

Seroprevalence of Dengue Fever Virus among Suspected Patients in Taiz Governorate-Yemen

Abstract

Background: Dengue Fever virus (DENV) considers one of the most important mosquito-borne viral disease in the world and it is endemic in more than 100 countries.

Objective: This study aimed to determine the seroprevalence of DENV infection among suspected patients and to investigate some associated risk factors with dengue fever infection in Taiz governorate, Yemen.

Methods: This study was cross-sectional, descriptive, and experimental, combining the use of a structured questionnaire and analysis of serum samples obtained from 300 suspected patients attending at many hospital and clinic centers in Taiz during the period from July to November 2016. The serum samples were tested for anti-dengue immunoglobulin (IgM) and (IgG) by Enzyme-linked Immunosorbent Assay (ELISA).

Results: Out of 300 suspected febrile cases, it was found that 49 (16.3%), 68 (22.7%), and 17 (5.7%) cases were showed positive for the IgM, IgG, and both IgM and IgG antibodies, respectively, while 166 (55.3%) cases were negative. The incidence rate was more in males than in females. The most affected age group with dengue fever infection were (21–30) years. Dengue Fever was more frequent among patients coming from the urban area, having secondary school, and low-income status people. Also, there was statistical significant between DENV infections with a place of residency and gender ($P < 0.05$) and not-statistical significance between DENV infections and other factors ($P > 0.05$). **In conclusion,** Taiz governorate become one of the endemic governorates in Yemen particularly the Taiz city which should be brought to the attention of public health authorities

Keywords: Dengue fever virus, IgG, IgM, Seroprevalence, Taiz governorate, Yemen.

INTRODUCTION

Dengue Fever virus (DENV) is one of the most important mosquito-borne viral disease in the world and it is endemic in more than 100 countries, with an estimated 100 million infected cases and about 25,000 deaths per year worldwide¹. DENV is an enveloped, positive-sense, single-stranded RNA genome that belongs to the family Flaviviridae, genus Flavivirus².

Dengue fever is caused by one of the four serotypes of dengue viruses (DENV1-4). These types of viruses are transmitted via infective female mosquitoes, namely *Aedes aegypti* (principal vector) and *Aedes albopictus*, through the bites or blood meals on human hosts³. Direct person-to-person transmission has not been documented. Although a few case reports have been published on the transmission of DENV through exposure to DENV-infected blood, organs, or other tissues from blood transfusions, solid organ or bone marrow transplants^{4,5}.

Dengue virus causes primary and secondary infections. Primary infection is an acute feverish illness known as Dengue Fever (DF) which mostly eliminate around seven days by a composite immune response, while the secondary infection is additional rigorous and causes Dengue Hemorrhagic Fever (DHF) or Dengue Shock Syndrome (DSS)^{6,7}. The

majority of deaths that result from dengue infection result from Dengue Hemorrhagic Fever (DHF) and Dengue Shock Syndrome (DSS) ⁸.

In Yemen, the early first recorded of dengue-like epidemics in Aden was reported by Hirsch between 1870 and 1873 years ⁹. Also, the prevalence of dengue fever was recorded in Al-Hudidah governorate in 1954 which affected 98% of the studied population ¹⁰. After that dengue fever became an epidemic in the coastal planes of the Tehama (Hudidah), then outbreaks increased since 2005 and the disease has spread to new governorates ¹¹.

All three dengue viral serotypes 1-3 were confirmed to be in Yemen by using PCR, but Dengue virus type 2 was the most predominant serotype in Yemen ¹². Interestingly DENV serotype 4 was identified and confirmed the first time in Al-Hudidah-Yemen by Alahdalet *al.* ¹³.

In the Taiz governorate, the infection rate varied from year to year. Recently, WHO ¹⁴ documented an extreme prevalence in cases of dengue fever in Taiz governorate were reached to 1328 suspected cases compared with last years. But seroprevalence of DENV is not well documented so far. Therefore, this study designed to determine the seroprevalence of dengue fever and to reveal some possible risk factors associated with DENV infection in Taiz governorate, Yemen.

MATERIALS AND METHODS

Study area

Taiz governorate is situated in the southwestern part of the Republic of Yemen between latitudes (12-14°) North of the equator and between longitudes (43-45°) East of GMT. Taiz is the third largest governorate in Yemen, it holds the first rank in terms of the population which rate 2,393,425 according to the census results for the year (2004) ¹⁵. Administratively, Taiz is among the largest governorates as it includes 23 districts in its frame. The climate of Taiz is characterized by diversity, where their highland parts are cold in winter and mild in summer, while low parts are mild in winter and warm to hot relatively in summer. The climate in the western parts of the governorate is the coastal desert climate that is warm in winter and hot in summer, and the average temperature in the governorate is up (21°C). The rain falls in the summer on all parts of the governorate and in some districts in the winter ¹⁶.

Study design

This is a cross-sectional, descriptive study was conducted from July to November 2016, performed on suspected patients suffering undifferentiated fever (age range, 1 to 65 years) attending many hospitals and health centers at nine main locations in the Taiz governorate, which most patients coming seeking medical care.

Ethical approval

Approval for this study was obtained from the Ethical Review Committee in the Biology Department, Faculty of Science, Sana'a University.

Data collection

A structured questionnaire was designed to collect required data from suspected patients regarding socio-demographics and risk-related data. The questionnaire was filled for each participant via face-to-face interview by a researcher to avoid any misunderstanding and confirm the accurate collection of data properly.

Sample collection

Five-mL blood sample was collected from 300 suspected patients suffering undifferentiated fever by venipuncture, transferred into a sterile anticoagulant-free sterile bottle, and allowed to clot. The clotted blood sample was centrifuged (3000 rpm, 5 min), and the serum (the supernatant) was put in separate Eppendorf tubes with a specific study number (SNO) transferred inside a cooling box and stored at -20°C until required for use.



Figure 1: Map of Yemen and Taiz governorate

Serological Assay

Samples of serum were tested for DENV-specific IgG and IgM antibodies by using IgG and IgM DRG Immunodiagnostic Kits (GmbH Germany) that performed by Enzyme-Linked Immunosorbent Assay (ELISA) (Absorbance Microplate Reader/ELIZA-IRE96, SFRI, French)¹⁷.

Statistical analysis

The obtained data were analyzed using version 18.0 SPSS (Statistical Package for Social Science). A significant difference between the proportions and the groups or variables was

determined by Chi-square test (≥ 3.9 considered significant) and *P*-value (< 0.05 considered significant).

RESULTS

In a total of 300 participants enrolled in this study, 202 (67.3%) were males and 98 (32.7%) were females. The highest participant groups in the present study were patients aged between (21-30) years with an average of 105 (35%). Most suspected patients 195 (65%) were living in urban areas. The suspected patients with secondary education levels were more than the third 113 (37.7%) of the total population in this study. On the other hand, more than half of the respondents 160 (53.3%) were of low-income status (Table 1).

Table 1: Characteristics of study cases

| Variable | | Number (%) | Variable | | Number (%) |
|-----------|---------|------------|-----------------|-------------|------------|
| Gender | Male | 202 (67.3) | Education level | Illiterate | 46 (15.3) |
| | Female | 98 (32.7) | | Primary | 56 (18.7) |
| Age | 1 – 10 | 24 (8) | | Secondary | 113 (37.7) |
| | 11 – 20 | 58 (19.3) | | University | 85 (28.3) |
| | 21 – 30 | 105 (35) | Income Status | Low Status | 160 (53.3) |
| | 31 – 40 | 64 (21.3) | | Med Status | 118 (39.3) |
| | 41 – 50 | 29 (9.7) | | High Status | 22 (7.3) |
| | > 50 | 20 (6.7) | | | |
| Residence | Rural | 105 (35) | | | |
| | Urban | 195 (65) | | | |

Out of 300 suspected febrile cases, 49 (16.3%) showed positive results for the IgM antibodies (acute infection) and 68 cases (22.7%) were positive for the IgG antibodies (chronic infection), and 17 cases (5.7%) were positive for both IgM and IgG (acute and chronic infection), while 166 (55.3%) cases were negative for anti DENV antibodies (Figure 2).

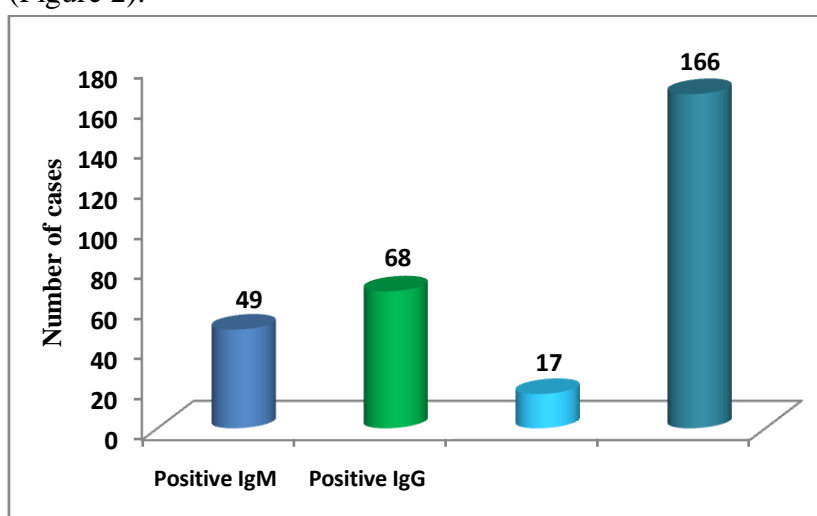


Figure 1: The positive and negative of anti DENV antibodies

In the present results, the DENV IgM, IgG, and both IgM and IgG seropositivity were mostly observed in male patients, 39 (19.3%), 53 (26.2%), and 14 (6.9%), respectively, while the female patients were lower from that 10 (10.2%), 15 (15.3%), and 3 (3.1%),

respectively. Also, the statistical analysis showed a significant association between the gender and seroprevalence of DENV IgM and DENV IgG ($P < 0.05$) (Table 2).

In the current work, the higher prevalence rate of DENV IgM, IgG, and both (IgM and IgG) seropositivity were observed among patients aged from 21-30 years with 22 (21%), 29 (27.6%), and 8 (7.6%), respectively. While the lowest from those was among (> 50) years, 2 (10%), 3 (15%) and 0 (0 %), respectively. There was an association between age groups and seroprevalence of DENV was not significant ($P > 0.05$) (Table 2).

The result regarding the resident area, it was found that 19.5% of positive anti-IgM, 26.2% of positive anti-IgG, and 8.2% of positive anti-IgM and anti-IgG antibodies were reported among patients coming from the urban area. On other hand, the seropositivity of DENV antibodies among cases living in the rural area was 10.5% for IgM, 16.2% for IgG, and 1% for both IgM and IgG. Also, there was a statistically significant association between the place of residence and seroprevalence of DENV antibodies ($P < 0.05$) (Table 2).

DENV IgM, IgG, and both (IgM and IgG) seropositive were the highest rates among participants who have secondary education level with rate 22 (19.5%), 31 (27.4%), and 7 (6.2%), respectively, whereas the lowest rate was found among illiterate participants with 5 (10.9%), 7 (15.2%) and 2 (4.3%), respectively. The result showed no significant association between any education levels and seroprevalence of DENV ($P > 0.05$) (Table 2).

Table 2 shows the highest rate of DENV seropositivity of IgM, IgG, and both IgM and IgG antibodies were recorded among cases with low-income status when compared to the lowest seropositivity of DENV antibodies among cases with high-income status.

Table 2: Distributions of anti-DENV with socio-demographical characteristic of the suspected patients

| Variables | | Number of cases | IgM positive | | IgG positive | | IgM and IgG positive | |
|-----------------|------------|-----------------|--------------|---------|--------------|---------|----------------------|---------|
| | | No. (%) | No. (%) | P-value | No. (%) | P-value | No. (%) | P-value |
| Gender | Male | 202 (67.3) | 39 (19.3) | 0.04 | 53 (26.2) | 0.03 | 14 (6.9) | 0.2 |
| | Female | 98 (32.7) | 10 (10.2) | | 15 (15.3) | | 3 (3.1) | |
| Age (years) | 1 – 10 | 24 (8) | 3 (12.5) | 0.7 | 4 (16.7) | 0.7 | 1 (4.2) | 0.8 |
| | 11 – 20 | 58 (19.3) | 9 (15.5) | | 12 (20.7) | | 3 (5.2) | |
| | 21 – 30 | 105 (35) | 22 (21) | | 29 (27.6) | | 8 (7.6) | |
| | 31 – 40 | 64 (21.3) | 9 (14.1) | | 14 (21.9) | | 4 (6.3) | |
| | 41 – 50 | 29 (9.7) | 4 (13.8) | | 6 (20.7) | | 1 (3.4) | |
| | >50 | 20 (6.7) | 2 (10) | | 3 (15) | | 0 (0) | |
| Residence | Rural | 105 (35) | 11 (10.5) | 0.04 | 17 (16.2) | 0.04 | 1 (1) | 0.01 |
| | Urban | 195 (65) | 38 (19.5) | | 51 (26.2) | | 16 (8.2) | |
| Education level | Illiterate | 46 (15.3) | 5 (10.9) | 0.6 | 7 (15.2) | 0.6 | 2 (4.3) | 0.9 |
| | Primary | 56 (18.7) | 8 (14.3) | | 11 | | 3 (4.5) | |

| | | | | | | | | |
|----------------------|-------------------|------------|-----------|------------|-----------|------------|----------|------------|
| | | | | | (19.6) | | | |
| | Secondary | 113 (37.7) | 22(19.5) | | 31(27.4) | | 7(6.2) | |
| | University | 85(28.3) | 14(16.5) | | 19(22.4) | | 5(4.8) | |
| Income Status | Low | 160 (53.3) | 30 (18.8) | 0.4 | 41 (25.6) | 0.2 | 10 (6.3) | 0.4 |
| | Med | 118 (39.3) | 17 (14.4) | | 25 (21.2) | | 7 (5.9) | |
| | High | 22 (7.3) | 2 (9.1) | | 2 (9.1) | | 0 (0) | |

χ^2 Chi-square ≥ 3.8 ; $P < 0.05$ (significant)

DENV IgM, IgG, and both (IgM and IgG) seropositivity case the most effective breeding sites of mosquitoes were the bogs that were 31 (19.6%), 42 (26.6%), and 11 (6.9%) of infected cases. The second source was trashes 36 (18.2%), 46 (23.4%) and 12 (6.1%), then open drums 24 (17.1%), 31 (22.1%) and 10 (7.1%), followed by, open sewage 10 (16.7%), 13 (21.7%) and 2 (3.3%). Next, pools 6 (13.6%), 9 (20.5%) and 4 (9.1%) respectively. The lower abundant site was tires which recorded 7 (12.5%), 10 (17.9%) and 3 (5.4%), respectively of the breeding sites. There was no significant association between the seroprevalence of DENV and all breeding sites of mosquitoes ($P > 0.05$) (Table 3).

Table 3: Seroprevalence of DENV antibodies according to breeding sites of mosquitoes.

| Breeding sites | Examined Cases | IgM positive | | | IgG positive | | | IgM and IgG positive | | |
|--------------------|----------------|--------------|----------|---------|--------------|----------|---------|----------------------|----------|---------|
| | No. (%) | No. (%) | χ^2 | P-value | No. (%) | χ^2 | P-value | No. (%) | χ^2 | P-value |
| Bogs | 159 (53) | 31 (19.6) | 2.4 | 0.1 | 42 (26.6) | 2.7 | 0.1 | 11 (6.9) | 0.9 | 0.3 |
| Open Sewage | 60 (20) | 10 (16.7) | 0.0 | 0.9 | 13 (21.7) | 0.0 | 0.8 | 2 (3.3) | 0.7 | 0.3 |
| Pools | 44 (14.7) | 6 (13.6) | 0.3 | 0.6 | 9 (20.5) | 0.1 | 0.7 | 4 (9.1) | 1.2 | 0.3 |
| Open drums | 140 (46.7) | 24 (17.1) | 0.1 | 0.7 | 31 (22.1) | 0.0 | 0.8 | 10 (7.1) | 1.1 | 0.3 |
| Trash | 197 (65.7) | 36 (18.2) | 1.5 | 0.2 | 46 (23.4) | 0.1 | 0.7 | 12 (6.1) | 0.2 | 0.7 |
| Tires | 56 (18.7) | 7 (12.5) | 0.7 | 0.4 | 10 (17.9) | 0.9 | 0.3 | 3 (5.4) | 0.01 | 0.9 |

χ^2 Chi-square ≥ 3.8 ; $P < 0.05$ (significant)

In the present finding, the most of infected cases that observed positive for IgM, IgG, and both (IgM and IgG) antibodies were recorded in Taiz city with 31 (19.4%), 43 (26.9%) and 10 (6.9%), followed by Al-Barh 8 (17.7%), 11 (24.4%) and 3 (6.7%), then Al-Rahedah 4 (16.7%), 5 (20.8%) and 2 (8.3%), after that, Al-Makha 2 (12.5%), 3 (18.8%) and 1 (6.2%). Next, Al-Demna 2 (10%), 4 (20%) and 1 (5%), respectively, the lowest of that were Hajdah 1 (10%) for IgM, 2 (20%) for IgG and Mawyah 1 (10%) for IgM. While two locations were free of infection namely Al-Turbah and Al-Nashamah which no recorded any cases for IgM or IgG seropositivity (Table 4).

Table 4: The distribution of DENV antibodies according to sampling location

| Area | Participants No. (%) | IgM positive No. (%) | IgG positive No. (%) | IgGandIgM positive No. (%) |
|--------------|-------------------------|-------------------------|-------------------------|----------------------------------|
| Taiz city | 160 (53.4) | 31 (19.4) | 43 (26.9) | 10 (6.9) |
| Al-Barh | 45 (15) | 8 (17.7) | 11 (24.4) | 3 (6.7) |
| Al-Rahedah | 24 (8) | 4 (16.7) | 5 (20.8) | 2 (8.3) |
| Al-Makha | 16 (5.3) | 2 (12.5) | 3 (18.8) | 1 (6.2) |
| Al-Demna | 20 (6.7) | 2 (10) | 4 (20) | 1 (5) |
| Hajdah | 10 (3.3) | 1 (10) | 2 (20) | 0 (0) |
| Mawyah | 10 (3.3) | 1 (10) | 0 (0) | 0 (0) |
| Al-Turbah | 10 (3.3) | 0 (0) | 0 (0) | 0 (0) |
| Al-Nashamah | 5 (1.7) | 0 (0) | 0 (0) | 0 (0) |
| Total | 300 (100) | 49 (16.3) | 68 (22.7) | 17 (5.7) |

DISCUSSION

The findings of this study revealed that the overall seroprevalence of DENV antibodies among suspected patients was 134 cases (44.7%). 49 cases (16.3%) were positive for acute DENV infection (IgM positive) and 68 (22.7%) of suspected cases were found positive for chronic DENV infection (IgG positive), The cases that were positive for both acute and chronic infection (IgM and IgG positive) were 17 (5.7%), while 166 (55.3%) of suspected cases were negative for dengue fever virus.

Interestingly, in this study, the level of DENV chronic infection (IgG seropositivity) was found (22.7%) more than acute infection (16.3%) and both (5.7%). These results were similar to a study in the Shabwah governorate by Al-Moyedet *al.*¹⁸ who showed that 438 (53.5%) of cases were positive for DENV antibodies. Similarly, Abdullah *et al.*¹⁹ found that 179 cases were positive for IgM antibody (42% of all suspected specimens) and 262 cases were positive for IgG antibody (61.6%). Also, 96 specimens were positive for both (22.5%). However, 83 (19.5%) of suspected cases were negative for DENV antibodies.

The high prevalence of dengue IgG among suspected cases in this study may be attributed to previous exposures and endemicity of infection, especially in Taiz city, which may be increasing the risk of complicated dengue infection through a phenomenon known as antibody-dependent enhancement depending on the number of DENV serotypes that circulating in the governorate. This observation is an agreement with a study by Madani *et al.*²⁰.

However, the war from 2015 until now in Yemen particularly in Taiz that lead to limited primary health care services, lack of water supply systems, sanitation services, and insufficient control on mosquitos breeding sites facilitating the spread of endemic diseases such as dengue fever.

Furthermore, the distribution of DENV IgM, IgG, and both (IgM and IgG) seropositivity in Taiz mostly observed in male cases, (19.3%), (26.2%) and (6.9%), respectively. While the female cases were lower from that (10.2%), (15.3%) and (3.1%), respectively. This result was supported with similar previous reports in different countries that noticed that the males were more affected by fever infection than females by Bin Gluth *et al.*²¹, Madani *et al.*²⁰, and Abdullah *et al.*¹⁹ in Yemen; Ayyub *et al.*²² in Saudi Arabia, Abdelhalim *et al.*²³ in Sudan.

The possible reason for the high number of dengue infected cases among males due to the habit of males in the summer season, they did not cover their body whether at home or

outside, spend more time outdoor and they have traveling history to that area where the dengue incidence is high. These habits make them more exposed to the bite of *Aedes aegypti*.

In the present study, it was revealed that the most susceptible age group for DENV infection were (21-30) years which showed a higher prevalence of dengue infection (IgM=21%, IgG=27.6% and both =7.6%). While, the less infected category with dengue infection was the older people aged (> 50) years with an average (10%), (15%) and (0%) for IgM, IgG, and both, respectively. Similar observations were also reported in Yemen by Madani *et al.*²⁰ and Qassim²⁴, in India by Akula and Kammili²⁵, and Pakistan by Muhammad *et al.*²⁶. This result suggesting that the individuals in these age groups were more actively outdoor during the day which increased their chances of exposure to the infective DENV vector bite.

According to the present study, the relationship between the dengue fever seropositivity and education levels which found higher in people with secondary education levels compared with lower among illiterates participants. This finding is similar to a study by Abdullah *et al.*¹⁹ conducted in Yemen. This finding may reflect their outdoor activity during the day, for playing, schooling, or picnic that increased their chances of exposure to the infective DENV vector bite.

The prevalence of dengue fever infection was noted significantly correlated with place of residence (rural or urban) areas. The prevalence rate of dengue fever infection in the urban area was more than the rural areas. This finding is in agreement with earlier studies in Yemen by Abdullah *et al.*¹⁹ and Bin Gluth *et al.*²¹.

In the present study, the income statuses were found non-significantly associated with dengue infection. But Dengue IgM, IgG and both seropositivity, were found highest with low-income statuses cases (4-7 \$ daily), it was rated (18.8%), (25.6%) and (6.3%), respectively. Whereas, the less infected people were with high-income statuses (>20 \$ daily), which rated (9.1%), (9.1%) and (0%) for IgM, IgG, and both, respectively. Similar results were reported by Al-Hemiree²⁷ and Abdullah *et al.*¹⁹ in Yemen and by Muhammad *et al.*²⁶ in Pakistan.

Furthermore, in this work the dengue mosquitoes breeding sites were found positive factors related to dengue IgM, IgG, and both seropositivity. The bogs were the most breeding that associated with most of the suspected cases compared to the lowest rate among the other factor such as trash, open drums, pools, open drainage and the tires. This result is an agreement with Abdullah *et al.*¹⁹ who noted that the highest breeding site factor was the bogs which rated (92.2%), while the lowest factor was the tires with an average (26.3%).

Interestingly, the distribution of dengue fever results according to districts revealed that the of the most hotspot of dengue fever virus infection in Taiz governorate was concentrated in the city of Taiz, followed by Al-Barh, Al-Rahedah, Al-Makha, Al-Demna, Hajdah, and Mawyah. Whereas two districts namely Al-Turbah and Al-Nashamah were found free from DENV infection that may be due to climate change and the lack of suitable conditions for the mosquitoes breeding cycle in these areas.

The low infected cases were found Al-Makha during this study that the infected case was low maybe contributing to the war most people replacement to another area within the governorate most infected case in Al-Barh which were coming from Alhodidah and Al-Makha.

CONCLUSION

In conclusion, the high prevalence of DENV antibodies in Taiz is becoming one of the most endemic governorates in Yemen which should be brought to the attention of public

health authorities. Warm climate, rainfall also the war since 2015 until now contributed to destroying Yemen's healthcare, presence of breeding sites, lack of water supply systems and mosquito control measure, low-income status, and insufficient sanitation systems are the reasons that attributed to the increase of suspected cases of DENV among study area. Therefore, continuous surveillance for outbreaks of DENV infection required to identify early and in order to prevent and control the spread of infections among the community.

ACKNOWLEDGMENTS

The authors would like to thank Laboratories and health care centers located at Taiz Governorate for their great help, also to all the team at Dar-Asaha Modern Medical Laboratory, Taiz, Yemen and Jordanian University, Sana'a, Yemen, for their cooperation.

REFERENCES

1. Arima Y, Chiew M, Matsui T, *et al.* Epidemiologic update on the dengue situation in the Western Pacific Region, 2012. *Western Pac Surveill Response J* 2015; 4: 47-54. DOI: [10.5365/WPSAR.2014.5.4.002](https://doi.org/10.5365/WPSAR.2014.5.4.002)
2. Lindenbach BD, Rice CM. Molecular biology of flaviviruses. *Adv Virus Res* 2003; 59: 23-61. DOI: [10.1016/s0065-3527\(03\)59002-9](https://doi.org/10.1016/s0065-3527(03)59002-9)
3. World Health Organization (WHO). Dengue: guidelines for diagnosis, treatment, prevention and control _New ed. Geneva. World Health Organization. 2009; 1-160.
4. Smith WA, Gubler J. Geographic expansion of dengue: the impact of international travel. *Med Clin North Am* 2008; 92(6):1377-1390. DOI: [10.1016/j.mcna.2008.07.002](https://doi.org/10.1016/j.mcna.2008.07.002)
5. Chen LH, Wilson, ME. Dengue and Chikungunya infections in travelers. *Curr Opin Infect Dis* 2010; 23(5): 438-444. DOI: [10.1097/QCO.0b013e32833c1d16](https://doi.org/10.1097/QCO.0b013e32833c1d16)
6. Naseem S, Farheen A, Muhammad A, Fauzia R. Dengue fever outbreak in Karachi, 2005—A clinical experience. *Infect Dis J* 2005; 14(4):115-117.
7. Almas A, Parkash O, Akhter J. Clinical factors associated with mortality in dengue infection at a tertiary care center. *Southeast Asian J Trop Med Public Health* 2010; 41(2): 333-340. PMID: 20578516
8. Rigau P, Clark G, Gubler D, Reiter P, Sanders E, Vorndam A. Dengue and dengue haemorrhagic fever. *Lancet* 1998; 352:971-977. DOI: [10.1016/s0140-6736\(97\)12483-7](https://doi.org/10.1016/s0140-6736(97)12483-7)
9. Van-Kleef E, Bambrick H. The geographic distribution of dengue fever and the potential influence of global climate change. *TropIK A.net* 2011; 1-22. DOI: [10.1289/isee.2011.00337](https://doi.org/10.1289/isee.2011.00337)
10. Jimenez-Lucho VE, Fisher EJ, Saravolatz LD. Dengue with hemorrhagic manifestations: an imported case from the Middle East. *The American Journal of Tropical Medicine And Hygiene* 1984; 33(4): 650-653. DOI: <https://doi.org/10.4269/ajtmh.1984.33.650>
11. World Health Organization (WHO). Global strategy for dengue prevention and control 2012-2020. 20 avenue Appia, 1211 Geneva 27, Switzerland. 2012.
12. Al-Garadi MA. Epidemiological review of dengue fever in Yemen. *IJAR* 2015; 7:1578-1584.
13. Alahdal M, Al-Shabi J, Ogaili M, Abdullah QY, Alghalibi S, Jumaan AO, Al-Kamarany MA. Detection of dengue fever virus serotype – 4 by using one-step real-time RT-PCR in Hodeidah, Yemen. *BMRJ*, 2016; 14(6): 1-7.
14. World Health Organization (WHO). Midterm epidemiological report electronic disease early warning and response system. 2016.

15. National Information Center (NIC). About Taiz province, 2012, [http://www.yemenic.info/english_site/./Home/Aboutyemen/governorates/taiz/About governorates](http://www.yemenic.info/english_site/./Home/Aboutyemen/governorates/taiz/About%20governorates). (Accessed 30/09/2020).
16. Central Statistical Organization (CSO). Statistical year book. Central Statistical Office. Sana'a. 2013.
17. Kuno G, Gomez I, Gubler DJ. An ELISA procedure for the diagnosis of dengue infections. *J Virol Methods* 1991; 33: 101–113.
18. Al-Moyed T, Khaled A, Ali AJ, Aisha OJ. Sero-prevalence of reported dengue fever in Shabwah governorate, Yemen. *Hadramout Journal of Medical Sciences* 2012; 1:82-87. DOI: [10.12816/0005940](https://doi.org/10.12816/0005940)
19. Abdullah QY, Ogaili M, Alahdal M, AL-Kamran AM. Dengue fever infection in Hodeidah, Yemen: Risk factors and socioeconomic indicators. *British Biomedical Bulletin*. 2015; 3(1): 058-065.
20. Madani TA, Abuelzein TE, Al-Bar, HS, *et al*. Outbreak of viral hemorrhagic fever caused by dengue virus type 3 in Al-Mukalla, Yemen. *Journal of BMC Infectious Diseases* 2013; 13:136. DOI: [10.1186/1471-2334-13-136](https://doi.org/10.1186/1471-2334-13-136)
21. Bin Ghouth AS, Amarasinghe A, Letson WG. Dengue outbreak in Hadramout, Yemen, 2010: An epidemiological perspective. *Am J Trop Med Hyg* 2012; 86(6): 1072–1076. doi: [10.4269/ajtmh.2012.11-0723](https://doi.org/10.4269/ajtmh.2012.11-0723)
22. Ayyub M, Khazindar AM, Lubbad EH, *et al*. Characteristics of dengue fever in a large public hospital, Jeddah, Saudi Arabia. *J Ayub Med Coll Abbottabad* 2006; 18(2): 9-13.
23. Abdelhalim KA, Kafi SK. Seroprevalence of West Nile fever and dengue fever viruses in Suburban areas in Khartoum State, Sudan. *American Journal of Research Communication* 2014; 2(8): 81-86.
24. Qassim M. Dengue fever outbreak investigation in Taiz governorate. First national Yemen field epidemiology training program conference, 26-27 February, 2014, Sana'a, Yemen. 2014; Pp 41.
25. Akula S, Kammili N. Serological and virological profile of dengue fever in a tertiary care hospital, southern part of Hyderabad, during 2011-12. *International Journal of Microbiology* 2015; 1-7.
26. Muhammad, Ali R, Akbar S, Khan I, Ahmad T. Outbreak of dengue in Khwazakhela district Swat during August-November 2013 *Bull. Env Pharmacol Life Sci* 2014; 3 (2): 26-28.
27. Al-Hemiree AR. Epidemiological study of dengue fever in Al-Rahedah, Taiz. MSc thesis, Department Of Medical Microbiology, Faculty Of Medicine And Health Science, Sanaa University, Yemen. 2008.