other Indian studies were independently done by Agarwal et al., Ray et al., and Wali et al., Deshkar ST et al. found a higher number of male patients infected with dengue compared to females. It might be due to differences in the socio-cultural environment where males are more exposed to outdoor activities and their bodies less covered as compared to females.

4.4. Month-wise distribution

In our study, an increase in the number of cases was observed in October and November 2018. In 2019 also, there was a gradual increase in the number of cases from July with a peak in September. Kumar A et al. also observed a gradual increase in cases from June with a peak...
in September. Peak was observed in August, September, October, and November in a study by Deshkar ST et al.\textsuperscript{2} Gunasekaran P et al. also reported a high percentage of IgM positivity during September and October in all the three years.\textsuperscript{21} Kumar M et al. also observed a marked rise in the number of dengue positive cases between August and December.\textsuperscript{15}

Less number of cases were reported in the year 2020. This reduction could be attributable to the mobilization of epidemiological surveillance teams to respond to the emergence of the COVID-19 pandemic, causing a delay or underreporting of cases of dengue. Furthermore, most of the febrile dengue cases overlap in health centers along with cases of COVID-19, sharing clinical features.\textsuperscript{22} We are looking forward to increasing the serosurveillance of dengue cases in upcoming years.

5. Conclusion

Despite several preventive measures by WHO, new outbreaks of dengue infection have been reported in several parts of the world during the post-monsoon season. To prevent such outbreaks, newer diagnostic techniques, public awareness programs, better education, and proper monitoring of vector control are required.

6. Source of Funding

None.

7. Conflict of Interest

The authors declare that there is no conflict of interest.

8. Acknowledgement

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References


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