

## REVIEW

# Characterizing dengue transmission in rural areas: A systematic review

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## Abstract

Dengue has historically been considered an urban disease associated with dense human populations and the built environment. Recently, studies suggest increasing dengue virus (DENV) transmission in rural populations. It is unclear whether these reports reflect recent spread into rural areas or ongoing transmission that was previously unnoticed, and what mechanisms are driving this rural transmission. We conducted a systematic review to synthesize research on dengue in rural areas and apply this knowledge to summarize aspects of rurality used in current epidemiological studies of DENV transmission given changing and mixed environments. We described how authors defined rurality and how they defined mechanisms for rural dengue transmission. We systematically searched PubMed, Web of Science, and Embase for articles evaluating dengue prevalence or cumulative incidence in rural areas. A total of 106 articles published between 1958 and 2021 met our inclusion criteria. Overall, 56% ( $n = 22$ ) of the 48 estimates that compared urban and rural settings reported rural dengue incidence as being as high or higher than in urban locations. In some rural areas, the force of infection appears to be increasing over time, as measured by increasing seroprevalence in children and thus likely decreasing age of first infection, suggesting that rural dengue transmission may be a relatively recent phenomenon. Authors characterized rural locations by many different factors, including population density and size, environmental and land use characteristics, and by comparing their context to urban areas. Hypothesized mechanisms for rural dengue transmission included travel, population size, urban infrastructure, vector and environmental factors, among other mechanisms. Strengthening our understanding of the relationship between rurality and dengue will require a more nuanced definition of rurality from the perspective of DENV transmission. Future studies should focus on characterizing details of study locations based on their

environmental features, exposure histories, and movement dynamics to identify characteristics that may influence dengue transmission.

### Author summary

Dengue is a widespread mosquito-borne viral disease with a large global burden. Previous research has focused on transmission in urban areas and only recently has been identified in rural populations. This systematic review that aims to (1) identify rural indicators used in current studies of dengue transmission; (2) characterize dengue seroprevalence and incidence in rural areas; and (3) summarize those studies that compare the risk of dengue in rural and urban locations.

## Introduction

Dengue is a significant public health challenge with roughly half of the world's population living in endemic countries and more than 3 billion people at risk [1,2]. It is estimated that 96 million symptomatic dengue cases occur per year, which is likely an underestimate due to underreporting and misdiagnosis, particularly in countries with insufficient surveillance systems or lack of diagnostic capacity [3]. Urban centers have historically been thought to be at elevated risk for dengue due to (i) increasing population density that allows vectors to transmit dengue virus (DENV) to large pools of susceptible individuals without flying long distances [4–8]; (ii) increased human movement to and from hot spots of transmission; (iii) infrastructure failure (i.e., breakdown of water and waste management systems); and (iv) human activities leading to an abundance of urban-associated *Aedes aegypti* oviposition sites such as abandoned tires, vessels that can hold small amounts of water, and larger containers to store water for household use [9]. These factors favor vector abundance within urban centers that have the population and other ingredients for sustained arboviral transmission.

Dengue research has therefore largely (but not exclusively) focused on virus transmission in urban areas [2,10–16]. However, an increasing number of studies have described high levels of DENV infections in rural populations [4,17–19]. Some have postulated that the increase in rural DENV transmission is due to the increase in human travel [20]. As transportation infrastructure has developed worldwide, travel between urban and rural areas as well as human mobility between remote regions has increased, facilitating virus spread into and out of rural settings [21]. Additionally, some of the urban features that are commonly considered to drive risk may increasingly be present in more rural areas, including higher population density and habitats suitable for *Ae. aegypti* development. Finally, because DENV infections are often asymptomatic [1], the burden of infection may be underestimated in poor or rural settings where surveillance systems are less developed. It is increasingly recognized that rural DENV transmission may be crucial for maintaining serotype diversity in populations and may also facilitate disease reemergence [22–24].

Definitions of what is rural vary widely in the dengue literature. For example, researchers have defined rural environments based on population size [25], population density [4,17, 25–27], housing density [28], infrastructure or surface cover type (impervious surfaces, vegetation) [29,30], access to urban areas or distance to a road/urban center [31,32], environmental changes (including changes to landscapes, rural production systems, climate, land use, and transportation infrastructure) [27,32,33], or agricultural practices [25]. A definition useful for surveillance and mitigation may be a function, subset, or combination of these factors and

may vary by region. Our overall goal is to summarize rurality indicators used in current epidemiological studies of DENV transmission given changing and mixed environments. To achieve this goal, we conducted 3 subanalyses: (1) to review how authors defined rural and the mechanisms for rural transmission; (2) to summarize those studies that compare the incidence of rural dengue to urban estimates; and (3) to assess dengue seroprevalence and incidence in rural areas.

## Methods

### Study registration

This protocol has been registered with the international prospective register of systematic reviews (PROSPERO) (registration number 92243). The Preferred Reporting Items for Systematic Reviews and Meta-analyses (PRISMA) Statement guided the conduct of this review [34].

### Search strategy

Literature search strategies were developed under the guidance of an information specialist from the Health Science Library at the University of Michigan (GR). Medical Subject Headings (MeSH) and text word searches related to dengue and urbanicity were used in PubMed, Embase, and Web of Science. The following MeSH and keyword terms were used in PubMed ( $n = 324$ ), Embase ( $n = 380$ ), and Web of Science ( $n = 30$ ) articles: (“rural population” or “rural health” or “urbanization”) AND (“dengue” or “Dengue virus”). The search was conducted on 18 November 2021.

### Eligibility

All studies and official reports published in English or Spanish were eligible for inclusion based on the criteria below. We did not exclude articles based on study design or quality, as the strength of evidence from each study was evaluated separately using an adapted version of the Newcastle–Ottawa Scale (NOS) [35]. We also did not limit our study by year of publication. Because there is no international consensus for the definitions of “rural” and “urban” [36–38], the United Nations recommends following definitions based on regional perceptions [38]. Therefore, we considered study sites to be “rural” or “urban” based on the original authors’ designations. It follows that we only considered “urban” study sites if authors had directly compared them to a “rural” location. An alternative approach to our eligibility criteria would be to classify studies based on a predefined list of specific rural characteristics; however, this could bias our review because not all features of rural environments may be reported by all papers.

### Inclusion criteria

We included studies that meet one or more of the following criteria:

- (1) Authors describe the study site as “rural,” or a similar descriptor, or describe a rural–urban continuum in the title or abstract and one or more of the following:
  - (a) mentions at least one positive or probable dengue case; or
  - (b) describes dengue seroprevalence; or
  - (c) describes an outbreak of dengue; or

(2) Studies or official reports of dengue incidence, prevalence, or outbreaks in humans of a larger geographic area where rural estimates could be extracted. For example, the Pan American Health Organization's Health Information Platform for the Americas provides estimates at fine geographic resolution such that rural estimates can be isolated.

### Exclusion criteria

We excluded studies if (1) the study did not provide human outcome data on dengue, including all studies that only mention mosquitoes and/or zoonotic transmission without any description of dengue incidence or prevalence; (2) news articles, commentaries, factsheets, or other source formats that do not provide dengue outcome data; or if (3) the authors described location as "a city" or "urban" in the title or abstract; or (4) if the authors mention the name of the study location and the location has a population greater than 100,000 without providing dengue estimates from a comparable rural study location. The addition of a 100,000-population cutoff was used to exclude large cities, reducing the number of irrelevant studies to be screened.

### Data collection process

Two reviewers (OM and RT) independently extracted data from the included sources on study population, methodology, and reported dengue outcomes. Reviewers resolved disagreements by discussion and a third reviewer (ANMK or JNSE) decided on any unresolved disagreements. See supporting information for more details on the data collection and quality assessment. In brief, the quality of each study was ranked using an adapted version of the NOS (see supporting information for more details).

### Analysis

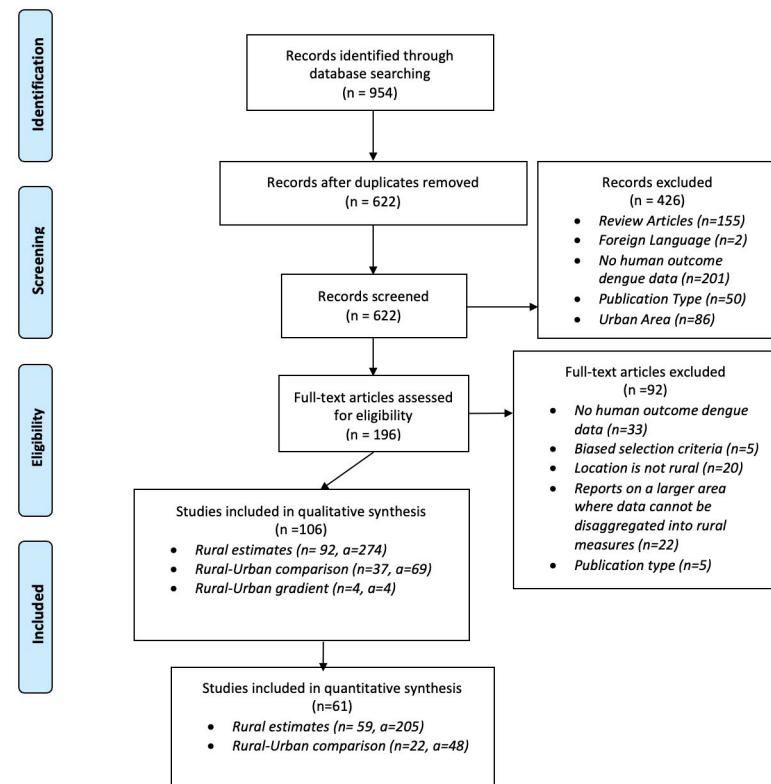
All studies that met our inclusion criteria were considered for qualitative review, including articles that described a relationship but did not provide a numeric point estimate. While we used our NOS subscale to determine which of our hypothesized rural characteristics were also mentioned by study authors, we also extracted characteristics mentioned by study authors separately to create a narrative summary of what has been found in the literature that was unbiased by our prior expectations. In addition to characteristics used to define rurality in the articles included in this review, we also separately extracted authors' hypothesized mechanisms for their measured rural incidence.

Studies were considered for the quantitative analysis if authors provided information about cases that could be approximated to dengue incidence, prevalence, or seroprevalence estimates. In order to include a larger variety of studies with varying research questions, we hand calculated estimates when necessary.

When studies could be grouped by comparable definitions of the exposure and outcome, we conducted a meta-analysis using a random effects model to quantify the overall measure of association for objectives (2) and (3). For rural dengue studies, we also considered how reported incidence might be influenced by key effect modifiers that varied across studies, including population size and the method of data collection.

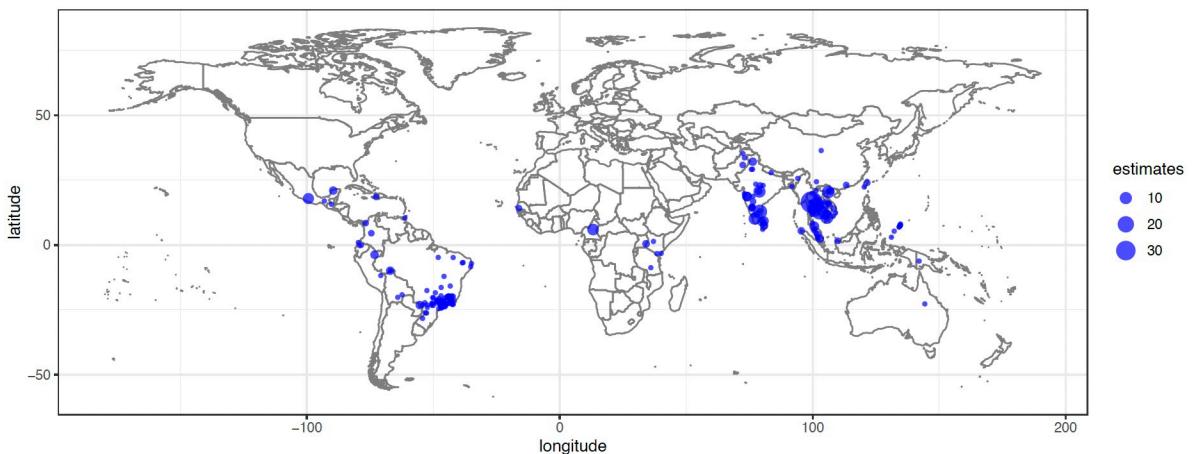
### Results

A total of 106 studies and 347 estimates were included in our qualitative synthesis [4–9,19,25,39–135] (Fig 1). Most estimates used data collected in tropical regions of Southeast Asia, Latin America, and Africa (Fig 2). These studies were divided into 2 categories: (1) articles



**Fig 1. PRISMA diagram of articles included in our analysis where “n” represents the number of articles and “a” represents the number of estimates.** From: Moher D, Liberati A, Tetzlaff J, Altman DG, The PRISMA Group (2009). Preferred Reporting Items for Systematic Reviews and Meta-Analyses: The PRISMA Statement. PLoS Med 6(7): e1000097. doi:10.1371/journal.pmed.1000097.

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**Fig 2. Map of study locations.** The size of dot corresponds to the number of estimates. The basemap is publicly available at <https://www.naturalearthdata.com/downloads/50m-raster-data/50m-bathymetry/>.

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that presented rural–urban comparison estimates ( $n = 37$ ) [4–7,9,25,39,42–44,49,53,59–62,65,69,72,75,78,83,85,86,88,93,95–97,101,109,119,120,125,126,133,135] and (2) articles that presented rural estimates ( $n = 92$ ) [4–9,19,25,40–71,73–78,80–82,84,85,87,89–94,97–100,102–109,111–119,121,122,124–132,134,135]. In addition, we identified 4 studies that modeled dengue along rural–urban gradients [79,82,123,134]. Of these, three observed higher dengue incidences in urban compared to rural areas [32,39,41], whereas one found the opposite [40]. We did not consider the gradient studies any further due to noncomparable analysis methods.

For the qualitative analysis, studies were assessed for reported definitions of rurality and author hypothesized mechanism for rural dengue transmission.

Of the 106 studies that met our inclusion criteria, 45 articles were excluded from the quantitative synthesis (Fig 1) [19,39,40,42,45,47,49,52–56,60,66,67,71,72,76,80,83,85–88,90,95,96,101,104,107,110,111,114,115,117,118,120,122,124,125,127–129,133,134]. Most articles were excluded because they lacked information on variance (73.3%,  $n = 33$ ) or they did not report the underlying population (e.g., they only reported the proportion of fever cases, 48.9%,  $n = 22$ ). Other reasons for exclusion include biased sampling methods and incomparable case information (e.g., a rise in cases after natural disasters). Articles included had data collection dates ranging from 1987 to 2016.

### Objective 1: Qualitative analysis ( $n = 106$ )

**Rural definitions.** For all studies included in our qualitative synthesis, we evaluated the rural definitions and study site descriptions used by the authors (Table 1). One-third of the articles did not provide justification for classifying their study site as rural (34.9%,  $n = 37$ ) [4,5,40–42,47,49,52,56,65–67,72,75,77,78,87,89–93,95–

**Table 1.** Characteristics used by authors in their rural definitions (106 studies included 177 cited characteristics; 37 studies had no definition of rural and many articles used more than one definition).

Definition of Rural	No. of Articles	No. of Articles (% within category)
Undefined	37	37 (100%)
<i>Population Characteristics</i>	49	
Population		28 (57.1%)
Population density		14 (28.6%)
Housing density or number of households		7 (14.3%)
<i>Environmental and Land Usage Characteristics</i>	40	
Terrain characteristics and presence of wildlife		21 (52.5%)
Agriculture		19 (47.5%)
<i>Relative to Urban Areas</i>	30	
Direct comparison to urban*		15 (50.0%)
Distance from urban area		11 (36.7%)
Travel time to urban area		4 (13.3%)
<i>Miscellaneous Characteristics</i>	21	
Description of transportation systems		7 (33.3%)
Limited water access or use		4 (19.1%)
Socioeconomic status		4 (19.1%)
Description of housing or road conditions		4 (19.1%)
Country-wide classification system		2 (9.5%)

\*Studies in the “direct comparison to urban” category defined locations as rural because they did not meet the authors’ criteria for an urban center.

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[97](#),[107](#),[110](#),[111](#),[113](#),[118](#),[120](#),[121](#),[124](#),[125](#),[127](#),[129](#),[130](#),[132](#)]. The remaining 69 studies provided a wide variety of rural definitions. We grouped these definitions into 4 broad categories: population, environment and land use, relationship to urban areas, and miscellaneous, which we further divided into 13 specific subcategories. An article often used a definition that allowed them to be placed in more than 1 category. When definitions related to population characteristics were used, it was more common for authors to consider population size (57.1%,  $n = 28$ ) [[8](#),[19](#),[44](#),[46](#),[48](#),[50](#),[61](#),[63](#),[64](#),[68](#),[70](#),[74](#),[76](#),[82](#),[83](#),[86](#),[94](#),[100](#),[102](#),[103](#),[105](#),[106](#),[108](#),[109](#),[116](#),[119](#),[122](#),[128](#)] than population density (28.6%,  $n = 14$ ) [[6](#),[8](#),[19](#),[48](#),[73](#),[76](#),[83](#),[100](#),[101](#),[103](#),[108](#),[109](#),[115](#),[134](#)] or housing density (14.3%,  $n = 7$ ) [[19](#),[44](#),[61](#),[63](#),[99](#),[126](#),[131](#)]. When authors used environmental and land use definitions of rural, they tended to use various descriptors. Examples include terrain, such as “tropical forests” or “rice fields,” proximity to increased wildlife biodiversity (52.5%,  $n = 21$ ) [[45](#),[48](#),[51](#),[54](#),[55](#),[57](#),[69](#),[71](#),[81](#),[88](#),[99](#),[100](#),[105](#),[106](#),[114](#),[115](#),[117](#),[119](#),[122](#),[134](#),[135](#)] or agricultural practices, including the types of crops and livestock, (47.5%,  $n = 19$ ) [[8](#),[43](#),[45](#),[57](#),[58](#),[62](#)–[64](#),[68](#),[70](#),[71](#),[83](#),[100](#),[104](#),[114](#),[115](#),[119](#),[128](#),[134](#)]. If rural was defined in relation to urban centers, authors commonly used direct comparisons and considered locations to be rural because (a) they were not urban (50.0%,  $n = 15$ ) [[7](#),[9](#),[25](#),[39](#),[43](#),[53](#),[59](#),[62](#),[69](#),[79](#),[83](#),[85](#),[86](#),[109](#),[133](#)]; (b) the distance from an urban location was above a threshold (36.7%,  $n = 11$ ) [[6](#),[50](#),[54](#),[55](#),[63](#),[64](#),[80](#),[82](#),[98](#),[120](#),[124](#)]; or (c) travel time to an urban location was above a threshold (13.3%,  $n = 4$ ) [[60](#),[64](#),[82](#),[84](#)]. Other miscellaneous definitions of rural included quality of transportation systems [[58](#),[60](#),[64](#),[84](#),[106](#),[112](#),[115](#)], water access or usage [[60](#),[64](#),[108](#),[119](#)], quality of housing or roads [[60](#),[88](#),[115](#),[119](#)], socioeconomic status [[68](#),[73](#),[99](#),[126](#)], or a country-wide classification system [[62](#),[126](#)].

**Reported modes of transmission.** We also searched each of the 106 articles used in this review for possible modes of DENV transmission in the rural study sites. Thirty (28.3%) articles did not discuss how transmission occurred (Table 2). Of the remaining 76 (71.7%) articles, one or more modes of transmission were suggested by study authors, most of which were included in author-specified definitions of rural areas (Table 1). We placed the articles into 19 general transmission categories (Table 2) that fell into 4 groups: (1) human population (travel, movement, and population size); (2) urban infrastructure (water storage, improper water management, proximity to roads, increasing infrastructure, and unspecified urbanization, mentioned by 30 articles); (3) vector and environmental factors (mentioned by 34 articles); and (4) other mechanisms (emergence in new areas, underreporting, low levels of prior immunity, and biological variability, mentioned by 18 articles). Some articles also mentioned the importance of local seroprevalence patterns, which define susceptibility of local populations, and disease underreporting associated with poor or rural areas.

## Objective 2: Rural–urban comparison

We identified 48 estimates from 22 articles comparing rural to urban dengue estimates [[4](#),[6](#),[7](#),[9](#),[25](#),[43](#),[44](#),[49](#),[53](#),[59](#)–[62](#),[65](#),[69](#),[75](#),[78](#),[93](#),[97](#),[109](#),[126](#),[135](#)]. The rural–urban comparison studies had an average NOS quality ranking of 3.5, with a standard deviation of 1.1. Eighteen articles and 40 estimates reported values that could be approximated as incidence ratios [[4](#),[6](#),[7](#),[9](#),[44](#),[49](#),[53](#),[60](#)–[62](#),[65](#),[69](#),[78](#),[93](#),[97](#),[109](#),[126](#),[135](#)], and 7 articles with 8 estimates reported prevalence or seroprevalence ratios [[25](#),[43](#),[59](#),[62](#),[65](#),[69](#),[119](#)]. While some of these articles used different study designs, the ratios from these articles could be compared across studies to assess rural versus urban patterns of disease. Overall, 56% ( $n = 22$ ) of these estimates found that dengue was as high or higher in rural areas (Fig 3)

[[4](#),[6](#),[7](#),[9](#),[25](#),[43](#),[44](#),[49](#),[53](#),[59](#),[61](#),[62](#),[65](#),[69](#),[75](#),[78](#),[93](#),[97](#),[109](#),[119](#),[126](#),[135](#)]. This pattern of similar or higher rural dengue compared with urban dengue was similar regardless of population size or

Table 2. Hypothesized mode of dengue virus transmission suggested in articles included in this systematic review.

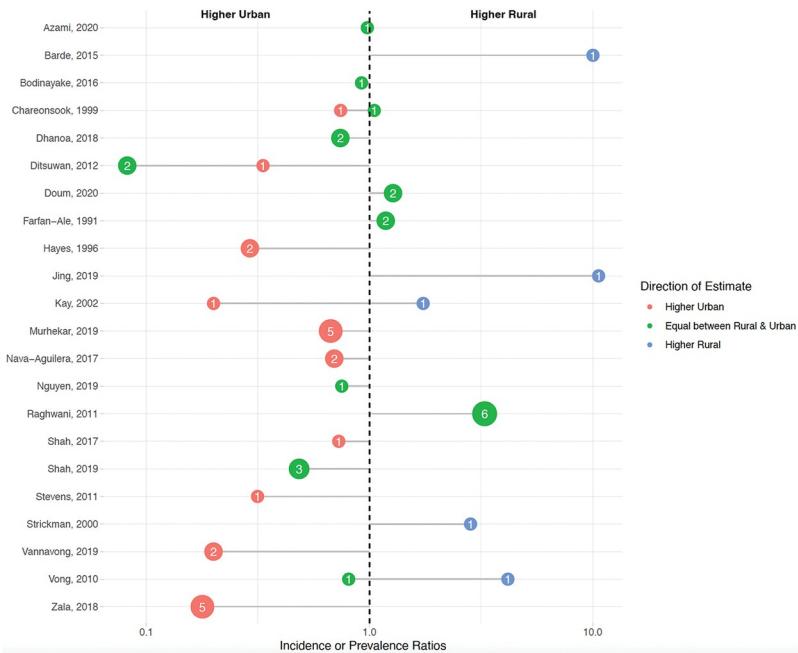
Hypothesized mode of transmission	No. of Articles	References
<b><i>Human population</i></b>	<b>42</b>	
Travel to urban centers	16	[6,7,46,48,49,57,63,69,79,80,82,108,110,119,120,123]
Human movement	18	[19,25,43,48,55,57,59,62,68,69,91,101,106,113,121,122,126,131]
Increasing population density or size	20	[4–6,9,25,39,43,45,55,69,84,93,100,109,110,121,123,125,134]
Introduction through travel, not sustained	3	[39,78,82]
<b><i>Urban infrastructure</i></b>	<b>30</b>	
Increases in water storage	15	[6,9,43,44,49,53,62,64,76,80,84,96,123,127]
Insufficient waste management (increasing container habitat availability)	6	[9,39,43,54,119,123]
Proximity to roads	5	[6,25,101,106,110]
Building infrastructure	7	[9,25,39,75,81,84,119]
Unspecified urbanization	10	[5,9,43,72,81,93,94,109,127,128]
<b><i>Vector and environmental mechanisms</i></b>	<b>34</b>	
Land use and agriculture	7	[59,68,76,96,106,108,134]
Exposure to natural environments	11	[43,51,54,58,68,69,84,88,95,101,134]
Sylvatic cycle	3	[51,63,69]
Climate Change	5	[60,68,93,121,134]
Other vector related mechanisms (i.e., alternate vectors, vector densities, vector spread)	22	[43–45,49,53,58–60,62,64,68,69,72,75,76,93,95,101,110,114,126]
<b><i>Other modes of DENV transmission</i></b>	<b>18</b>	[59,71,72,75,77,80,82,95,98,104,109,111,112,123,125,127,131,135]
Prior underreporting or data collection limitations	7	[59,75,83,95,109,117,123]
Emergence in new area	9	[71,72,80,98,104,111,112,127,135]
Emergence of a new serotype	2	[82,98]
Higher susceptibility in rural areas (due to low prior exposure)	2	[82,131]
Biological variability uncaptured by standard variables	1	[74]
<b>None</b>	<b>30</b>	[40–42,47,50,52,56,61,65–67,70,73,74,77,87,89,90,92,97,99,102,103,107,115,116,124,130,132,133]

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the method of data collection (active versus passive). Several studies assessed dengue cases overtime, showing either that values in rural and urban areas eventually became more similar or that cases were higher in urban populations after years of circulation [4,6,53,61,78].

### Objective 3: Rural dengue

We identified 59 articles (n) with 205 estimates (a) estimating dengue in rural areas [4–9,25,41,43,44,46,48,50,51,57–59,61–65,68–70,73–77,81,82,84,89,91–94,97–100,102–



**Fig 3. Distribution of rural to urban dengue incidence and prevalence ratios.** Each dot represents the average incidence or prevalence ratio grouped by direction of the estimate. The size and number of each dot represents the number of estimates. Red-colored estimates signify that incidence/prevalence was higher in urban areas; green-colored estimates signify that incidence/prevalence was equal between rural and urban areas (i.e., had confidence intervals that crossed the null value); and blue-colored estimates signify that incidence/prevalence was higher in rural areas.

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104,106,108,109,113,116,119,121,126,130–132,135]. These estimates were further categorized into measurements of seroprevalence ( $n = 18$ ,  $a = 37$ ) [25,43,46,50,51,57–59,62,65,68,69,75,81,91,93,109,119], incidence ( $n = 32$ ,  $a = 119$ ) [4–9,41,48,61,62,65,69,70,73,76,77,89,92,99,100,102,104,108,109,113,116,121,126,130–132,135], prevalence ( $n = 5$ ,  $a = 8$ ) [74,97,106,131,132], and outbreak studies ( $n = 8$ ,  $a = 40$ ) [44,63,64,82,84,94,98,103].

**Seroprevalence.** Studies that looked at rural dengue seroprevalence tended to have higher NOS scores (mean = 4.2, standard deviation = 0.6) when compared to the other rural result subgroupings. Among these studies, seroprevalence was not statistically related to population size ( $p = 0.1832$ ). Five studies that stratified rural seroprevalence by age show increasing seroprevalence among children over time [25,57,65,109,119] (S1 Fig). Several seroprevalence articles surveyed large geographic areas, one of which could be divided into smaller villages.

**Rural incidence.** Studies that looked at rural incidence had lower NOS scores (mean = 2.5, standard deviation = 1.1) when compared to other rural result subgroupings. Most of these were community-based studies that included active case finding rather than passive surveillance. The average follow-up time for active surveillance studies was 381.2 days [5–7,62,65,69,70,73,78,99,100,109,126,130–132] compared to the average follow-up time of 1,118.7 days for passive surveillance studies [4,8,9,41,48,61,76,77,89,92,102,108,113,116,121,135]. Estimates collected from passive methods (i.e., hospital referrals) reported lower incidence rates than studies that used more active case finding (i.e., cohort, community surveillance). Passive data collection studies ( $n = 16$ ,  $a = 63$ ) had a pooled annual incidence estimate of 142 cases per 100,000 [95% CI: –537, 822, range: 0, 2,133] (S2 Fig). Active data collection studies ( $n = 16$ ,  $a = 56$ ) had a pooled incidence

estimate of 2,469 cases per 100,000 [95% CI: 1,471, 3,466, range: 0, 17,066] ([S3 Fig](#)). For some studies, the incidence of rural dengue appeared to be increasing over time, but this trend was inconsistent. Moreover, while several articles presented dengue incidence for consecutive years [4,6,61,70,78,121], they did not report information on the predominant serotype (DENV1–4). If the same serotype was circulating during sequential years, yearly estimates may capture multiple or incomplete transmission cycles. Rural incidence appeared to be similar regardless of population size.

**Rural prevalence.** Studies that collected information on rural prevalence had low NOS quality scores (mean = 2.5, standard deviation 0.8). Due to the small number of articles ( $n = 5$ ,  $a = 8$ ) and variable definitions of dengue (e.g., self-reported household dengue or prior history of dengue), we did not further evaluate the effect of potential drivers of rural prevalence [[74](#),[97](#),[106](#),[131](#),[132](#)].

**Rural outbreaks.** Articles that evaluated outbreaks of dengue in rural areas had an average NOS quality score of 2.2 (standard deviation 1.1). There were 8 articles on outbreaks with 40 estimates [[44](#),[63](#),[64](#),[82](#),[84](#),[94](#),[98](#),[103](#)]. Further assessment of rural outbreaks was difficult due to variations in follow-up time, sampling methods, and definitions of probable dengue infections. For example, one article only considered positive cases during a 2-week period [[103](#)], while another study only considered cases with positive neurologic side effects (e.g., altered mental status) [[63](#)]. Additionally, it is unclear if data collection spanned the entirety of the outbreak. Although some articles stratified by age group [[98](#),[103](#)], stratifications were inconsistent.

## Discussion

In this review, we compiled literature on rural dengue with 3 objectives in mind: (1) to identify definitions and mechanisms of rural dengue transmission; (2) to summarize those studies that compare dengue risk in rural areas with urban areas; and (3) to assess the burden of dengue in rural areas. Though 37% of 106 studies did not define the term “rural” at all, the remaining studies most commonly defined rural dengue based on population characteristics and environmental and land usage characteristics ([Table 1](#)). The overall level of incidence in rural areas was commonly related to human population, the level of local infrastructure, and environmental and vector characteristics ([Table 2](#)) (Objective 1). Based on studies comparing urban and rural dengue, we find evidence that the risk DENV transmission in rural areas is substantial and is often similar to or sometimes higher than the risk in urban areas ([Fig 3](#)). This suggests that rural regions might be an important source of dengue cases (Objective 2). Based on those articles focusing on rural areas alone, burden was nontrivial, and some data suggested that rural dengue incidence might be increasing over time ([S1 Fig](#)) (Objective 3).

### Objective 1: Identify definitions and mechanisms of rural dengue transmission

Study authors used a variety of rural definitions based on political definitions, population characteristics, environmental and land usage patterns, distance/travel time to urban centers, and transportation systems ([Table 1](#)). Underlying these study definitions of rurality are proposed or assumed mechanisms and factors related to rural DENV transmission ([Table 2](#)). The varied definitions and wide range of proposed mechanisms highlight complexities in the understanding of rural DENV transmission dynamics and ability to compare results across studies. This finding is consistent with prior research that has shown that definitions of “rural” vary across regions [[36](#)–[38](#)] and that multiple features beyond those, which have been typically classified as “rural” characteristics, tend to impact risk.

While many features were only mentioned by a few authors, environmental exposures and local infrastructure were mentioned frequently as both definitions of rural areas (Table 1) and subsequent drivers of dengue incidence. However, in most instances, rural drivers of DENV transmission appeared to be locally specific and did not show a relationship at aggregate levels. For example, we did not observe a quantifiable relationship between population size and rural dengue seroprevalence. This suggests that population size may be a driver of DENV infection in some rural locations but is not sufficient to explain the existing global trends in the literature. Thus, population may not be an adequate proxy for rurality alone. Local differences in population size might be more useful in defining geographic patterns of risk rather than defining absolute transmission levels globally. Unfortunately, we were not powered to assess the impact of population size for incidence studies. Follow-up studies that assess the impact of population size might provide more insight on this relationship.

Most articles we reviewed also described changing features of rural environments associated with increasing transmission. These include (i) a higher rural incidence among politically defined rural areas with multiple “urban” characteristics [6,44,78]; (ii) improvements in infrastructure and vector control mechanisms in urban centers, such as a regular water supply [6,62,127], which shift the disease burden further towards more rural settings without parallel improvements in infrastructure; and (iii) other changes to rural environments that may promote DENV transmission, including the increased introduction of plastics [9,44,49,119], the proliferation of new roads [6,25,106,110], and changes in land use [59,76,106,108]. These changing rural environments and evolving ways in which humans interact with their environments may contribute to the differential DENV transmission across various rural and urban settings.

### Objective 2: Compare dengue risk in rural and urban areas

We found that the risk of DENV transmission in rural areas is often as high or higher than the risk in urban areas (Fig 3). While many papers included in this review suggested that rural dengue is an emerging challenge, our review also illustrates the complexity of dengue in rural regions, showing that dengue transmission is not best explained by a rural versus urban dichotomy. As rural transmission increases, the articles included in our review suggest that ongoing urbanization may create a feedback loop where new land and network development may bring rural sectors in closer proximity to urban sectors and increase movement of humans, goods, and services between urban and rural areas. The role of urbanization on dengue transmission in our review has specifically been attributed to regional development, such as the construction of new roads [6,25] and increasing transportation systems [6,7,49,53,69,78,135], which may allow importation of infected individuals [121] and the expansion of population densities or sizes [6,9,25,58,100]. This has also resulted in communities encountering new challenges that may potentiate DENV transmission, such as inadequate waste management systems [9,119] and changing water storage practices [6,9,76]. Other changes, such as to housing infrastructure [119] or land use [58,59,68,76,108], may contribute to increasing vector densities [76] through further changes to breeding sites and human exposure to the environment. These factors may create environmental patches of high and low risk condition that cross the politically defined rural and urban sectors [21].

### Objective 3: Assess dengue burden in rural areas

While our results are insufficient to confirm that incidence is increasing in rural areas, several other lines of evidence from this review also support this conclusion. Four studies that stratified rural seroprevalence by age show increasing seroprevalence among children over time [57,109,119] (S1 Fig). This implies that the average age of first infection may be decreasing and

rural incidence may be increasing over time. Some studies suggested an increasing incidence over time; however, comparison across regions was difficult as DENV transmission can cycle and overall patterns may vary depending on the time scale used. Generally, longer time scales would be less likely to be susceptible to this bias. Thus, future studies may obtain more reliable estimates when using longer follow-up studies.

Additionally, the incidence of dengue we observe here may be related to a true increase or result from other factors. For example, improved surveillance systems may lead to improved case reporting, even as incidence remains unchanged. We find that relatively few studies provided sufficient information to distinguish between these 2 possibilities. More detailed laboratory data that combine seroprevalence and prospective incidence data from cohort studies could help disentangle these 2 mechanisms. The methods used in case ascertainment may have a large impact on an observed association with dengue in the literature. For example, many articles captured by our review come from passive hospital-based surveillance systems [4,6,9,49,53,61,78,135], which may disproportionately underestimate rural cases. These systems fail to identify people who do not seek healthcare and those with milder presentations [44,78], which may be amplified by transportation challenges in more remote or poorer areas. As expected, the overall incidence of dengue for the articles in our review was generally lower among studies using passive surveillance systems compared with studies with more active case finding methods (S2 and S3 Figs).

Other factors contributing to inconsistent DENV rural estimates may include study size, varied definitions of rurality, and local differences in mechanisms. These inconsistent estimates may be related to the sample sizes and aggregation used in many studies. Many studies had small sample sizes, making their quantitative estimates less reliable. Moreover, several studies assessed dengue seroprevalence in large rural areas or combined rural estimates from several different villages, resulting in relatively large rural populations with variations in cases between specific sites [58,119]. While such studies might be more representative of rural populations in general, pooling communities in this way made it difficult for us to quantify the impact of individual community size on dengue risk.

While it seems unlikely that isolated rural areas captured by the articles in our review have a sufficient population to sustain endemic levels of dengue, some authors hypothesized that transportation networks among rural communities resulted in population dynamics that mimic urban centers [69]. Others have noted the role of a sylvatic cycle and exposure to natural environments on rural DENV transmission [58,59,68,69]. Areas with close contact to heavily vegetated areas and primates may create the opportunity for spillback sylvatic transmission and periodic reintroduction of arboviruses. While none of the articles reviewed found definitive evidence of sylvatic DENV transmission to humans [69,81], some have discussed how changes in rural land use may increase the potential for human exposure to the existing natural transmission cycles [58,59,68]. If rural dengue is independently sustained by larger networks of rural populations or through reintroduction by sylvatic cycles, it is possible that rural circulation may be sufficient to maintain a reservoir. When immunity in urban centers is high and growing transportation networks may facilitate the reintroduction of DENV into urban centers once population immunity declines.

## Limitations

The primary limitation of this review was the inconsistent quality of surveillance systems in the literature and a lack of longitudinal data, which made it difficult to assess changes in dengue burden over time. Secondly, while our inclusion criteria focused on studies in rural sites including urban areas only if they were compared directly with rural sites by the same research

group, this strategy a large proportion of available incidence data available from urban sites, potentially limiting our ability to compare incidence in urban and rural areas more generally. Moreover, given that there is no international consensus on what is considered “rural” or “urban,” we used regional definitions, as recommended by the UN [38]. Thus, a study in one region reporting rural dengue might have been classified as urban in another context. We are hopeful that the detailed data captured in this review can help the field move towards standardized definitions in future studies.

## Conclusions and public health implications

There is a pressing need to better understand the changing and potentially increasing burden of DENV transmission in rural areas. While population size may be related to dengue risk, our results suggest that understanding the environmental and infrastructure features that drive risk may be just as important—if not more important—than absolute population size. As population size continues to increase in rural areas, our review suggests that infrastructural improvements are needed to minimize population vulnerability. For example, reliable water systems and safe waste disposal might help mitigate concerns about growing dengue incidence by limiting breeding sites. The increases in transportation networks may connect smaller rural regions, increasing the effective population size. Quality surveillance systems to track dengue incidence in rural areas can help identify new surges in population risk that might accompany changing infrastructure such as transportation. In the absence of such interventions, the often underresourced rural regions may play a major role in the spread of dengue infection, serving as sources of new infections as well as sinks to introduced from other regions without being noticed. Because these same infrastructure features change with ongoing urbanization, a feedback loop may be created that drives incidence to continue to increase in rural areas. Better identifying key features of dengue risk in rural locations will promote surveillance and support the development of intervention strategies that are most appropriate in this context.

## Supporting information

**S1 Fig. Rural dengue seroprevalence (%) in children over time.** Each association is from a different study.

(DOCX)

**S2 Fig. Yearly incidence per 100,000 from studies that used passive methods to collect rural dengue data.**

(DOCX)

**S3 Fig. Yearly incidence per 100,000 from studies that used active methods to collect rural dengue data.**

(DOCX)

**S1 Table. Adapted Newcastle–Ottawa Scale (NOS) criteria for systematic reviews.**

(DOCX)

**S1 Text. Supplementary text.**

(DOCX)

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## References

1. Shepard DS, Undurraga EA, Halasa YA, Stanaway JD. The global economic burden of dengue: a systematic analysis. *Lancet Infect Dis.* 2016; 16(8):935–941. [https://doi.org/10.1016/S1473-3099\(16\)00146-8](https://doi.org/10.1016/S1473-3099(16)00146-8) PMID: 27091092
2. Alves JA, Santos JR, de Mendonça EN, Abud AFC, Nunes MDS, Fakhouri R, et al. Epidemiological aspects of dengue in Aracaju, State of Sergipe, Brazil. *Rev Soc Bras Med Trop.* 2011; 44(6):670–673. <https://doi.org/10.1590/s0037-86822011000600004> PMID: 22231240
3. Bhatt S, Gething PW, Brady OJ, Messina JP, Farlow AW, Moyes CL, et al. The global distribution and burden of dengue. *Nature.* 2013; 496(7446):504–507. <https://doi.org/10.1038/nature12060> PMID: 23563266
4. Raghwani J, Rambaut A, Holmes EC, Hang VT, Hien TT, Farrar J, et al. Endemic dengue associated with the co-circulation of multiple viral lineages and localized density-dependent transmission. *PLoS Pathog.* 2011; 7(6):e1002064. <https://doi.org/10.1371/journal.ppat.1002064> PMID: 21655108
5. Morales-Pérez A, Nava-Aguilera E, Legorreta-Soberanis J, Cortés-Guzmán AJ, Balanzar-Martínez A, Harris E, et al. "Where we put little fish in the water there are no mosquitoes": a cross-sectional study on biological control of the Aedes aegypti vector in 90 coastal-region communities of Guerrero, Mexico. *BMC Public Health.* 2017; 17(Suppl 1):433. <https://doi.org/10.1186/s12889-017-4302-z> PMID: 28699557
6. Vong S, Khieu V, Glass O, Ly S, Duong V, Huy R, et al. Dengue incidence in urban and rural Cambodia: results from population-based active fever surveillance, 2006–2008. *PLoS Negl Trop Dis.* 2010; 4(11):e903. <https://doi.org/10.1371/journal.pntd.0000903> PMID: 21152061
7. Nava-Aguilera E, Morales-Pérez A, Balanzar-Martínez A, Rodríguez-Ramírez O, Jiménez-Alejo A, Flores-Moreno M, et al. Dengue occurrence relations and serology: cross-sectional analysis of results from the Guerrero State, Mexico, baseline for a cluster-randomised controlled trial of community mobilisation for dengue prevention. *BMC Public Health.* 2017; 17(Suppl 1):435. <https://doi.org/10.1186/s12889-017-4291-y> PMID: 28699560
8. Tozan Y, Ratanawong P, Sewe MO, Wilder-Smith A, Kittayapong P. Household costs of hospitalized dengue illness in semi-rural Thailand. *PLoS Negl Trop Dis.* 2017; 11(9):e0005961. <https://doi.org/10.1371/journal.pntd.0005961> PMID: 28937986
9. Stevens AM, Carter K, Stevenson K, Schneeweiss R. The epidemiology of dengue fever in Palau. *Pac Health Dialog.* 2011; 17(1):119–128. PMID: 23008976
10. Balmaseda A, Standish K, Mercado JC, Matute JC, Tellez T, Saborío S, et al. Trends in patterns of dengue transmission over 4 years in a pediatric cohort study in Nicaragua. *J Infect Dis.* 2010; 201(1):5–14. <https://doi.org/10.1086/648592> PMID: 19929380
11. Chau TN, Anders KL, Lien LB, Hung NT, Hieu LTM, Tuan NM, et al. Clinical and virological features of Dengue in Vietnamese infants. *PLoS Negl Trop Dis.* 2010; 4(4):e657. <https://doi.org/10.1371/journal.pntd.0000657> PMID: 20405057
12. Gubler DJ. The global emergence/resurgence of arboviral diseases as public health problems. *Arch Med Res.* 2002; 33(4):330–342. [https://doi.org/10.1016/s0188-4409\(02\)00378-8](https://doi.org/10.1016/s0188-4409(02)00378-8) PMID: 12234522
13. Gubler DJ. Cities spawn epidemic dengue viruses. *Nat Med.* 2004; 10(2):129–130. <https://doi.org/10.1038/nm0204-129> PMID: 14760418

14. Guzman MG, Alvarez A, Vazquez S, Alvarez M, Rosario D, Pelaez O, et al. Epidemiological studies on dengue virus type 3 in Playa municipality, Havana, Cuba, 2001–2002. *Int J Infect Dis.* 2012; 16(3): e198–e203. <https://doi.org/10.1016/j.ijid.2011.09.026> PMID: 22277259
15. Kuan G, Gordon A, Avilés W, Ortega O, Hammond SM, Elizondo D, et al. The Nicaraguan pediatric dengue cohort study: study design, methods, use of information technology, and extension to other infectious diseases. *Am J Epidemiol.* 2009; 170(1):120–129. <https://doi.org/10.1093/aje/kwp092> PMID: 19435864
16. Reyes M, Mercado JC, Standish K, Matute JC, Ortega O, Moraga B, et al. Index cluster study of dengue virus infection in Nicaragua. *Am J Trop Med Hyg.* 2010; 83(3):683–689. <https://doi.org/10.4269/ajtmh.2010.10-0023> PMID: 20810839
17. Schmidt WP, Suzuki M, Thiem VD, White RG, Tsuzuki A, Yoshida LM, et al. Population density, water supply, and the risk of dengue fever in Vietnam: cohort study and spatial analysis. *PLoS Med.* 2011; 8(8):e1001082. <https://doi.org/10.1371/journal.pmed.1001082> PMID: 21918642
18. Cuong HQ, Vu NT, Cazelles B, Boni MF, Thai KTD, Rabaa MA, et al. Spatiotemporal dynamics of dengue epidemics, southern Vietnam. *Emerg Infect Dis.* 2013; 19(6):945–953. <https://doi.org/10.3201/eid1906.121323> PMID: 23735713
19. Rabaa MA, Klungthong C, Yoon IK, Holmes EC, Chinnawirotisan P, Thaisomboonsuk B, et al. Frequent in-migration and highly focal transmission of dengue viruses among children in Kamphaeng Phet, Thailand. *PLoS Negl Trop Dis.* 2013; 7(1):e1990. <https://doi.org/10.1371/journal.pntd.0001990> PMID: 23350000
20. Stoddard ST, Morrison AC, Vazquez-Prokopec GM, Soldan VP, Kochel TJ, Kitron U, et al. The role of human movement in the transmission of vector-borne pathogens. *PLoS Negl Trop Dis.* 2009; 3(7): e481. <https://doi.org/10.1371/journal.pntd.0000481> PMID: 19621090
21. Kraay ANM, Trostle J, Brouwer AF, Cevallos W, Eisenberg JNS. Determinants of Short-term Movement in a Developing Region and Implications for Disease Transmission. *Epidemiology.* 2018; 29(1):117–125. <https://doi.org/10.1097/EDE.0000000000000751> PMID: 28901976
22. Perez-Castro R, Castellanos JE, Olano VA, et al. Detection of all four dengue serotypes in *Aedes aegypti* female mosquitoes collected in a rural area in Colombia. *Mem Inst Oswaldo Cruz.* 2016; 111(4):233–240. <https://doi.org/10.1590/0074-02760150363> PMID: 27074252
23. Weaver SC, Vasilakis N. Molecular evolution of dengue viruses: contributions of phylogenetics to understanding the history and epidemiology of the preeminent arboviral disease. *Infect Genet Evol.* 2009; 9(4):523–540. <https://doi.org/10.1016/j.meegid.2009.02.003> PMID: 19460319
24. Vasilakis N, Cardosa J, Hanley KA, Holmes EC, Weaver SC. Fever from the forest: prospects for the continued emergence of sylvatic dengue virus and its impact on public health. *Nat Rev Microbiol.* 2011; 9(7):532–541. <https://doi.org/10.1038/nrmicro2595> PMID: 21666708
25. Shah PS, Deoshatwar A, Karad S, Mhaske S, Singh A, Bachal RV, et al. Seroprevalence of dengue in a rural and an urbanized village: A pilot study from rural western India. *J Vector Borne Dis.* 2017; 54(2):172–176. <https://doi.org/10.48883> PMID: 28748839
26. Braks MA, Honório NA, Lourenço-De-Oliveira R, Juliano SA, Lounibos LP. Convergent habitat segregation of *Aedes aegypti* and *Aedes albopictus* (Diptera: Culicidae) in southeastern Brazil and Florida. *J Med Entomol.* 2003; 40(6):785–794. <https://doi.org/10.1603/0022-2585-40.6.785> PMID: 14765654
27. Honório NA, Castro MG, Barros FSM, Magalhães MAFM, Sabroza PC. The spatial distribution of *Aedes aegypti* and *Aedes albopictus* in a transition zone, Rio de Janeiro, Brazil. *Cad Saude Publica.* 2009; 25(6):1203–1214. <https://doi.org/10.1590/s0102-311x200900600003> PMID: 19503951
28. Cox J, Grillet ME, Ramos OM, Amador M, Barrera R. Habitat segregation of dengue vectors along an urban environmental gradient. *Am J Trop Med Hyg.* 2007; 76(5):820–826. <https://doi.org/10.4216/ajtmh.17488898> PMID: 17488898
29. Murdock CC, Evans MV, McClanahan TD, Miazgowicz KL, Tesla B. Fine-scale variation in microclimate across an urban landscape shapes variation in mosquito population dynamics and the potential of *Aedes albopictus* to transmit arboviral disease. *PLoS Negl Trop Dis.* 2017; 11(5):e0005640. <https://doi.org/10.1371/journal.pntd.0005640> PMID: 28558030
30. Fournet F, Rican S, Vaillant Z, Roudot A, Meunier-Nikiema A, Kassié D, et al. The Influence of Urbanization Modes on the Spatial Circulation of Flaviviruses within Ouagadougou (Burkina Faso). *Int J Environ Res Public Health.* 2016; 13(12). <https://doi.org/10.3390/ijerph13121226> PMID: 27973402
31. Barmak DH, Dorso CO, Otero M, Solari HG. Dengue epidemics and human mobility. *Phys Rev E Stat Nonlin Soft Matter Phys.* 2011; 84(1 Pt 1):011901. <https://doi.org/10.1103/PhysRevE.84.011901> PMID: 21867207
32. Lana RM, Gomes M, Lima TFM, Honório NA, Codeço CT. The introduction of dengue follows transportation infrastructure changes in the state of Acre, Brazil: A network-based analysis. *PLoS Negl Trop Dis.* 2017; 11(11):e0006070. <https://doi.org/10.1371/journal.pntd.0006070> PMID: 29149175

33. Kraemer MU, Perkins TA, Cummings DA, Zakar R, Hay SI, Smith DL, et al. Big city, small world: density, contact rates, and transmission of dengue across Pakistan. *J R Soc Interface*. 2015; 12(111):20150468. <https://doi.org/10.1098/rsif.2015.0468> PMID: 26468065
34. Shamseer L, Moher D, Clarke M, Ghersi D, Liberati A, Petticrew M, et al. Preferred reporting items for systematic review and meta-analysis protocols (PRISMA-P) 2015: elaboration and explanation. *BMJ*. 2015; 350:g7647. <https://doi.org/10.1136/bmj.g7647> PMID: 25555855
35. Wells GA, Shea B, O'Connell D, Peterson J, Welch V, Losos M, et al. The Newcastle-Ottawa scale (NOS) for assessing the quality of nonrandomized studies in meta-analyses. [http://www.ohri.ca/programs/clinical\\_epidemiology/oxford.asp](http://www.ohri.ca/programs/clinical_epidemiology/oxford.asp). [cited 2018 May 21].
36. World Health Organization. Increasing access to health workers in remote and rural areas through improved retention. [http://apps.who.int/iris/bitstream/handle/10665/44369/9789241564014\\_eng.pdf;jsessionid=34485001B63A8D7318C4EF53DA82260B?sequence=1](http://apps.who.int/iris/bitstream/handle/10665/44369/9789241564014_eng.pdf;jsessionid=34485001B63A8D7318C4EF53DA82260B?sequence=1). [cited 2018 May 21].
37. United Nations. Principles and Recommendations for Population and Housing Censuses. 2008.
38. Centers for Disease Control and Prevention. NCHS Urban-Rural Classification Scheme for Counties. 2017.
39. Ahmad N, Khan T, Jamal SM. A Comprehensive Study of Dengue Epidemics and Persistence of Anti-Dengue Virus Antibodies in District Swat, Pakistan. *Intervirology*. 2020; 63(1–6):46–56. <https://doi.org/10.1159/000510347> PMID: 33049733
40. Aikat A, Sarkar A, Tudu N, Moitra S, Pal PP, Sarkar TK. Diagnostic dilemma of dengue fever: How much reliability are there in rapidity. *J Indian Med Assoc*. 2011; 109(8):543–545. PMID: 22315859
41. Aldstadt J, Yoon IK, Tannitisupawong D, Jarman RG, Thomas SJ, Gibbons RV, et al. Space-time analysis of hospitalised dengue patients in rural Thailand reveals important temporal intervals in the pattern of dengue virus transmission. *Trop Med Int Health*. 2012; 17(9):1076–1085. <https://doi.org/10.1111/j.1365-3156.2012.03040.x> PMID: 22808917
42. Arroyave E, Londono AF, Quintero JC, et al. Etiology and epidemiological characterization of non-malarial febrile syndrome in three municipalities of Urabá (Antioquia), Colombia. *Biomedica*. 2013; 33:99–107.
43. Azami NAM, Moi ML, Salleh SA, Neoh HM, Kamaruddin MA, Jalal NA, et al. Dengue epidemic in Malaysia: urban versus rural comparison of dengue immunoglobulin G seroprevalence among Malaysian adults aged 35–74 years. *Trans R Soc Trop Med Hyg*. 2020; 114(11):798–811. <https://doi.org/10.1093/trstmh/traa056> PMID: 32735681
44. Barde PV, Kori BK, Shukla MK, Bharti PK, Chand G, Kumar G, et al. Maiden outbreaks of dengue virus 1 genotype III in rural central India. *Epidemiol Infect*. 2015; 143(2):412–418. <https://doi.org/10.1017/S0950268814000612> PMID: 24667083
45. Bartley L, Carabin H, Chau NV, Ho V, Luxemburger C, Hien TT, et al. Assessment of the factors associated with flavivirus seroprevalence in a population in Southern Vietnam. *Epidemiol Infect*. 2002; 128(2):213–220. <https://doi.org/10.1017/s0950268801006495> PMID: 12002539
46. Bartoloni A, Remoli ME, Farchi F, Zammarchi L, Fortuna C, Benedetti E, et al. Seroprevalence of West Nile and dengue virus in the human population of the Bolivian Chaco. *J Med Virol*. 2019; 91(1):146–150. <https://doi.org/10.1002/jmv.25283> PMID: 30118540
47. Baruah HC, Mohapatra PK, Kire M, Pegu DK, Mahanta J. Haemorrhagic manifestations associated with dengue virus infection in Nagaland. *J Commun Dis*. 1996; 28(4):301–303. PMID: 9057456
48. Bhoomiboonchoo P, Gibbons RV, Huang A, Yoon IK, Buddhari D, Nisalak A, et al. The spatial dynamics of dengue virus in Kamphaeng Phet, Thailand. *PLoS Negl Trop Dis*. 2014; 8(9):e3138. <https://doi.org/10.1371/journal.pntd.0003138> PMID: 25211127
49. Bodinayake CK, Tillekeratne LG, Nagahawatte A, Devasiri V, Arachichi WK, Strouse JJ, et al. Emergence of Epidemic Dengue-1 Virus in the Southern Province of Sri Lanka. *PLoS Negl Trop Dis*. 2016; 10(10):e0004995. <https://doi.org/10.1371/journal.pntd.0004995> PMID: 27711206
50. Buddhari D, Anderson KB, Gromowski GD, Jarman RG, Iamsirithaworn S, Thaisomboonsuk B, et al. Correlation between reported dengue illness history and seropositivity in rural Thailand. *PLoS Negl Trop Dis*. 2021; 15(6):e0009459. <https://doi.org/10.1371/journal.pntd.0009459> PMID: 34129599
51. Catenacci LS, Ferreira MS, Fernandes D, Padda H, Travassos-da-Rosa ES, Deem SL, et al. Individual, household and environmental factors associated with arboviruses in rural human populations, Brazil. *Zoonoses Public Health*. 2021; 68(3):203–212. <https://doi.org/10.1111/zph.12811> PMID: 33538403
52. Chaaithanya IK, Kulkarni R, Salvi N, Nahiri KA, Kantela NN, Patil S, et al. Sentinel Surveillance of Dengue Virus Infection in a Tribal District of Maharashtra, India. *J Assoc Physicians India*. 2021; 69(9):11–12.

53. Chareonsook O, Foy H, Teeraratkul A, Silarug N. Changing epidemiology of dengue hemorrhagic fever in Thailand. *Epidemiol Infect*. 1999; 122(1):161–166. <https://doi.org/10.1017/s0950268898001617> PMID: 10098800
54. Cheah W, Chang M, Wang Y. Spatial, environmental and entomological risk factors analysis on a rural dengue outbreak in Lundu District in Sarawak, Malaysia. *Trop Biomed*. 2006; 23(1):85–96. PMID: 17041556
55. Chen W, Chen S, Chien L, Chen CC, King CC, Harn MR, et al. Silent transmission of the dengue virus in southern Taiwan. *Am J Trop Med Hyg*. 1996; 55(1):12–16. <https://doi.org/10.4269/ajtmh.1996.55.12> PMID: 8702015
56. Cohen AL, Dowell SF, Nisalak A, Mammen MP Jr, Petkanchanapong W, Fisk TL. Rapid diagnostic tests for dengue and leptospirosis: antibody detection is insensitive at presentation. *Trop Med Int Health*. 2007; 12(1):47–51. <https://doi.org/10.1111/j.1365-3156.2006.01752.x> PMID: 17207147
57. Da Silva-Nunes M, De Souza VAF, Pannuti CS, Sperança MA, Terzian ACB, Nogueira ML, et al. Risk factors for dengue virus infection in rural amazonia: Population-based cross-sectional surveys. *Am J Trop Med Hyg*. 2008; 79(4):485–494. PMID: 18840734
58. Da Silva-Nunes M, Malafronte RDS, Luz BDA, de Souza EA, Martins LC, Rodrigues SG, et al. The Acre Project: The epidemiology of malaria and arthropod-borne virus infections in a rural Amazonian population. *Cad Saude Publica*. 2006; 22(6):1325–1334. <https://doi.org/10.1590/s0102-311x2006000600021> PMID: 16751971
59. Dhanoa A, Hassan SS, Jahan NK, Reidpath DD, Fatt QK, Ahmad MP, et al. Seroprevalence of dengue among healthy adults in a rural community in Southern Malaysia: a pilot study. *Infect Dis Poverty*. 2018; 7(1):1. <https://doi.org/10.1186/s40249-017-0384-1> PMID: 29335021
60. Dickstein Y, Neuberger A, Golus M, Schwartz E. Epidemiologic profile of patients seen in primary care clinics in an urban and a rural setting in Haiti, 2010–11. *Int Health*. 2014; 6(3):258–262. <https://doi.org/10.1093/inthealth/ihu033> PMID: 24969645
61. Ditsuwan T, Liabsuetrakul T, Ditsuwan V, Thammapalo S. Cost of standard indoor ultra-low-volume space spraying as a method to control adult dengue vectors. *Trop Med Int Health*. 2012; 17(6):767–774. <https://doi.org/10.1111/j.1365-3156.2012.02997.x> PMID: 22943302
62. Doum D, Overgaard HJ, Mayxay M, Suttiprapa S, Saichua P, Ekalaksananan T, et al. Dengue Sero-prevalence and Seroconversion in Urban and Rural Populations in Northeastern Thailand and Southern Laos. *Int J Environ Res Public Health*. 2020; 17(23). <https://doi.org/10.3390/ijerph17239134> PMID: 33297445
63. Dubot-Pérès A, Vongphrachanh P, Denny J, Phetsouvanh R, Linthavong S, Sengkeopraseuth B, et al. An epidemic of dengue-1 in a remote village in rural Laos. *PLoS Negl Trop Dis*. 2013; 7(8):e2360. <https://doi.org/10.1371/journal.pntd.0002360> PMID: 23951379
64. Eamchan P, Nisalak A, Foy H, Chareonsook O. Epidemiology and control of dengue virus infections in Thai villages in 1987. *Am J Trop Med Hyg*. 1989; 41(1):95–101. PMID: 2764233
65. Farfan-Ale JA, Lorono-Pino MA. Incidencia de infección por virus dengue en niños de 8 a 14 años de edad radicados en las áreas urbana y rural del municipio de Mérida, Yucatán. *Bol Med Hosp Infant Mex*. 1991; 48(11):780–784.
66. Gilani SA, Khan MA, Malik AA, Burq AI. Sero-positive rate of hepatitis-c viral infection in patients of communicable and non-communicable diseases in rural communities of Punjab Pakistan. *Medical Forum Monthly*. 2017; 28(6):21–23.
67. Goel MK, Malik JS, Khanna P, Gaur DR, Behera BK, Yadav RK. Sero—Epidemiological investigation of an outbreak of fever in rural area of Rohtak. *J Commun Dis*. 2010; 42(1):77–79. PMID: 22468557
68. Grossi-Soyster EN, Cook EAJ, de Glanville WA, Thomas LF, Krystosik AR, Lee J, et al. Serological and spatial analysis of alphavirus and flavivirus prevalence and risk factors in a rural community in western Kenya. *PLoS Negl Trop Dis*. 2017; 11(10):e0005998. <https://doi.org/10.1371/journal.pntd.0005998> PMID: 29040262
69. Hayes CG, Phillips IA, Callahan JD, Griebenow WF, Hyams KC, Wu SJ, et al. The epidemiology of dengue virus infection among urban, jungle, and rural populations in the Amazon region of Peru. *Am J Trop Med Hyg*. 1996; 55(4):459–463. <https://doi.org/10.4269/ajtmh.1996.55.459> PMID: 8916809
70. Hem S, Ly S, Votsi I, Vogt F, Asgari N, Buchy P, et al. Estimating the Burden of Leptospirosis among Febrile Subjects Aged below 20 Years in Kampong Cham Communities, Cambodia, 2007–2009. *PLoS ONE*. 2016; 11(4):e0151555. <https://doi.org/10.1371/journal.pone.0151555> PMID: 27043016
71. Hercik C, Cosmas L, Mogeni OD, Wamola N, Kohi W, Houpt E, et al. A Combined Syndromic Approach to Examine Viral, Bacterial, and Parasitic Agents among Febrile Patients: A Pilot Study in Kilombero, Tanzania. *Am J Trop Med Hyg*. 2018; 98(2):625–632. <https://doi.org/10.4269/ajtmh.17-0421> PMID: 29280432

72. Hu TS, Zhang HL, Feng Y, Fen JH, Tang T, Liu YH, et al. Epidemiological and molecular characteristics of emergent dengue virus in Yunnan Province near the China-Myanmar-Laos border, 2013–2015. *BMC Infect Dis.* 2017; 17(1):331. <https://doi.org/10.1186/s12879-017-2401-1> PMID: 28482813

73. Huy R, Wichmann O, Beatty M, Ngan C, Duong S, Margolis HS, et al. Cost of dengue and other febrile illnesses to households in rural Cambodia: a prospective community-based case-control study. *BMC Public Health.* 2009; 9:155. <https://doi.org/10.1186/1471-2458-9-155> PMID: 19473500

74. Ir P, Men C, Lucas H, Meessen B, Decoster K, Bloom G, et al. Self-reported serious illnesses in rural Cambodia: a cross-sectional survey. *PLoS ONE.* 2010; 5(6):e10930. <https://doi.org/10.1371/journal.pone.0010930> PMID: 20532180

75. Jing Q, Li Y, Liu J, Jiang L, Chen Z, Su W, et al. Dengue Underestimation in Guangzhou, China: Evidence of Seroprevalence in Communities With No Reported Cases Before a Large Outbreak in 2014. *Open Forum Infect Dis.* 2019; 6(7). <https://doi.org/10.1093/ofid/ofz256> PMID: 31304186

76. Joshi R, Mishra PK, Joshi D, Santhosh JR, Parida MM, Desikan P, et al. Clinical presentation, etiology, and survival in adult acute encephalitis syndrome in rural Central India. *Clin Neurol Neurosurg.* 2013; 115(9):1753–1761. <https://doi.org/10.1016/j.clineuro.2013.04.008> PMID: 23643180

77. Kartini K, Sofia S, Nasrullah N. Development of Smartphone-Based Early Alerts and Mosquito Monitoring System and Geographic Instrument System Applications. *Maced J Med Sci.* 2021; 9. <https://doi.org/10.3889/oamjms.2021.6127>

78. Kay B, Nam V, Tien T, Yen NT, Phong TV, Diep VTB, et al. Control of aedes vectors of dengue in three provinces of Vietnam by use of Mesocyclops (Copepoda) and community-based methods validated by entomologic, clinical, and serological surveillance. *Am J Trop Med Hyg.* 2002; 66(1):40–48. <https://doi.org/10.4269/ajtmh.2002.66.40> PMID: 12135266

79. Kraemer MU, Sinka ME, Duda KA, Mylne AQN, Shearer FM, Barker CM, et al. The global distribution of the arbovirus vectors *Aedes aegypti* and *Ae. albopictus*. *Elife.* 2015; 4:e08347. <https://doi.org/10.7554/elife.08347> PMID: 26126267

80. Kumar A, Sharma SK, Padbidri VS, Thakare JP, Jain DC, Datta KK. An outbreak of dengue fever in rural areas of northern India. *J Commun Dis.* 2001; 33(4):274–281. PMID: 12561505

81. Kuniholm M, Wolfe N, Huang C, Mpoudi-Ngole E, Tamoufe U, LeBreton M, et al. Seroprevalence and distribution of Flaviviridae, Togaviridae, and Bunyaviridae arboviral infections in rural Cameroonian adults. *Am J Trop Med Hyg.* 2006; 74(6):1078–1083. PMID: 16760524

82. Le Viet T, Choisy M, Bryant JE, Trong DV, Quang TP, Horby P, et al. A dengue outbreak on a floating village at Cat Ba Island in Vietnam. *BMC Public Health.* 2015; 15:940. <https://doi.org/10.1186/s12889-015-2235-y> PMID: 26395076

83. Leal-Castellanos CB, Garcia-Suárez R, González-Figueroa E, Fuentes-Allen JL, Escobedo-de la Peñal J. Risk factors and the prevalence of leptospirosis infection in a rural community of Chiapas, Mexico. *Epidemiol Infect.* 2003; 131(3):1149–1156. <https://doi.org/10.1017/s0950268803001201> PMID: 14959783

84. Lee GO, Vasco MS, Zuniga-Moya JC, Van Engen A, Uruchima J, et al. A dengue outbreak in a rural community in Northern Coastal Ecuador: An analysis using unmanned aerial vehicle mapping. *PLoS Negl Trop Dis.* 2021; 15(9). <https://doi.org/10.1371/journal.pntd.0009679> PMID: 34570788

85. Li HM, Huang YK, Su YC, Kao CH. Increased risk of autoimmune diseases in dengue patients: A population-based cohort study. *J Infect.* 2018; 77(3):212–219. <https://doi.org/10.1016/j.jinf.2018.03.014> PMID: 29746944

86. Mai VQ, Mai TTX, Tam NLM, Nghia LT, Komada K, Murakami H. Prevalence and Risk Factors of Dengue Infection in Khanh Hoa Province, Viet Nam: A Stratified Cluster Sampling Survey. *J Epidemiol.* 2018; 28(12):488–497. <https://doi.org/10.2188/jea.JE20170090> PMID: 29780057

87. Marques-Toledo CA, Degener CM, Vinhal L, Coelho G, Meira W, Codeço CT, et al. Dengue prediction by the web: Tweets are a useful tool for estimating and forecasting Dengue at country and city level. *PLoS Negl Trop Dis.* 2017; 11(7):e0005729. <https://doi.org/10.1371/journal.pntd.0005729> PMID: 28719659

88. Masika MM, Korhonen EM, Smura T, Uusitalo R, Vapalahti K, Mwaengi D, et al. Detection of dengue virus type 2 of Indian origin in acute febrile patients in rural Kenya. *PLoS Negl Trop Dis.* 2020; 14(3): e0008099. <https://doi.org/10.1371/journal.pntd.0008099> PMID: 32126086

89. Mayxay M, Castonguay-Vanier J, Chansamouth V, Dubot-Pérès A, Paris DH, Phetsouvanh R, et al. Causes of non-malarial fever in Laos: a prospective study. *Lancet Glob Health.* 2013; 1(1):e46–e54. [https://doi.org/10.1016/S2214-109X\(13\)70008-1](https://doi.org/10.1016/S2214-109X(13)70008-1) PMID: 24748368

90. Mayxay M, Sengvilaipaseuth O, Chanthongthip A, Dubot-Pérès A, Rolain JM, Parola P, et al. Causes of Fever in Rural Southern Laos. *Am J Trop Med Hyg.* 2015; 93(3):517–520. <https://doi.org/10.4269/ajtmh.14-0772> PMID: 26149859

91. Mease LE, Coldren RL, Musila LA, Prosser T, Ogolla F, Ofula VO, et al. Seroprevalence and distribution of arboviral infections among rural Kenyan adults: a cross-sectional study. *Virol J.* 2011; 8:371. <https://doi.org/10.1186/1743-422X-8-371> PMID: 21794131
92. Mueller TC, Siv S, Khim N, Kim S, Fleischmann E, Arey F, et al. Acute undifferentiated febrile illness in rural Cambodia: a 3-year prospective observational study. *PLoS ONE.* 2014; 9(4):e95868. <https://doi.org/10.1371/journal.pone.0095868> PMID: 24755844
93. Murhekar MV, Kamaraj P, Kumar MS, Khan SA, Allam RR, Barde P, et al. Burden of dengue infection in India, 2017: a cross-sectional population based serosurvey. *Lancet Glob Health.* 2019; 7(8):e1065–e1073. [https://doi.org/10.1016/S2214-109X\(19\)30250-5](https://doi.org/10.1016/S2214-109X(19)30250-5) PMID: 31201130
94. Murray-Smith S, Weinstein P, Skelly C. Field epidemiology of an outbreak of dengue fever in Charters Towers, Queensland: are insect screens protective? *Aust N Z J Public Health.* 1996; 20(5):545–547. <https://doi.org/10.1111/j.1467-842x.1996.tb01637.x> PMID: 8987228
95. Mutucumarana CP, Bodinayake CK, Nagahawatte A, Devasiri V, Kurukulasooriya R, Anuradha R, et al. Geospatial analysis of dengue emergence in rural areas in the Southern Province of Sri Lanka. *Trans R Soc Trop Med Hyg.* 2020; 114(6):408–414. <https://doi.org/10.1093/trstmh/trz123> PMID: 31885050
96. Nagao Y, Svasti P, Tawatsin A, Thavara U. Geographical structure of dengue transmission and its determinants in Thailand. *Epidemiol Infect.* 2008; 136(6):843–851. <https://doi.org/10.1017/S0950268807008990> PMID: 17624231
97. Nguyen PV, Vo TQ, Nguyen TD, Chung Phan TT, Ho Fan NV. Dengue fever in Southern of Vietnam: A survey of reported knowledge, attitudes, and practices. *J Pak Med Assoc.* 2019; 69(6). PMID: 31369542
98. Norman G, Theodre A, Joseph A. An insular outbreak of dengue fever in a rural south Indian village. *J Commun Dis.* 1991; 23(3):185–190. PMID: 1812164
99. Olson D, Lamb M, Lopez MR, Colborn K, Paniagua-Avila A, Zacarias A, et al. Performance of a Mobile Phone App-Based Participatory Syndromic Surveillance System for Acute Febrile Illness and Acute Gastroenteritis in Rural Guatemala. *J Med Internet Res.* 2017; 19(11):e368. <https://doi.org/10.2196/jmir.8041> PMID: 29122738
100. Overgaard HJ, Alexander N, Matiz MI, Jaramillo JF, Olano VA, Vargas S, et al. A Cluster-Randomized Controlled Trial to Reduce Diarrheal Disease and Dengue Entomological Risk Factors in Rural Primary Schools in Colombia. *PLoS Negl Trop Dis.* 2016; 10(11):e0005106. <https://doi.org/10.1371/journal.pntd.0005106> PMID: 27820821
101. Phanitchat T, Zhao B, Haque U, Pientong C, Ekalaksananan T, Aromseree S, et al. Spatial and temporal patterns of dengue incidence in northeastern Thailand 2006–2016. *BMC Infect Dis.* 2019; 19(1):743. <https://doi.org/10.1186/s12879-019-4379-3> PMID: 31443630
102. Phuong HL, de Vries PJ, Nagelkerke N, Giao PT, Hung LQ, Binh TQ, et al. Acute undifferentiated fever in Binh Thuan province, Vietnam: imprecise clinical diagnosis and irrational pharmaco-therapy. *Trop Med Int Health.* 2006; 11(6):869–879. <https://doi.org/10.1111/j.1365-3156.2006.01636.x> PMID: 16772009
103. Poblap T, Nitatpattana N, Chaimarin A, Barbazan P, Chauvancy G, Yoksan S, et al. Silent transmission of virus during a dengue epidemic, Nakhon Pathom Province, Thailand 2001. *Southeast Asian J Trop Med Public Health.* 2006; 37(5):899–903. PMID: 17333731
104. Raina S, Raina R, Agarwala N, Raina S, Sharma R. Coinfections as an aetiology of acute undifferentiated febrile illness among adult patients in the sub-Himalayan region of north India. *J Vector Borne Dis.* 2018; 55(2):130–136. <https://doi.org/10.4103/0972-9062.242560> PMID: 30280711
105. Reller ME, Bodinayake C, Nagahawatte A, Devasiri V, Kodikara-Arachchige W, Strouse JJ, et al. Unsuspected dengue and acute febrile illness in rural and semi-urban southern Sri Lanka. *Emerg Infect Dis.* 2012; 18(2):256–263. <https://doi.org/10.3201/eid1802.110962> PMID: 22304972
106. Riley-Powell AR, Lee GO, Naik NS, Jensen KE, O’Neal C, Salmón-Mulankovich G, et al. The Impact of Road Construction on Subjective Well-Being in Communities in Madre de Dios, Peru. *Int J Environ Res Public Health.* 2018; 15(6). <https://doi.org/10.3390/ijerph15061271> PMID: 29914119
107. Säll O, Thulin Hedberg S, Neander M, Tiwari S, Dornon L, Bom R, et al. Etiology of Central Nervous System Infections in a Rural Area of Nepal Using Molecular Approaches. *Am J Trop Med Hyg.* 2019; 101(1):253–259. <https://doi.org/10.4269/ajtmh.18-0434> PMID: 31162021
108. Samuel PP, Govindarajan R, Leo SVJ, Krishnamoorthy R, Paramasivan R, Arunachalam N. Emergence of dengue in the tribal pockets of Nilgiris district, Tamil Nadu. *Trop Doct.* 2017; 47(2):170–173. <https://doi.org/10.1177/0049475516677158> PMID: 27838633
109. Shah PS, Alagarasu K, Karad S, Deoshatwar A, Jadhav SM, Raut T, et al. Seroprevalence and incidence of primary dengue infections among children in a rural region of Maharashtra, Western India. *BMC Infect Dis.* 2019; 19(1):296. <https://doi.org/10.1186/s12879-019-3937-z> PMID: 30940086

110. Sharma KD, Mahabir RS, Curtin KM, Sutherland JM, Agard JB, Chadee DD. Exploratory space-time analysis of dengue incidence in Trinidad: A retrospective study using travel hubs as dispersal points, 1998–2004. *Parasit Vectors*. 2014; 7(1). <https://doi.org/10.1186/1756-3305-7-341> PMID: 25052242
111. Shukla MK, Singh N, Sharma RK, Barde PV. Utility of dengue NS1 antigen rapid diagnostic test for use in difficult to reach areas and its comparison with dengue NS1 ELISA and qRT-PCR. *J Med Virol*. 2017; 89(7):1146–1150. <https://doi.org/10.1002/jmv.24764> PMID: 28042883
112. Silarug N, Foy HM, Kupradinon S, Rojanasuphot S, Nisalak A, Pongsuwan Y. Epidemic of fever of unknown origin in rural Thailand, caused by influenza A (H1N1) and dengue fever. *Southeast Asian J Trop Med Public Health*. 1990; 21(1):61–67. PMID: 2169651
113. Sippy R, Herrera D, Gaus D, Gangnon RE, Patz JA, Osorio JE. Seasonal patterns of dengue fever in rural Ecuador: 2009–2016. *PLoS Negl Trop Dis*. 2019; 13(5):e0007360. <https://doi.org/10.1371/journal.pntd.0007360> PMID: 31059505
114. Smith C. The distribution of antibodies to Japanese encephalitis, dengue, and yellow fever viruses in five rural communities in Malaya. *Trans R Soc Trop Med Hyg*. 1958; 52(3):237–252. [https://doi.org/10.1016/0035-9203\(58\)90083-x](https://doi.org/10.1016/0035-9203(58)90083-x) PMID: 13556886
115. Sohan L, Shyamal B, Kumar TS, Malini M, Ravi K, Venkatesh V et al. Studies on leptospirosis outbreaks in Peddamandem Mandal of Chittoor District, Andhra Pradesh. *J Commun Dis*. 2008; 40(2):127–132. PMID: 19301697
116. Sokhna C, Mediannikov O, Fenollar F, Bassene H, Ditta G, Tall A, et al. Point-of-care laboratory of pathogen diagnosis in rural Senegal. *PLoS Negl Trop Dis*. 2013; 7(1):e1999. <https://doi.org/10.1371/journal.pntd.0001999> PMID: 23350001
117. Spicer PE, Taufa T, Benjamin AL. Scrub typhus (*Orientia tsutsugamushi*), spotted fever (*Rickettsia australis*) and dengue fever as possible causes of mysterious deaths in the Strickland Gorge area of Southern Highlands and West Sepik Provinces of Papua New Guinea. *P N G Med J*. 2007; 50(3–4):172–183. PMID: 19583101
118. Srey VH, Sadones H, Ong S, Mam M, Yim C, Sor S, et al. Etiology of encephalitis syndrome among hospitalized children and adults in Takeo, Cambodia, 1999–2000. *Am J Trop Med Hyg*. 2002; 66(2):200–207. <https://doi.org/10.4269/ajtmh.2002.66.200> PMID: 12135294
119. Strickman D, Sithiprasasna R, Kittayapong P, Innis B. Distribution of dengue and Japanese encephalitis among children in rural and suburban Thai villages. *Am J Trop Med Hyg*. 2000; 63(1–2):27–35. <https://doi.org/10.4269/ajtmh.2000.63.27> PMID: 11357991
120. Sun J, Lu L, Wu H, Yang J, Xu L, Sang S, et al. Epidemiological trends of dengue in mainland China, 2005–2015. *Int J Infect Dis*. 2017; 57:86–91. <https://doi.org/10.1016/j.ijid.2017.02.007> PMID: 28214563
121. Sun W, Xue L, Xie X. Spatial-temporal distribution of dengue and climate characteristics for two clusters in Sri Lanka from 2012 to 2016. *Sci Rep*. 2017; 7(1):12884. <https://doi.org/10.1038/s41598-017-13163-z> PMID: 29018222
122. Swoboda P, Fuehrer HP, Ley B, Starzengruber P, Ley-Threimer K, Jung M, et al. Evidence of a major reservoir of non-malarial febrile diseases in malaria-endemic regions of Bangladesh. *Am J Trop Med Hyg*. 2014; 90(2):377–382. <https://doi.org/10.4269/ajtmh.13-0487> PMID: 24420774
123. Tipayamongkholgu M, Lisakulruk S. Socio-geographical factors in vulnerability to dengue in Thai villages: A spatial regression analysis. *Geospat Health*. 2011; 5(2):191–198. <https://doi.org/10.4081/gh.2011.171> PMID: 21590669
124. Van Damme W, Van Leemput L, Por I, Hardeman W, Meessen B. Out-of-pocket health expenditure and debt in poor households: evidence from Cambodia. *Trop Med Int Health*. 2004; 9(2):273–280. <https://doi.org/10.1046/j.1365-3156.2003.01194.x> PMID: 15040566
125. Vanitha SS, Angadi N. A Cross Sectional Study on Social Profile of Dengue among Children in Davangere. *Indian J Public Health Res Dev*. 2018; 9(2).
126. Vannavong N, Seidu R, Stenström TA, Dada N, Overgaard HJ. Dengue-like illness surveillance: a two-year longitudinal survey in suburban and rural communities in the Lao People's Democratic Republic and in Thailand. *Western Pac Surveill Response J*. 2019; 10(1):15–24. <https://doi.org/10.5365/wpsar.2017.8.4.001> PMID: 31110838
127. Vedpathak VL, Soundale SG, Lakde RN, Deo DS, Khadilkar HA. Clinico-epidemiological profile of an outbreak of dengue fever in rural area of Ambajogai Mandai, district Beed. *J Commun Dis*. 2011; 43(1):69–72. PMID: 23785885
128. Victor TJ, Malathi M, Gurusamy D, Desai A, Ravi V, Narayanasamy G, et al. Dengue fever outbreaks in two villages of Dharmapuri district in Tamil Nadu. *Indian J Med Res*. 2002; 116:133–139. PMID: 12674826
129. Yoon IK. Focal Dengue Virus Transmission In Kamphaeng Phet, Thailand And Implications For Management. *Southeast Asian J Trop Med Public Health* 2015; 46 Suppl 1:17–25. PMID: 26506728

130. Yoon IK, Getis A, Aldstadt J, Rothman AL, Tannitisupawong D, Koenraadt CJM, et al. Fine scale spatiotemporal clustering of dengue virus transmission in children and *Aedes aegypti* in rural Thai villages. *PLoS Negl Trop Dis.* 2012; 6(7):e1730. <https://doi.org/10.1371/journal.pntd.0001730> PMID: 22816001
131. Yoon IK, Rothman AL, Tannitisupawong D, Srikiatkachorn A, Jarman RG, Aldstadt J, et al. Underrecognized mildly symptomatic viremic dengue virus infections in rural Thai schools and villages. *J Infect Dis.* 2012; 206(3):389–398. <https://doi.org/10.1093/infdis/jis357> PMID: 22615312
132. Yoon IK, Srikiatkachorn A, Hermann L, Buddhari D, Scott TW, Jarman RG, et al. Characteristics of mild dengue virus infection in Thai children. *Am J Trop Med Hyg.* 2013; 89(6):1081–1087. <https://doi.org/10.4299/ajtmh.13-0424> PMID: 24127167
133. Zafar H, Hayyat A, Akhtar N, Rizwan S. Prevalence of undifferentiated fever in adults of Rawalpindi having primary dengue fever. *J Pak Med Assoc.* 2013; 63(6):770–771. PMID: 23901683
134. Zafar S, Shipin O, Paul RE, Rocklöv J, Haque U, Rahman MS, et al. Development and Comparison of Dengue Vulnerability Indices Using GIS-Based Multi-Criteria Decision Analysis in Lao PDR and Thailand. *Int J Environ Res Public Health.* 2021; 18(17). <https://doi.org/10.3390/ijerph18179421> PMID: 34502007
135. Zala DB, Khan V, Kakadiya M, Sanghai AA, Das VK. Circulation of dengue serotypes in the Union Territory of Dadra & Nagar Haveli (India). *Parasite Epidemiol Control.* 2018; 3(3):e00069. <https://doi.org/10.1016/j.PAREPI.2018.e00069> PMID: 29988334