

RESEARCH ARTICLE

Dengue in Java, Indonesia: Relevance of Mosquito Indices as Risk Predictors

Siwi P. M. Wijayanti^{1,2}, Sunaryo Sunaryo³, Suprihatin Suprihatin⁴, Melanie McFarlane¹, Stephanie M. Rainey¹, Isabelle Dietrich¹, Esther Schnettler¹, Roman Biek^{1,5}, Alain Kohl^{1*}

1 MRC-University of Glasgow Centre for Virus Research, Glasgow, Scotland, United Kingdom, **2** Public Health Department, Faculty of Health Sciences, University of Jenderal Soedirman, Purwokerto, Indonesia, **3** Vector Borne Disease Research and Development Unit, Banjarnegara, Indonesia, **4** Parasitology Department, University of Gadjah Mada, Yogyakarta, Indonesia, **5** Boyd Orr Centre for Population and Ecosystem Health, Institute of Biodiversity, Animal Health and Comparative Medicine, College of Medical Veterinary and Life Sciences, University of Glasgow, Glasgow, United Kingdom

* alain.kohl@glasgow.ac.uk



 OPEN ACCESS

Citation: Wijayanti SPM, Sunaryo S, Suprihatin S, McFarlane M, Rainey SM, Dietrich I, et al. (2016) Dengue in Java, Indonesia: Relevance of Mosquito Indices as Risk Predictors. *PLoS Negl Trop Dis* 10(3): e0004500. doi:10.1371/journal.pntd.0004500

Editor: Michael J Turell, United States Army Medical Research Institute of Infectious Diseases, UNITED STATES

Received: June 3, 2015

Accepted: February 9, 2016

Published: March 11, 2016

Copyright: © 2016 Wijayanti et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the paper.

Funding: This work was funded by the Higher Education Ministry of Education, Indonesia (Directorate General of Higher Education Postgraduate Scholarship to SPMW) and the UK Medical Research Council (AK). RB is supported by the RAPIIDD programme of the Science and Technology Directorate of the Department of Homeland Security and NIH Fogarty International Center. The funders had no role in study design, data

Abstract

Background

No vaccine is currently available for dengue virus (DENV), therefore control programmes usually focus on managing mosquito vector populations. Entomological surveys provide the most common means of characterising vector populations and predicting the risk of local dengue virus transmission. Despite Indonesia being a country strongly affected by DENV, only limited information is available on the local factors affecting DENV transmission and the suitability of available survey methods for assessing risk.

Methodology/principal findings

We conducted entomological surveys in the Banyumas Regency (Central Java) where dengue cases occur on an annual basis. Four villages were sampled during the dry and rainy seasons: two villages where dengue was endemic, one where dengue cases occurred sporadically and one which was dengue-free. In addition to data for conventional larvae indices, we collected data on pupae indices, and collected adult mosquitoes for species identification in order to determine mosquito species composition and population density. Traditionally used larval indices (House indices, Container indices and Breteau indices) were found to be inadequate as indicators for DENV transmission risk. In contrast, species composition of adult mosquitoes revealed that competent vector species were dominant in dengue endemic and sporadic villages.

Conclusions/significance

Our data suggested that the utility of traditional larvae indices, which continue to be used in many dengue endemic countries, should be re-evaluated locally. The results highlight the need for validation of risk indicators and control strategies across DENV affected areas here and perhaps elsewhere in SE Asia.

collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

Author Summary

Geographically and economically, Indonesia is one of the most prominent countries in SE Asia, yet many of its endemic infectious diseases are poorly managed, controlled and understood. This includes dengue virus (DENV), which can result in serious human disease and is transmitted by mosquitoes. Dengue risk assessment is a key factor in managing the impact of infection on public health, and this often relies on assessing the presence and nature of mosquitoes through a number of indices associated with the occurrence of larvae and the location/availability of breeding containers. Here we assessed traditionally used indices in combination with other indicators including pupae indices, and the presence of adult mosquitoes in areas with different dengue status: endemic, sporadic or free. Our data suggested that traditional indices were poor indicators of reported local DENV transmission. This has important consequences for design and focus of risk management strategies and efforts to control DENV locally as well as elsewhere in the region.

Introduction

Dengue virus (DENV) is considered to be the most important arbovirus world wide, with a heavy disease burden in humans [1]. It is transmitted mainly by *Aedes aegypti* mosquitoes, but *Ae. albopictus* can also act as a vector [2–5]. Dengue is endemic in many countries around the world, especially in the tropics; moreover the number of endemic areas is increasing [6]. DENV belongs to the genus *Flavivirus* in the family *Flaviviridae* and consists of four antigenically distinct and medically relevant serotypes, with a possible fifth recently described (DENV1, 2, 3, 4 and 5) [1,6–9]. The clinical spectrum of DENV infection can vary from asymptomatic to more severe forms such as dengue hemorrhagic fever (DHF) and dengue shock syndrome (DSS) [8]. DENV is transmitted to humans following mosquito bite (horizontal transmission), and mosquitoes can become infected by ingestion of a DENV-containing blood meal [2]. However, DENV can also be maintained via vertical transmission i.e. passed into eggs and subsequently into the next generation of mosquitoes, thereby maintaining outbreaks in human populations. This has already been documented in both *Ae. aegypti* and *Ae. albopictus* in different countries including in SE Asia [10–13].

Since it was first identified in 1968 in the cities of Jakarta (capital of Indonesia) and Surabaya (East Java), dengue disease has been recognised as an important public health problem in Indonesia. Periodic outbreaks have occurred in Indonesia with an increasing number of cases and severity [14]. DENV incidence in Indonesia has been shown to peak during the rainy season (between the months of October and April) [15]. From 2004 onwards, Indonesia reported the highest number of DENV cases in the region. All four serotypes of DENV have been found to be circulating since and DENV3 infections associated with the most severe disease [16,17]. Despite dengue being a major concern remarkably little is known or done to control this virus in Indonesia, in spite of its size (in surface and population, as the world's largest island nation but with high levels of poverty) and important economical position in SE Asia and the world [18].

As there are no vaccines or drugs available for DENV, control programmes for DENV transmission are often focused on managing mosquito populations. To determine the nature of mosquito populations, entomological surveys are usually conducted within routine control programmes [19]. For many years, the standard protocol has relied on traditional sampling (*Stegomyia* indices) which is based solely on the presence of larvae [20]. Indicators of DENV vector abundance (mainly *Ae. aegypti*) were based on larval surveys of container habitats and

the calculation of various indices, namely House Index (HI: percentage of houses infested with larvae or pupae), Breteau Index (BI: number of larvae or pupae positive containers per 100 houses examined) and Container Index (CI: percentage of water-holding containers found to be infested with larvae or pupae) [20]. These indices can facilitate understanding of vector ecology in a given control area, but also serve as useful measures to determine the success of intervention strategies. However, these traditional sampling methods have shortcomings by measuring only the abundance of larvae and not determining species, which therefore may be poor predictors of the abundance of adult vector mosquitoes that are responsible for transmission [20]. Following consideration of these issues, Focks (2003) suggested that pupal/demographic survey methods were developed to replace the more traditional larval indices [20]. The pupae index is based on counting the number of pupae per container and identifying which container types are responsible for the largest number of adult mosquitoes. It assumes the ability to predict the potential of DENV epidemics more accurately than the traditional HI, which does not necessarily correlate with DENV transmission [21,22].

Despite the limitation of traditional sampling methods, many studies have continued to focus on indices from larval stages of the mosquito, e.g. [23–28]. However, the importance of developing improved and locally appropriate entomological surveys in DENV endemic areas is increasingly recognised [20]. In Indonesia, variations in dengue disease reporting make it important to further understand the entomological differences between areas with different DENV risk i.e., in endemic areas (defined here as an area that has regularly reported DENV cases in the three years preceding this study, 2009–2011), sporadic areas (defined here as an area which has had an irregular number of DENV cases reported in the three years preceding this study, 2009–2011) and compare these to dengue free areas (defined here as an area with no reports of dengue disease in the three years preceding this study, 2009–2011). Comparisons of mosquito populations and local habits in these different areas are more likely to indicate the key entomological differences that can inform potential points of intervention, and the validity of the various indices and survey methods.

In this study we applied traditional larvae indices, the pupae index as well as adult mosquito collections and species identification to compare and enhance the validity of entomological survey results in villages with different dengue endemicity in the Banyumas Regency of Java. By comparing these traditional indices to newer indicators with respect to their ability to predict dengue risk, we aimed to better understand the local dengue transmission processes. Overall our data help to fill important gaps in our knowledge of dengue transmission and associated ecology/human behaviour in this area of SE Asia and inform local prevention strategies. Our observations may be relevant beyond the study area by informing entomological surveys elsewhere. By determining the most representative factors to predict/analyse mosquito populations and transmission risk, entomology surveys can be done in an effective and more efficient manner.

Methods

Description of the study area

The study site used in this analysis is the Banyumas Regency, located in the southwest of Central Java Province, Indonesia (Fig 1). Coordinates for this location are as follows: 108° 39' 17"–109° 27' 15" East longitude, and 7° 15' 05"–7° 37' 10" South latitude. The total area is 132,760 km², with a population of 1.85 Million inhabitants at a male to female ratio of 50:50. Banyumas Regency consists of 27 sub districts, and has 39 community health centres and a total of 331 villages. The environment in Java is characterized by a tropical monsoonal climate, with a dry season lasting approximately 6 months and a heavy monsoon the rest of year. Total annual

Location of Entomology Survey in Banyumas Regency

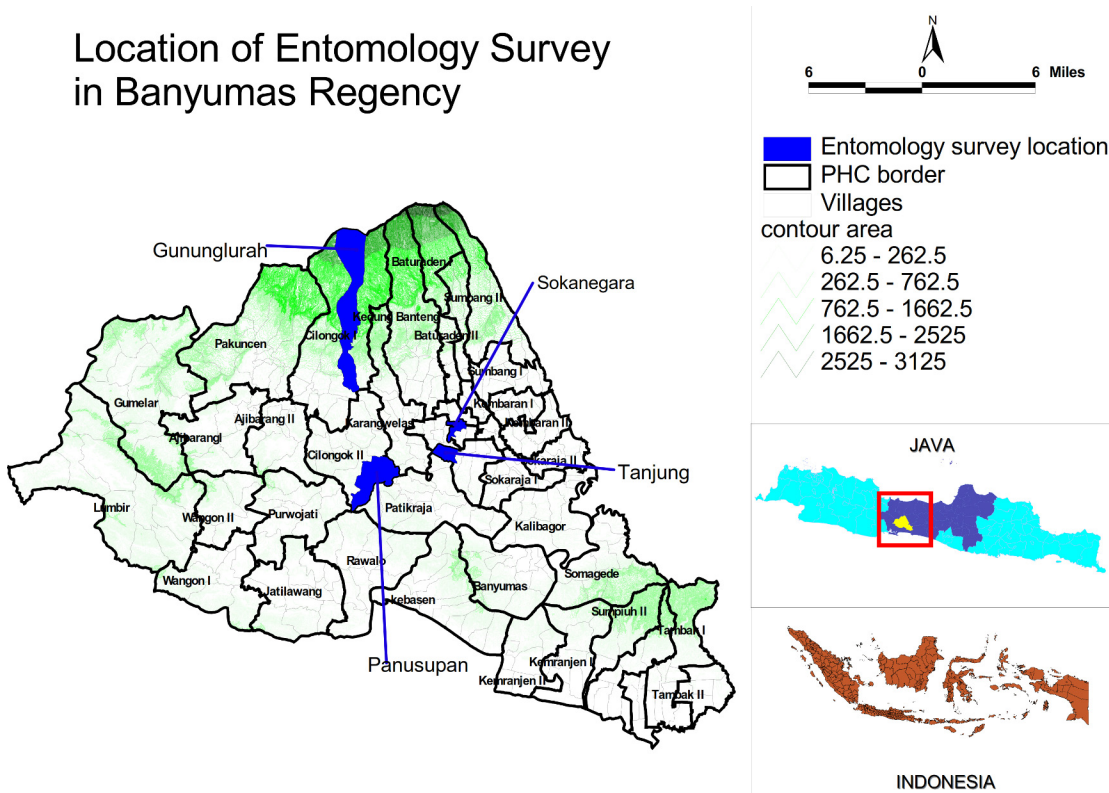


Fig 1. Location of the entomology survey areas in the Banyumas regency. Black areas indicate the locations of the entomology survey: Tanjung, Sokanegara, Panusupan and Gunung Lurah. This map also shows the contours of the area (in meters). Lower right, map of Java with Banyumas regency (in white) and also whole map of Indonesia.

doi:10.1371/journal.pntd.0004500.g001

precipitation averages at 1755 mm (69.1 inches) and there are 2975 hours of sunshine on average per year.

Household studies and seasonal analysis

In order to determine the differences in mosquito population density between seasons, entomological surveys were carried out twice, once in the dry and once in the rainy season. During the dry season three independent villages were selected on the basis of differing endemicity status and their spread across the region (Fig 1): DENV endemic: Tanjung village, South Purwokerto Community Health Centre and Sokanegara, East Purwokerto Community Health Centre; DENV sporadic: Panusupan village, Cilongok I Community Health Centre; DENV free: Gunung Lurah village, Cilongok II Community Health Centre.

The endemicity status criteria are based on “The Technical Manual Eradication of Dengue Mosquito-borne Diseases, Indonesian Ministry of Health” (1992) [29]. The determination of DENV status was made before the survey began, based on reported dengue cases from the Banyumas Regency Health Office. All suspected DENV cases are reported to the health office, however not all cases are confirmed. It is important to note that all reported cases are severe and require hospitalisation. Therefore it is likely there is under reporting of actual cases across the regency. The dry season survey was conducted from May-June 2012, while the rainy season survey was conducted from January-February 2013. One additional village, Sokanegara village (DENV endemic) was added to the survey in the rainy season. In each village, 100 houses were chosen by simple random sampling for the entomological surveys, resulting in a total of 300

houses being analysed in the dry season and 400 houses in the rainy season. The individual locations of the entomological surveys are shown in Fig 1. An overview of environmental conditions and characteristics of the four villages is shown in Table 1.

Larval/pupae surveys and adult mosquito collections

Larvae collection was carried out in every container both inside and outside the participating houses. To measure the entomological parameters the House Index (HI: percentage of houses infested with larvae and/or pupae), Container Index (CI: percentage of water-holding containers infested with larvae or pupae), Breteau Index (BI: number of positive containers per 100 houses inspected), Pupae Index (PI: number of pupae per 100 houses inspected) and Free Larvae Index (FLI: the percentage of houses without larvae) were determined [20]. 100 houses is the sample size recommended by the Indonesian Ministry of Health, in “The Technical Manual Eradication of Dengue Mosquito-borne Diseases, Indonesian Ministry of Health” (1992) [29]. The interpretation of transmission risk levels of each village was made based on the larvae index, as described in the WHO document “A review of entomological sampling methods and indicators for dengue vectors” [20]. The survey also included a description of all containers, both artificial and natural in each participant’s house. Identification of the recovered larvae was based on the key identification criteria as described by Stojanovich and Scott, 1965 [30].

Insect collections were carried out using back-pack aspirators to capture adult mosquitoes in resting and flying positions. Areas inside the house where mosquitoes normally rest were focused on. For example, *Ae. aegypti* mosquitoes prefer to rest in dark, shielded, humid areas on hanging objects such as clothes and curtains and on walls. Adult mosquito capture was carried out between 8–11 AM for around 20 minutes per house (100 houses per village). Identification of adult mosquitoes was conducted by using key identification criteria as described earlier [30]. Sample sizes of 100 houses were used as recommended.

Statistical tests

Confidence intervals (C.I.) 95%, t test, chi square test were calculated by using IBM SPSS Statistic 21.

Table 1. Description of the study sites in Banyumas Regency, Java. Details of the ecology and population of the four study sites are provided.

| Study sites | Ecological description | | | | | | | | Dengue cases in the years before the survey | | |
|---------------------------|------------------------|------------|------------------------------------|--------------------|--|---|------------------------------------|-----------------------|---|------|------|
| | Urban/rural | Population | Elevation (metres above sea level) | Coverage area (ha) | Rainfall in dry season (Apr-Sept (mm)) | Rainfall in rainy season (Oct-March) (mm) | Range of temperature (in °Celsius) | Range of humidity (%) | 2009 | 2010 | 2011 |
| Tanjung (DENV endemic) | Urban | 9696 | 65 | 149 | 796 | 1858 | 28–32 | 73–91 | 7 | 13 | 14 |
| Sokanegara (DENV endemic) | Urban | 7987 | 75 | 119 | 731 | 1705 | 30–38 | 57–83 | 9 | 17 | 6 |
| Panusupan (DENV sporadic) | Rural | 7627 | 200 | 775 | 885 | 2065 | 29–36 | 70–87 | 1 | 2 | 0 |
| Gunung Lurah (DENV free) | Rural | 7120 | 400–700 | 878 | 765 | 1785 | 25–34 | 57–83 | 0 | 0 | 0 |

doi:10.1371/journal.pntd.0004500.t001

Table 2. Risk level of dengue transmission based on larvae indices in the 4 study areas in the dry and rainy seasons. The following table summarizes House index (HI), Breteau Index (BI), Container Index (CI) and Free Larvae Index (FLI). Risk level determination according to AHA Brown [20]; C.I.: confidence interval.

| Village | DENV status | Larvae index, dry season | | | | | Larvae Index, rainy season | | | | |
|--------------|-------------|--------------------------|---------------|---------------|----------------|------------|----------------------------|---------------|---------------|----------------|------------|
| | | HI (95% C.I.) | BI (95% C.I.) | CI (95% C.I.) | FLI (95% C.I.) | Risk Level | HI (95% C.I.) | BI (95% C.I.) | CI (95% C.I.) | FLI (95% C.I.) | Risk Level |
| Tanjung | Endemic | 16 (8–23) | 16 (8–23) | 10 (5–14) | 84 (76–91) | Medium | 18 (10–26) | 18 (10–26) | 4 (2–5) | 82(74–90) | Medium |
| Sokanegara | Endemic | - | - | - | - | - | 18 (10–26) | 18 (10–26) | 5 (2–6) | 82 (74–90) | Medium |
| Panusupan | Sporadic | 25 (16–33) | 35 (25–44) | 22 (15–28) | 75 (66–83) | Medium | 44 (34–53) | 71 (62–79) | 13 (9–15) | 56 (46–65) | High |
| Gunung Lurah | free | 3 (0–6) | 3 (0–6) | 3 (0–5) | 97 (93–100) | Low | 16 (8–23) | 19 (11–26) | 3 (1–4) | 84 (76–91) | Medium |
| Average | | 15 | 18 | 11 | 85 | | 24 | 31 | 6 | 76 | |

doi:10.1371/journal.pntd.0004500.t002

Ethical statement

Studies conducted here (data collection of mosquito breeding sites, mosquito egg collections) were carried out with ethical approval from the University of Glasgow (Project Number: 2012082) and the Ministry of National Education, Faculty of Medicine Gadjah Mada University, Medical and Health Research Ethics Committee (KE/FK/323/FC). No data involving human participants were collected in this study.

Results

Larvae and pupae: survey results inside and outside houses

Following the field surveys conducted during the dry and rainy seasons, the HI and BI indices in the rainy season were found to be higher than in the dry season: the average HI and BI in all villages in the rainy season were 24 and 31, respectively, higher than in the dry season (15 and 18, respectively). On the other hand, the CI was lower in the rainy season (Table 2) probably because more containers were found in the rainy seasons in all villages. Thus compared to the dry season, more mosquito larvae were found during the rainy season. Panusupan (DENV sporadic) showed the lowest free larvae indices (FLI), and was classed as a high risk level of DENV transmission compared to other villages using this index.

To determine whether larvae density correlated with number of DENV cases occurring after the survey, updated information on the number of dengue cases from the Banyumas Regency Health Office was obtained (Table 3).

Based on the report, Tanjung and Sokanegara (DENV endemic) which were classified as medium risk level according to the indices above, continued to have more dengue cases in 2012

Table 3. Mosquito larvae density in dry and rainy seasons, and the number of dengue cases in 2012 and 2013. Dengue case numbers following the surveys were obtained from Banyumas Health Officer’s Report.

| Village | Endemicity status | Mosquito larvae density, dry season (May-June 2012) | Mosquito larvae density, rainy season (Jan-Feb 2013) | Dengue cases, 2012 (July-December) | Dengue cases, 2013 (March-December) |
|--------------|-------------------|---|--|------------------------------------|-------------------------------------|
| Tanjung | Endemic | Medium | Medium | 2 | 15 |
| Sokanegara | Endemic | - | Medium | 13 | 9 |
| Panusupan | Sporadic | Medium | High | 0 | 0 |
| Gunung Lurah | Free area* | Low | Medium | 0 | 1 |

* Despite one case, classed as free based on years preceding these surveys (see Table 1). Larval densities were measured in accordance to Focks et al. [20].

doi:10.1371/journal.pntd.0004500.t003

and 2013 compared to the sporadic and free area. Meanwhile, Panusupan (DENV sporadic, but classed as high risk) reported no dengue cases in 2012 and 2013 and Gunung Lurah (reported as DENV free before 2012 and with a low or medium risk depending on index used) reported one dengue case in 2013. In addition to calculating the various indices as outlined above, species identification of the collected larvae was performed (Fig 2).

Numbers of larvae in all villages were higher in the rainy season than in the dry season. *Ae. aegypti* and *Ae. albopictus* were the dominant species in Tanjung, Sokanegara (DENV endemic) and Panusupan (DENV sporadic). *Culex sp.* were identified in low numbers only in Panusupan (DENV sporadic) and Gunung Lurah (DENV free). Next pupae were assessed in each village and the Pupae Indices (PI) used in order to improve the entomology survey. Details of PI (house and container pupae indices) are indicated in Table 4.

The DENV endemic and sporadic areas had higher CPI and HPI than the DENV free area (Gunung Lurah) ($X^2 = 6.60$, $df = 1$, $p\text{-value} = 0.01$). This indicated that in the endemic/sporadic areas, mosquitoes tend to have a more conducive environment to survive from eggs to become

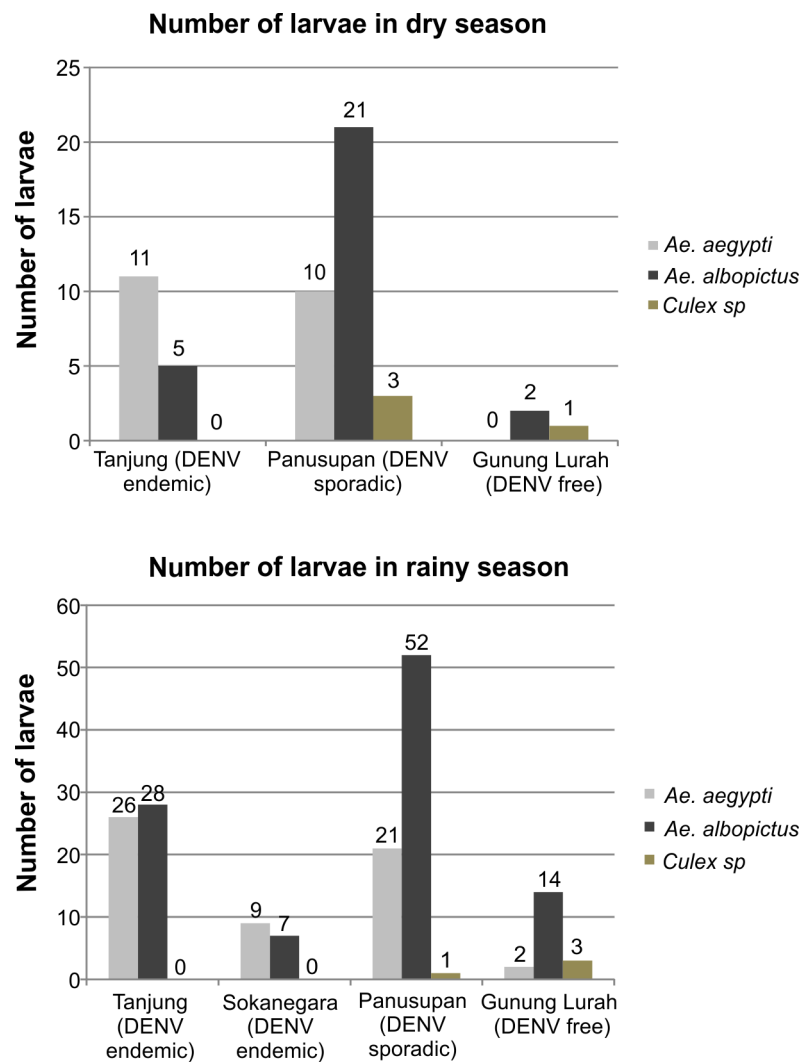


Fig 2. Number of mosquito species within the larvae identified in each of the four villages in dry and rainy season. Each bar represents the mosquito species as indicated in the legend.

doi:10.1371/journal.pntd.0004500.g002

Table 4. Pupae indices for the four villages included in this study.

| Name of Villages | status | CPI (%) (95% C.I.) | | HPI (%) (95% C.I.) | | Pupae/person | | Pupae/house | | Pupae/container | |
|------------------|----------|--------------------|---------|--------------------|-----------|--------------|-------|-------------|-------|-----------------|-------|
| | | dry | rainy | dry | rainy | dry | rainy | dry | rainy | dry | rainy |
| Tanjung | endemic | 4 (0–8) | 4 (0–8) | 5 (0.7–9) | 9(3–15) | 0.02 | 0.061 | 0.09 | 0.26 | 0.05 | 0.05 |
| Sokanegara | endemic | - | 5 (1–9) | - | 11 (5–17) | - | 0.04 | - | 0.21 | - | 0.05 |
| Panusupan | sporadic | 6 (1–11) | 1 (0–3) | 7 (2–12) | 5 (1–9) | 0.03 | 0.02 | 0.11 | 0.11 | 0.07 | 0.01 |
| Gunung Lurah | free | 3 (0–6) | 1 (0–3) | 2 (0–5) | 2 (0–5) | 0.01 | 0.007 | 0.04 | 0.03 | 0.03 | 0.005 |

*CPI: Container Pupae Index; HPI: House Pupae Index; C.I.: confidence interval.

doi:10.1371/journal.pntd.0004500.t004

pupae, and environments with greater survival of mosquitoes to the pupal stage correlated to a higher number of reported dengue cases in endemic areas (Table 3) compared to sporadic and free areas; the high CPI and HPI did not however, correlate with the zero reported cases in Panusupan in the months after the survey. According to Focks (2003) the threshold of dengue transmission is when the pupae/person index (Table 4) ranged between 0.5–1.5 with an optimum air temperature 28°C [20]. Containers found in each house were recorded in order to determine what the dominant mosquito breeding containers were in the various study areas. The main finding of the container survey was that more artificial containers were found in the four villages surveyed compared to natural containers (paired t-test, mean 195, SD 310, $p = 0.003$) (summarized in Table 5). We found more natural containers in sporadic and free areas (Panusupan and Gunung Lurah) compared to endemic areas (Sokanegara and Tanjung), although there was no significant difference ($p = 0.5$) (Table 5). Endemic areas (Tanjung and Sokanegara) are more urbanised (less vegetation, and more densely populated), as described in Table 1. Buckets, water storage containers and traditional bath-tubs were found to be the dominant breeding containers observed in all four villages. In fact, discarded tyres were the containers which had the highest proportion of infestation (53%), this finding is also consistent with other studies [31,32]. Moreover, other artificial containers such as aquariums, water dispensers and flower pots also showed high infestation rates.

Adult mosquito field collections: larval indices did not always correspond to abundance of adult vectors

Measuring adult mosquito numbers is considered to be the most representative quantitative estimate to obtain information about mosquito abundance, as immature stages need to go through several developmental stages to become adult mosquitoes before they can transmit DENV [20]. After identification of the mosquito species in the dry and rainy seasons, the numbers of each species in each area are shown in Fig 3.

The dominant adult mosquito species captured (both seasons combined) during the survey in Tanjung (where *Culex sp.* were dominant only in the rainy season) and Sokanegara (DENV endemic) were *Ae. aegypti*; in Panusupan (DENV sporadic), non dengue transmitting *Culex sp.* were identified as the dominant species (12 in the dry season and 144 in the rainy season); although comparable numbers (to endemic areas) of *Ae. aegypti* were identified. Moreover the combined numbers of *Ae. aegypti* and *Ae. albopictus* were higher in Panusupan than in the dengue endemic areas. However, because of its feeding preference, the role of *Ae. albopictus* has been called into question [3]. This adult collection result is in contrast with the larvae identification, where *Ae. albopictus* was found to be the dominant species in Panusupan and this is likely due to breeding behavior, as *Ae. albopictus* (and *Ae. aegypti*) species frequently breed in

Table 5. Proportion of water-holding containers infested with larvae and/or pupae in the four villages of the study area. Both rainy and dry seasons are taken into account.
Containers are described by type, as indicated.

| Type of water-holding container | Container observed | | | | | | | | | | | | Container positive with larvae | | | | | | Total container positive with larvae | | Total container without larvae | | Proportion infested (%) | | |
|---------------------------------|--------------------|-------|-------|------------|-----|-------|-----------|------|-------|--------------|-----|-------|--------------------------------|----|---------|------|------------|-------|--------------------------------------|-----|--------------------------------|------|-------------------------|--------------------------------------|--------------------------------|
| | Tanjung | | | Sokanegara | | | Panusupan | | | Gunung Lurah | | | Total observed | | Tanjung | | Sokanegara | | Panusupan | | Gunung Lurah | | | Total container positive with larvae | Total container without larvae |
| | dry | | rainy | dry | | rainy | dry | | rainy | dry | | rainy | dry | | rainy | dry | | rainy | dry | | rainy | dry | | | |
| | dry | rainy | 75 | 88 | 63 | 63 | 48 | 48 | 48 | 48 | 480 | 3 | 8 | 11 | 4 | 12 | 1 | 0 | 421 | 39 | 62 | 1034 | | 8 | |
| Traditional bath tub | 55 | 198 | 152 | 73 | 344 | 62 | 212 | 1096 | 6 | 5 | 2 | 17 | 24 | 1 | 7 | 1034 | 6 | 6 | 16 | 8 | 41 | 16 | 6 | | |
| Dispenser | 8 | 16 | 16 | 0 | 8 | 0 | 1 | 49 | 2 | 2 | 0 | 0 | 3 | 0 | 1 | 41 | 16 | 16 | 8 | 8 | 41 | 16 | 16 | | |
| Leaf Mirdrib | 0 | 10 | 1 | 1 | 13 | 1 | 49 | 75 | 0 | 0 | 0 | 0 | 2 | 0 | 5 | 68 | 9 | 9 | 7 | 7 | 68 | 9 | 9 | | |
| Used bottles | 0 | 57 | 61 | 1 | 14 | 2 | 105 | 240 | 0 | 0 | 0 | 0 | 5 | 0 | 3 | 232 | 3 | 3 | 8 | 8 | 232 | 3 | 3 | | |
| Refrigerator | 8 | 10 | 17 | 0 | 3 | 0 | 0 | 38 | 1 | 0 | 0 | 0 | 1 | 0 | 0 | 36 | 5 | 5 | 2 | 2 | 36 | 5 | 5 | | |
| Flower pot | 7 | 11 | 1 | 18 | 6 | 3 | 0 | 46 | 1 | 0 | 9 | 1 | 1 | 0 | 0 | 34 | 26 | 26 | 12 | 12 | 34 | 26 | 26 | | |
| Water storage/container | 10 | 57 | 47 | 2 | 91 | 4 | 130 | 341 | 3 | 2 | 1 | 4 | 16 | 0 | 2 | 313 | 8 | 8 | 28 | 28 | 313 | 8 | 8 | | |
| Aquarium | 3 | 5 | 4 | 0 | 4 | 0 | 0 | 16 | 0 | 0 | 1 | 0 | 1 | 0 | 0 | 14 | 13 | 13 | 2 | 2 | 14 | 13 | 13 | | |
| Discarded Tires | 0 | 3 | 4 | 1 | 4 | 0 | 7 | 19 | 0 | 0 | 3 | 1 | 5 | 0 | 1 | 9 | 53 | 53 | 10 | 10 | 9 | 53 | 53 | | |
| Drum | 0 | 40 | 0 | 0 | 0 | 1 | 0 | 41 | 0 | 1 | 0 | 0 | 1 | 0 | 0 | 39 | 5 | 5 | 2 | 2 | 39 | 5 | 5 | | |
| coconut shells | 0 | 0 | 0 | 1 | 7 | 0 | 0 | 8 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 8 | 0 | 0 | 0 | 0 | 8 | 0 | 0 | | |
| | 166 | 482 | 391 | 160 | 557 | 121 | 552 | 2429 | 16 | 18 | 18 | 35 | 71 | 3 | 19 | 2249 | 7 | 7 | 180 | 180 | 2249 | 7 | 7 | | |

doi:10.1371/journal.pntd.0004500.t005

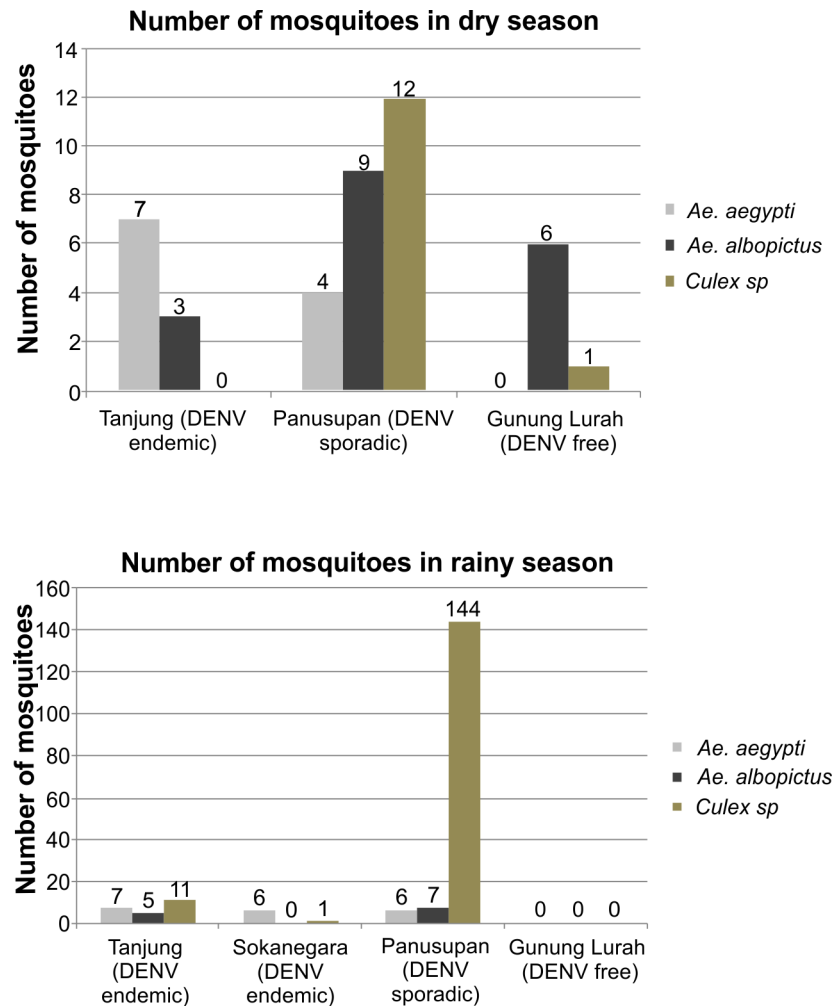


Fig 3. Number of adult mosquito species collected in each of the villages studied in the dry and rainy seasons. Each bar represents the mosquito species indicated as in the legend. Dengue endemicity status is also indicated.

doi:10.1371/journal.pntd.0004500.g003

containers around housing while culicine mosquitoes use different types of habitats. In Gunung Lurah (DENV free), we captured very few adult mosquitoes, and *Ae. albopictus* was the dominant species collected while *Ae. aegypti* was completely absent.

Discussion

Many DENV-endemic countries such as Indonesia, Malaysia and Thailand use entomological surveys as a routine method recommended by WHO to record mosquito populations [19]. Information on mosquito density can then be used in mosquito control efforts and in prevention of DENV transmission [33,34]. Areas with high mosquito populations have usually been treated with larvicides such as organophosphates or temephos in an attempt to prevent outbreaks of DENV. In Indonesia, a Ministry of Health programme encourages community participation in carrying out routine entomology surveys in their homes [35]. Some villages in Indonesia also have trained village health volunteers (VHV) who regularly conduct entomological surveys. Traditional sampling methods i.e. larvae indices were routinely applied over many years to determine mosquito densities in defined areas and the subsequent risk of DENV

transmission. However there can be limitations associated with traditional indices [20–22,36]. To assess the validity and usefulness of these methods, and improve the characterisation of vector populations in our study area, we combined the traditional larvae indices together with the pupae index, species identification and adult mosquito collections.

Our results suggest that traditional larvae indices might not always be an appropriate way of quantifying mosquito populations and dengue transmission risk, as has been previously reported [36]. From the adult mosquito collections (and subsequent species identification), the high larvae indices in Gunung Lurah village (DENV free area; one recent case likely to have been imported from an area where transmission occurred) did not support the transmission of DENV as very few adult mosquitoes were captured in this village. Nonetheless it is important to point out that both *Ae. albopictus* and *Ae. aegypti* were present in DENV-sporadic Panusupan and perhaps differences in vectorial capacity come into play locally. Clearly, our data indicated that larvae density was not always in accordance with the number of DENV cases reported in villages. The pupae survey in this study (Table 4), would suggest that the area of study has a low risk for dengue transmission according to factors previously defined by Focks (2003). The high presence of *Ae. albopictus* larvae in the dengue free area also suggests that presence of vectors alone may not predict transmission; *Ae. albopictus* in this area may have reduced capacity due to their feeding behaviour etc. or possibly reduced competence for DENV. That very few adult mosquitoes were found in 100 houses in Gunung Lurah might be due to a generally unfavourable environment for mosquitoes. These results suggested that high levels of adult *Ae. aegypti* in endemic (and sporadic) areas were a potential indicator of DENV transmission risk. These findings were in agreement with the real numbers of DENV cases which occurred in Gunung Lurah; one reported case in 2013. Adult mosquito numbers (and species identification) may be a useful estimate to obtain information on dengue disease risk, at least in this part of Indonesia as immature stages need to go through several developmental stages in order to become adult mosquitoes able to transmit DENV [20,36]. However, these methods require specialist skills [37] and are not easily transferable to local surveillance programmes. Moreover, while our observations suggest that the usefulness of several indices should be questioned at local level, we stress that underreporting of dengue cases needs to be taken into consideration in the discussion of our results. Improved patient data collection and dengue diagnostics need to be developed, implemented and combined with future mosquito surveillance work in the Regency to support entomological surveillance studies whose accuracy relies on such data. Our findings may encourage such efforts and lead to a more in depth re-evaluation of the observations reported here.

Based on the results of this study, mosquito populations in the regency are higher in the rainy season than in the dry season, for example more mosquito larvae and also adult mosquitoes in three villages were found during the rainy season compared to the dry season. This suggests that health officers and the community should focus their efforts on the beginning of the rainy season. Not surprisingly we also found more potential breeding containers in the rainy season with buckets and water storage containers as predominant water sources in all four villages surveyed. Our findings indicate that villagers can minimize the potential breeding sites for mosquitoes by reducing the presence of artificial containers such as traditional bath-tubs and buckets.

It can be assumed that by reducing the number of these containers, DENV incidence could be minimized. The results from adult mosquito captures in the four villages indicated that *Ae. aegypti* still preferred urban areas (Tanjung and Sokanegara), although in Panusupan (DENV sporadic, rural), *Ae. aegypti* was also observed although the numbers of *Culex sp.* mosquitoes in this village were far higher. *Ae. albopictus* is more likely to be found in rural or suburban areas. These observations are also emphasized by our container survey, where we observed that

natural containers were found more frequently in rural areas (Panusupan and Gunung Lurah) and *Ae. albopictus* is more prevalent than *Ae. aegypti*. These findings are in accordance with previous reviews on the differences in distribution and ecology between *Ae. aegypti* and *Ae. albopictus* which stated that *Ae. albopictus* prefers natural containers [38].

Species identification is important but rarely applied in the field, and often only for research purposes. *Culex* sp. mosquitoes were identified as the dominant adult mosquito type in Panusupan village; this is of interest since this species has not been shown to be a vector of DENV. Vazeille and colleagues stated that *Ae. aegypti* is the most effective vector for dengue viruses and is highly receptive to oral infection; they also demonstrated that *Cx. quinquefasciatus* can be infected by the parenteral route with DENV type 2 but the virus replicated to very low levels, therefore the authors concluded that *Cx. quinquefasciatus* should not be considered a biological vector of DENV [39].

A recent study carried out in Taiwan suggested that various vector indices alone were poor DENV outbreak indicators and each country should evaluate its own situation [40]. We agree with this statement, although we emphasize that better diagnostics needs to be implemented as part of any future studies on this subject in Java. The transmission risk by adult mosquitoes can be influenced by a number of factors that affect the extrinsic incubation period (EIP) and arbovirus/ vector interactions. Indeed, virus and vector genetics, but also gut microbiota and host responses are important factors in DENV-vector interactions [41–53]. Moreover, climatic factors such as temperature and humidity come into play. The influence of temperature on EIPs associated with DENV for example, has been analysed, and was shown to be important for EIP duration [54–56]. These, and other risk factors may vary locally, and could also change over time highlighting the importance of local assessments. At least in the case of Banyumas Regency, our findings also suggest that more prevention efforts should be carried out in the beginning of the rainy season to reduce dengue virus transmission, for example by clearing artificial containers. In summary the observations of this study can form the basis of a better understanding of dengue vector ecology in this part of Indonesia.

Acknowledgments

We gratefully acknowledge the people in Tanjung, Sokanegara, Panusupan and Gunung Lurah villages for cooperating during the entomological surveys in and around their homes. We would also like to thank the staff of the Vector Borne Disease Research and Development Unit Banjarnegara for field research support, as well as the students of the Public Health Department, University of Jenderal Soedirman who participated in the field work. We also thank Triwibowo AG of the Vector Borne Disease Research and Development Unit Salatiga for his advice.

Author Contributions

Conceived and designed the experiments: SPMW MM SMR ES RB AK. Performed the experiments: SPMW SSun SSup. Analyzed the data: SPMW MM SMR ID ES RB AK. Wrote the paper: SPMW MM SMR ES RB AK.

References

1. Bhatt S, Gething PW, Brady OJ, Messina JP, Farlow AW, et al. (2013) The global distribution and burden of dengue. *Nature* 496: 504–507. doi: [10.1038/nature12060](https://doi.org/10.1038/nature12060) PMID: [23563266](https://pubmed.ncbi.nlm.nih.gov/23563266/)
2. Halstead SB (2008) Dengue virus-mosquito interactions. *Annu Rev Entomol* 53: 273–291. PMID: [17803458](https://pubmed.ncbi.nlm.nih.gov/17803458/)

3. Lambrechts L, Scott TW, Gubler DJ (2010) Consequences of the expanding global distribution of *Aedes albopictus* for dengue virus transmission. *PLoS Negl Trop Dis* 4: e646. doi: [10.1371/journal.pntd.0000646](https://doi.org/10.1371/journal.pntd.0000646) PMID: [20520794](https://pubmed.ncbi.nlm.nih.gov/20520794/)
4. Weaver SC, Reisen WK (2010) Present and future arboviral threats. *Antiviral Res* 85: 328–345. doi: [10.1016/j.antiviral.2009.10.008](https://doi.org/10.1016/j.antiviral.2009.10.008) PMID: [19857523](https://pubmed.ncbi.nlm.nih.gov/19857523/)
5. Guzman MG, Halstead SB, Artsob H, Buchy P, Farrar J, et al. (2010) Dengue: a continuing global threat. *Nat Rev Microbiol* 8: S7–16. doi: [10.1038/nrmicro2460](https://doi.org/10.1038/nrmicro2460) PMID: [21079655](https://pubmed.ncbi.nlm.nih.gov/21079655/)
6. Kyle JL, Harris E (2008) Global spread and persistence of dengue. *Annu Rev Microbiol* 62: 71–92. doi: [10.1146/annurev.micro.62.081307.163005](https://doi.org/10.1146/annurev.micro.62.081307.163005) PMID: [18429680](https://pubmed.ncbi.nlm.nih.gov/18429680/)
7. Gubler DJ (2010) The Global Threat of Emergent/Re-emergent Vector-Borne Diseases. *Vector Biology, Ecology and Control*. Washington DC: National Academy of Sciences, Courtesy of the National Academies Press. pp. 39–62.
8. Guzman MG, Harris E (2015) Dengue. *Lancet* 385: 453–465. doi: [10.1016/S0140-6736\(14\)60572-9](https://doi.org/10.1016/S0140-6736(14)60572-9) PMID: [25230594](https://pubmed.ncbi.nlm.nih.gov/25230594/)
9. Mustafa MS, Rasotgi V, Jain S, Gupta V (2015) Discovery of fifth serotype of dengue virus (DENV-5): A new public health dilemma in dengue control. *Med J Armed Forces India* 71: 67–70. doi: [10.1016/j.mjafi.2014.09.011](https://doi.org/10.1016/j.mjafi.2014.09.011) PMID: [25609867](https://pubmed.ncbi.nlm.nih.gov/25609867/)
10. Rohani A, Aidil Azahary AR, Malinda M, Zurainee MN, Rozilawati H, et al. (2014) Eco-virological survey of *Aedes* mosquito larvae in selected dengue outbreak areas in Malaysia. *J Vector Borne Dis* 51: 327–332. PMID: [25540966](https://pubmed.ncbi.nlm.nih.gov/25540966/)
11. Thavara U, Siriyasatien P, Tawatsin A, Asavadachanukorn P, Anantapreecha S, et al. (2006) Double infection of heteroserotypes of dengue viruses in field populations of *Aedes aegypti* and *Aedes albopictus* (Diptera: Culicidae) and serological features of dengue viruses found in patients in southern Thailand. *Southeast Asian J Trop Med Public Health* 37: 468–476. PMID: [17120966](https://pubmed.ncbi.nlm.nih.gov/17120966/)
12. Thongrunkiat S, Maneekan P, Wasinpiyamongkol L, Prummongkol S (2011) Prospective field study of transovarial dengue-virus transmission by two different forms of *Aedes aegypti* in an urban area of Bangkok, Thailand. *J Vector Ecol* 36: 147–152. doi: [10.1111/j.1948-7134.2011.00151.x](https://doi.org/10.1111/j.1948-7134.2011.00151.x) PMID: [21635652](https://pubmed.ncbi.nlm.nih.gov/21635652/)
13. Khin MM, Than KA (1983) Transovarial transmission of dengue 2 virus by *Aedes aegypti* in nature. *Am J Trop Med Hyg* 32: 590–594. PMID: [6859404](https://pubmed.ncbi.nlm.nih.gov/6859404/)
14. Karyanti MR, Uiterwaal CS, Kusriastuti R, Hadinegoro SR, Rovers MM, et al. (2014) The changing incidence of dengue haemorrhagic fever in Indonesia: a 45-year registry-based analysis. *BMC Infect Dis* 14: 412. doi: [10.1186/1471-2334-14-412](https://doi.org/10.1186/1471-2334-14-412) PMID: [25064368](https://pubmed.ncbi.nlm.nih.gov/25064368/)
15. Setiati ET, F. P. Wagenaar J, D. de Kruijff M, T.A Mairuhu A, C.M van Grop E, et al. (2006) Changing Epidemiology of Dengue Haemorrhagic Fever in Indonesia. *Dengue Bulletin* 2006 1–4.
16. Suwandono A, Kosasih H, Nurhayati, Kusriastuti R, Harun S, et al. (2006) Four dengue virus serotypes found circulating during an outbreak of dengue fever and dengue haemorrhagic fever in Jakarta, Indonesia, during 2004. *Trans R Soc Trop Med Hyg* 100: 855–862. PMID: [16507313](https://pubmed.ncbi.nlm.nih.gov/16507313/)
17. Ong SH, Yip JT, Chen YL, Liu W, Harun S, et al. (2008) Periodic re-emergence of endemic strains with strong epidemic potential—a proposed explanation for the 2004 Indonesian dengue epidemic. *Infect Genet Evol* 8: 191–204. doi: [10.1016/j.meegid.2007.12.005](https://doi.org/10.1016/j.meegid.2007.12.005) PMID: [18243816](https://pubmed.ncbi.nlm.nih.gov/18243816/)
18. Tan M, Kusriastuti R, Savioli L, Hotez PJ (2014) Indonesia: an emerging market economy beset by neglected tropical diseases (NTDs). *PLoS Negl Trop Dis* 8: e2449. doi: [10.1371/journal.pntd.0002449](https://doi.org/10.1371/journal.pntd.0002449) PMID: [24587452](https://pubmed.ncbi.nlm.nih.gov/24587452/)
19. WHO (2009) *Dengue: Guidelines for Diagnosis, Treatment, Prevention and Control*. Geneva, Switzerland: WHO. pp. 1–160.
20. Focks DA (2003) *A Review of Entomological Sampling Methods and Indicators for Dengue Vectors*. Florida, USA: World Health Organization on behalf of the Special Programme for Research and Training in Tropical Diseases.
21. Barbazan P, Tuntaprasart W, Souris M, Demoraes F, Nitatpattana N, et al. (2008) Assessment of a new strategy, based on *Aedes aegypti* (L.) pupal productivity, for the surveillance and control of dengue transmission in Thailand. *Ann Trop Med Parasitol* 102: 161–171. doi: [10.1179/136485908X252296](https://doi.org/10.1179/136485908X252296) PMID: [18318938](https://pubmed.ncbi.nlm.nih.gov/18318938/)
22. Focks D, Alexander N (2006) *Multicountry study of *Aedes aegypti* pupal productivity survey methodology: findings and recommendations*. Geneva: World Health Organization on behalf of the Special Programme for Research and Training in Tropical Diseases.
23. Sanchez L, Cortinas J, Pelaez O, Gutierrez H, Concepcion D, et al. (2010) Breteau Index threshold levels indicating risk for dengue transmission in areas with low *Aedes* infestation. *Trop Med Int Health* 15: 173–175. doi: [10.1111/j.1365-3156.2009.02437.x](https://doi.org/10.1111/j.1365-3156.2009.02437.x) PMID: [20409286](https://pubmed.ncbi.nlm.nih.gov/20409286/)

24. Sanchez L VV, Alfonso L, Marquetti MC, Guzman MG, Bisset J, et al (2006) *Aedes aegypti* larval indices and risk for dengue epidemics. *Emerg Infect Dis* 12: 800–806. PMID: [16704841](#)
25. Gurtler RE, Garelli FM, Coto HD (2009) Effects of a five-year citywide intervention program to control *Aedes aegypti* and prevent dengue outbreaks in northern Argentina. *PLoS Negl Trop Dis* 3: e427. doi: [10.1371/journal.pntd.0000427](#) PMID: [19399168](#)
26. Chadee DD, Williams FL, Kitron UD (2005) Impact of vector control on a dengue fever outbreak in Trinidad, West Indies, in 1998. *Trop Med Int Health* 10: 748–754. PMID: [16045461](#)
27. Sulaiman S, Pawanchee ZA, Arifin Z, Wahab A (1996) Relationship between Breteau and House indices and cases of dengue/dengue hemorrhagic fever in Kuala Lumpur, Malaysia. *J Am Mosq Control Assoc* 12: 494–496. PMID: [8887232](#)
28. Correa PR, Franca E, Bogutchi TF (2005) *Aedes aegypti* infestation and occurrence of dengue in the city of Belo Horizonte, Brazil. *Rev Saude Publica* 39: 33–40. PMID: [15654458](#)
29. Indonesia MoH (1992) Technical manual eradication of dengue mosquito-borne diseases. Jakarta: Indonesian Ministry of Health.
30. Stojanovich CHS, H.G (1965) Illustrated Key to Mosquitoes of Vietnam. Atlanta: U.S. Department of Health, Education and Welfare, Public Health Service, CDC Atlanta.
31. Williams CR, Johnson PH, Ball TS, Ritchie SA (2013) Productivity and population density estimates of the dengue vector mosquito *Aedes aegypti* (*Stegomyia aegypti*) in Australia. *Med Vet Entomol* 27: 313–322. doi: [10.1111/j.1365-2915.2012.01051.x](#) PMID: [23205694](#)
32. Nyamah MA, Sulaiman S, Omar B (2010) Categorization of potential breeding sites of dengue vectors in Johor, Malaysia. *Trop Biomed* 27: 33–40. PMID: [20562811](#)
33. Ooi E- E, Goh K- T, Gubler DJ (2006) Dengue Prevention and 35 Years of Vector Control in Singapore. *Emerging Infectious Diseases* 12: 887–893. PMID: [16707042](#)
34. Scott TW, Morrison AC (2010) Vector dynamics and transmission of dengue virus: implications for dengue surveillance and prevention strategies: vector dynamics and dengue prevention. *Curr Top Microbiol Immunol* 338: 115–128. doi: [10.1007/978-3-642-02215-9_9](#) PMID: [19802582](#)
35. Ministry of Health I (2011) Module of Dengue Fever Control. In: Handoko DP, E.B; Hartoyo S, editor. Jakarta: Ministry of Health, Indonesia. pp. 120.
36. Bowman LR, Runge-Ranzinger S, McCall PJ (2014) Assessing the relationship between vector indices and dengue transmission: a systematic review of the evidence. *PLoS Negl Trop Dis* 8: e2848. doi: [10.1371/journal.pntd.0002848](#) PMID: [24810901](#)
37. Sivagnaname N, Gunasekaran K (2012) Need for an efficient adult trap for the surveillance of dengue vectors. *Indian J Med Res* 136: 739–749. PMID: [23287120](#)
38. Higa Y (2011) Dengue Vectors and their spatial distribution. *Tropical Medicine and Health* 39: 17–27.
39. Vazeille M RL, Mousson L, Failloux AB (2003) Low oral receptivity for dengue type 2 viruses of *Aedes albopictus* from Southeast Asia compared with that of *Aedes aegypti*. *Am J Trop Med Hyg* 68: 203–208. PMID: [12641412](#)
40. Chang F-S, Tseng Y-T, Hsu P-S, Chen C-D, Lian I-B, et al. (2015) Re-assess Vector Indices Threshold as an Early Warning Tool for Predicting Dengue Epidemic in a Dengue Non-endemic Country. *PLoS Negl Trop Dis* 9: e0004043. doi: [10.1371/journal.pntd.0004043](#) PMID: [26366874](#)
41. Fansiri T, Fontaine A, Diancourt L, Caro V, Thaisomboonsuk B, et al. (2013) Genetic mapping of specific interactions between *Aedes aegypti* mosquitoes and dengue viruses. *PLoS Genet* 9: e1003621. doi: [10.1371/journal.pgen.1003621](#) PMID: [23935524](#)
42. Pongsiri A, Ponlawat A, Thaisomboonsuk B, Jarman RG, Scott TW, et al. (2014) Differential susceptibility of two field *aedes aegypti* populations to a low infectious dose of dengue virus. *PLoS One* 9: e92971. doi: [10.1371/journal.pone.0092971](#) PMID: [24664142](#)
43. Bennett KE, Flick D, Fleming KH, Jochim R, Beaty BJ, et al. (2005) Quantitative trait loci that control dengue-2 virus dissemination in the mosquito *Aedes aegypti*. *Genetics* 170: 185–194. PMID: [15781707](#)
44. Bosio CF, Fulton RE, Salasek ML, Beaty BJ, Black WCt (2000) Quantitative trait loci that control vector competence for dengue-2 virus in the mosquito *Aedes aegypti*. *Genetics* 156: 687–698. PMID: [11014816](#)
45. Gomez-Machorro C, Bennett KE, del Lourdes Munoz M, Black WCt (2004) Quantitative trait loci affecting dengue midgut infection barriers in an advanced intercross line of *Aedes aegypti*. *Insect Mol Biol* 13: 637–648. PMID: [15606812](#)
46. Behura SK, Gomez-Machorro C, deBruyn B, Lovin DD, Harker BW, et al. (2014) Influence of mosquito genotype on transcriptional response to dengue virus infection. *Funct Integr Genomics* 14: 581–589. doi: [10.1007/s10142-014-0376-1](#) PMID: [24798794](#)

47. Charan SS, Pawar KD, Severson DW, Patole MS, Shouche YS (2013) Comparative analysis of midgut bacterial communities of *Aedes aegypti* mosquito strains varying in vector competence to dengue virus. *Parasitol Res* 112: 2627–2637. doi: [10.1007/s00436-013-3428-x](https://doi.org/10.1007/s00436-013-3428-x) PMID: [23636307](https://pubmed.ncbi.nlm.nih.gov/23636307/)
48. Chauhan C, Behura SK, DeBruyn B, Lovin DD, Harker BW, et al. (2012) Comparative expression profiles of midgut genes in dengue virus refractory and susceptible *Aedes aegypti* across critical period for virus infection. *PLoS One* 7: e47350. doi: [10.1371/journal.pone.0047350](https://doi.org/10.1371/journal.pone.0047350) PMID: [23077596](https://pubmed.ncbi.nlm.nih.gov/23077596/)
49. Behura SK, Gomez-Machorro C, Harker BW, deBruyn B, Lovin DD, et al. (2011) Global cross-talk of genes of the mosquito *Aedes aegypti* in response to dengue virus infection. *PLoS Negl Trop Dis* 5: e1385. doi: [10.1371/journal.pntd.0001385](https://doi.org/10.1371/journal.pntd.0001385) PMID: [22102922](https://pubmed.ncbi.nlm.nih.gov/22102922/)
50. Xi Z, Ramirez JL, Dimopoulos G (2008) The *Aedes aegypti* toll pathway controls dengue virus infection. *PLoS Pathog* 4: e1000098. doi: [10.1371/journal.ppat.1000098](https://doi.org/10.1371/journal.ppat.1000098) PMID: [18604274](https://pubmed.ncbi.nlm.nih.gov/18604274/)
51. Souza-Neto JA, Sim S, Dimopoulos G (2009) An evolutionary conserved function of the JAK-STAT pathway in anti-dengue defense. *Proc Natl Acad Sci U S A* 106: 17841–17846. doi: [10.1073/pnas.0905006106](https://doi.org/10.1073/pnas.0905006106) PMID: [19805194](https://pubmed.ncbi.nlm.nih.gov/19805194/)
52. Sanchez-Vargas I, Scott JC, Poole-Smith BK, Franz AW, Barbosa-Solomieu V, et al. (2009) Dengue virus type 2 infections of *Aedes aegypti* are modulated by the mosquito's RNA interference pathway. *PLoS Pathog* 5: e1000299. doi: [10.1371/journal.ppat.1000299](https://doi.org/10.1371/journal.ppat.1000299) PMID: [19214215](https://pubmed.ncbi.nlm.nih.gov/19214215/)
53. Ramirez JL, Souza-Neto J, Torres Cosme R, Rovira J, Ortiz A, et al. (2012) Reciprocal tripartite interactions between the *Aedes aegypti* midgut microbiota, innate immune system and dengue virus influences vector competence. *PLoS Negl Trop Dis* 6: e1561. doi: [10.1371/journal.pntd.0001561](https://doi.org/10.1371/journal.pntd.0001561) PMID: [22413032](https://pubmed.ncbi.nlm.nih.gov/22413032/)
54. Carrington LB, Armijos MV, Lambrechts L, Scott TW (2013) Fluctuations at a low mean temperature accelerate dengue virus transmission by *Aedes aegypti*. *PLoS Negl Trop Dis* 7: e2190. doi: [10.1371/journal.pntd.0002190](https://doi.org/10.1371/journal.pntd.0002190) PMID: [23638208](https://pubmed.ncbi.nlm.nih.gov/23638208/)
55. Chan M, Johansson MA (2012) The incubation periods of Dengue viruses. *PLoS One* 7: e50972. doi: [10.1371/journal.pone.0050972](https://doi.org/10.1371/journal.pone.0050972) PMID: [23226436](https://pubmed.ncbi.nlm.nih.gov/23226436/)
56. Carrington LB, Seifert SN, Armijos MV, Lambrechts L, Scott TW (2013) Reduction of *Aedes aegypti* vector competence for dengue virus under large temperature fluctuations. *Am J Trop Med Hyg* 88: 689–697. doi: [10.4269/ajtmh.12-0488](https://doi.org/10.4269/ajtmh.12-0488) PMID: [23438766](https://pubmed.ncbi.nlm.nih.gov/23438766/)