Occurrence of Japanese Encephalitis Virus Mosquito Vectors in Relation to Urban Pig Holdings

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Abstract. Japanese encephalitis virus (JEV) is transmitted to humans from pigs or birds by mosquitoes. In this study, the association between urban pig keeping and mosquito vectors was analyzed. A total of 7,419 mosquitoes were collected overnight in urban households with and without pigs in Can Tho City, Vietnam. The most prevalent vectors were *Culex tritaeniorhynchus* (36%), *Cx. gelidus* (24%), and *Cx. quinquefasciatus* (15%), which were present in all parts of the city. Pigs were associated with increased numbers of *Cx. tritaeniorhynchus*. Traps close to pigs had higher numbers of *Cx. tritaeniorhynchus* and *Cx. gelidus* than traps close to humans. Increased number of persons in the household was associated with increased numbers of *Cx. quinquefasciatus*. We demonstrate that JEV vector species are present at urban households with and without pigs, and show that keeping pigs in an urban area increase the number of mosquitoes competent as vectors for JEV.

INTRODUCTION

Japanese encephalitis (JE) is a zoonotic disease caused by Japanese encephalitis virus (JEV), a flavivirus that is transmitted by mosquitoes, particularly *Culex* species. Wading birds, such as herons and egrets, are reservoir hosts for the virus. Pigs serve as amplifying hosts, and a high density of pigs in an area can be associated with increased number of cases of JE. In spite of the high viremia in swine, infection is usually asymptomatic, except for abortions and mummified fetuses in pregnant sows. Humans and horses are dead-end hosts for the virus but may occasionally develop encephalitis, with case fatality rates of up to 30%.

The disease is present in southern and eastern Asia, and shows a tendency to emerge into new areas. In Vietnam, the first human case of JE was recognized in the 1960s and it has since become a serious health issue. During 1986–1994, there were more than 18,000 human cases reported, and JEV is currently endemic to the southern parts of the country. In southern Vietnam, the area used for irrigated rice production is increasing along with the pig production, which are factors likely to contribute further to the emergence of JEV. The Mekong delta region in Vietnam, has extensive pig farming and rice production, and is one of the regions in the country with the most JE cases. In Can Tho City Province, 60% of the pigs on farms have been shown to be seropositive for JEV.

The transmission of vector-borne diseases is regulated by the host, the pathogen, and the biology of vectors and their vectorial capacity, which makes the epidemiology complex and also dependent on environmental factors, e.g., climate and anthropogenic influences, such as changes in agricultural practices, irrigation schemes, and urbanization. The most important vectors of JEV, such as *Culex tritaeniorhynchus*, commonly undergo larval development in rice fields in rural areas. Therefore, JE has been considered a rural disease, in contrast to the closely-related dengue fever virus, which is well known for its outbreaks in urban areas, and spread by an urban dwelling mosquito (*Aedes aegypti*). However, JE may also expand into densely populated areas because of rapid urbanization, coupled with growing demands for urban agriculture involving livestock. In the cities, poultry, pigs, or small ruminants are preferred. Thus, in such areas, the close proximity to animals may result in increasing risks for transmission of zoonotic diseases, including JE.

To date, the influence of population density and pig husbandry on JEV vector distribution in urban areas has not been extensively investigated, although it has been shown that JEV infections occur in humans in urban settings. Therefore, the aim of the present study was to assess the effect of urban pig rearing and other factors on the presence of JEV vectors at households in Can Tho City in southern Vietnam.

MATERIAL AND METHODS

Description of urban agriculture in study area. Can Tho City is the central province of the Mekong delta in southern Vietnam. Data on the population of humans and animals and the area of rice fields and fish ponds were collected by visiting the local veterinary authorities in the different districts in the province during autumn of 2008, and using official government statistics. The study site in Ninh Kieu District is the most urban part of Can Tho City Province, and has a human population of approximately 217,000, which corresponded to a population density of 7,500 persons/km² in 2008. The rest of the province is mainly rural, with population densities of 380–1,400 persons/km² in the districts. The pig population in the rural districts varied between 65 and 123 pigs/km². Although Ninh Kieu had a human population density five times as high as the second most densely populated district, it had 94 pigs/km². The mean number of sows per pig holding, i.e., a household with domestic pigs, in Ninh Kieu was 1.4 (range = 0–4.5). Including piglets and boars, the mean number was 11 pigs per pig holding (range = 4.7–24). There were only nine hectares of rice paddies/km² in the urban district, compared with the other districts, which had 35–205 hectares/km².

Ninh Kieu contains 13 wards. Seven of these wards had cattle or buffaloes, and the density of large ruminants was 0.6–29/km². Only two of the most central wards, Tan An and An Hoi, had no records of pigs, and the ward reporting the
most pigs, An Binh, had a density of 340 pigs/km², the lowest human population density, and a ratio of 0.21 pigs/inhabitant. Pig and human populations are shown in Figure 1 (ArcMap software, ESRI, Redlands, CA).

**Study design and mosquito collection and identification.**
Wards representing a transect of the city with human densities ranging from 1,598 to 31,447 persons/km² and pig densities ranging from 0 to 340 pigs/km² were selected. In these wards, households within the urban area with and without pigs, which enabled sampling and where mosquito traps could be affixed, were included.

Mosquito collection was conducted during three-week periods from the middle of February to the beginning of March and from the end of October to the middle of November in 2009. Collection of mosquitoes was planned to be conducted on a weekly basis with at least two repetitions during each sampling period. Because of circumstances, such as the household moving, or trap failures, all households could not be sampled at both sampling periods or sampling could not be repeated in two subsequent weeks. A total of 17 households in 10 wards were studied (Table 1 and Figure 1).

Data were collected on factors potentially important for presence of vectors, i.e., the number of persons in the household; the number of pigs, pets, poultry, and other livestock present; and the presence of rice fields or fish ponds on the property.

Mosquitoes were collected by using un-baited CDC mini light traps (Bioquip Products, Compton, CA) equipped with a 4-watt light bulb. At each household, two traps were operated between dusk and dawn (6:00 PM to 7:00 AM). One trap was placed within two meters of the entrance to the human dwelling, or alternatively within two meters of the bed, if the bed was outside or in the same building as the pigs. In households with pigs, the second trap was placed either above or next to the pen. In households without pigs, the traps were placed 2–10 meters apart, where traps could be affixed. The traps were placed irrespective of other pets and livestock.

Mosquitoes were killed by freezing and identified according to the keys of Reuben and others. Because of difficulties...
in distinguishing *Cx. vishnui* and *Cx. pseudovishnui* within the *Cx. vishnui* subgroup, only *Cx. tritaeniorhynchus* was identified to species.

In collections containing more than 300 mosquitoes, a random sample of 300 specimens was identified, and the remaining specimens were counted. This amount always comprised more than 10% of the specimens in all the collections, and was thus estimated to be a representative sample size. The proportions of identified species and blood-filled specimens were then applied to the total number of mosquitoes to provide an extrapolated estimate of the composition of species in the total collection.

**Statistical analyses.** Statistical analysis was performed by using SAS for Windows 9.2 (SAS Institute Inc., Cary, NC). To improve the normality of the data, the number of mosquitoes was logarithmically transformed (log (n + 0.01)). The number of pigs and people at the household level were treated as continuous variables, and close proximity to livestock, including poultry, and the presence of pets, were classified as two separate dichotomous variables. Univariable analyses for the association of rice fields or fish ponds with the number of mosquitoes were performed, and both variables were found to have similar associations. The two variables were merged into one variable, representing the presence of large water bodies adjacent to the household.

Data on ward level was calculated as densities. The density of persons, pigs, and large ruminants were calculated as individuals/km² and used as continuous independent variables in the analyses. The area of rice fields and fish ponds was combined and divided by the total area of the ward.

To determine potential risk factors the dependent variables, the total number of mosquitoes, the total extrapolated number of *Cx. tritaeniorhynchus*, all members of the *Cx. vishnui* subgroup, *Cx. gelidus*, *Cx. quinquefasciatus*, *Culex* males, and the proportion of blood-filled females per trap per night were analyzed in two multivariable effect models. Statistical analyses were performed for *Cx. tritaeniorhynchus* and the whole *Cx. vishnui* subgroup. The first set of models included only traps placed near human dwellings. A subset of models were also made that used pigs in the household as a categorical, present or not present, variable instead of being continuous. The second set of models included only households with pigs and also whether the collections were made close to pigs versus close to humans.

Mixed effects modeling was performed using the SAS Mixed procedure for all analyses, except for the proportion of blood-filled females, for which the Glimmix procedure was used. Mixed modeling was used to account for clustering of collections at the ward and household level. In both procedures, the household was a random effect within the ward, and sampling period was a random effect within the household. Satterthwaite denominator degrees of freedom were used. All ward and household variables were included and then backward elimination was used for factors with a *P* value < 0.05 were kept in the model. Factors with a confounding effect were only kept in the model if they had an effect of more than 25% on the coefficient β of another factor that continued to be significant. Interactions between the remaining factors were tested and discarded if the *P* value was > 0.05.

**Table 2**

<table>
<thead>
<tr>
<th>Species/ward</th>
<th>An Binh</th>
<th>An Hoa</th>
<th>An Khanh</th>
<th>An Nghiep</th>
<th>An Phu</th>
<th>Cai Khe</th>
<th>Hung Loi</th>
<th>Thoi Binh</th>
<th>Xuan Khanh</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Culex tritaeniorhynchus</em></td>
<td>278</td>
<td>75</td>
<td>4</td>
<td>548</td>
<td>5</td>
<td>72</td>
<td>124</td>
<td>179</td>
<td>110</td>
<td>344</td>
</tr>
<tr>
<td><em>Cx. vishnui</em> subgroup*</td>
<td>45</td>
<td>12</td>
<td>1</td>
<td>6</td>
<td>–</td>
<td>–</td>
<td>6</td>
<td>26</td>
<td>12</td>
<td>14</td>
</tr>
<tr>
<td><em>Culex gelidus</em></td>
<td>132</td>
<td>64</td>
<td>2</td>
<td>95</td>
<td>2</td>
<td>37</td>
<td>24</td>
<td>237</td>
<td>93</td>
<td>266</td>
</tr>
<tr>
<td><em>Culex quinquefasciatus</em></td>
<td>18</td>
<td>33</td>
<td>1</td>
<td>18</td>
<td>1</td>
<td>8</td>
<td>1</td>
<td>5</td>
<td>4</td>
<td>12</td>
</tr>
<tr>
<td><em>Culex fuscoccephala</em></td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>5</td>
</tr>
<tr>
<td><em>Lutzia</em> spp.</td>
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<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>1</td>
</tr>
<tr>
<td><em>Anopheles</em> spp.</td>
<td>63</td>
<td>7</td>
<td>1</td>
<td>85</td>
<td>1</td>
<td>4</td>
<td>28</td>
<td>53</td>
<td>16</td>
<td>36</td>
</tr>
<tr>
<td><em>Uranotaenia</em> spp.</td>
<td>22</td>
<td>1</td>
<td>–</td>
<td>13</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>4</td>
<td>1</td>
<td>–</td>
</tr>
<tr>
<td><em>Mansonia</em> spp.</td>
<td>20</td>
<td>3</td>
<td>1</td>
<td>20</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>56</td>
<td>3</td>
<td>48</td>
</tr>
<tr>
<td><em>Aedes</em> spp.</td>
<td>1</td>
<td>–</td>
<td>–</td>
<td>1</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>8</td>
<td>1</td>
<td>–</td>
</tr>
<tr>
<td><em>Culex</em> males</td>
<td>29</td>
<td>111</td>
<td>10</td>
<td>15</td>
<td>14</td>
<td>54</td>
<td>82</td>
<td>71</td>
<td>80</td>
<td>547</td>
</tr>
<tr>
<td><em>Anopheles</em> males</td>
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<td>–</td>
<td>1</td>
<td>2</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>1</td>
<td>2</td>
<td>8</td>
</tr>
<tr>
<td><em>Mansonia</em> males</td>
<td>5</td>
<td>–</td>
<td>–</td>
<td>1</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>6</td>
<td>7</td>
<td>26</td>
</tr>
<tr>
<td><em>Aedes</em> males</td>
<td>1</td>
<td>–</td>
<td>–</td>
<td>1</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>2</td>
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<tr>
<td>Unidentified</td>
<td>22</td>
<td>1</td>
<td>1,605†</td>
<td>1</td>
<td>15</td>
<td>6</td>
<td>20</td>
<td>877†</td>
<td>2,547</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>840</td>
<td>379</td>
<td>32</td>
<td>2,405</td>
<td>35</td>
<td>234</td>
<td>476</td>
<td>717</td>
<td>522</td>
<td>1,779</td>
</tr>
</tbody>
</table>

*Excluding *Culex tritaeniorhynchus*.†At household S in An Khanh and household I in Xuan Khanh, more mosquitoes were collected per trap than could be identified in the local laboratory. A sample of 300 specimens were identified per trap, and the proportion of each species identified were then applied to the entire number collected to achieve an estimate of the mosquito population.
RESULTS

Description of households. There were 4–60 persons living in the investigated households without pigs (median = 4 persons). In the households with pigs, there were 1–10 persons (median = 6 persons). The latter households kept 1–110 pigs (median = 15 pigs). One of the households with pigs had no sows and seven households kept one or two sows. Four households had 6–13 sows. There was only one household where the distance between the nearest pigs and living quarters for humans was more than 10 meters. Only two households, I and S, had cattle, and household S also had crocodiles. Households A, F, I, M, N, and S had fish ponds by the house and two of these households (I and S) also had rice fields.

Mosquito collections. A total of 7,419 mosquitoes were collected (Table 2). The most prevalent species, Cx. tritaeniorhynchus (36%), Cx. gelidus (24%) and Cx. quinquefasciatus (15%), were found in all wards. The distribution of the number of these species at different collection sites is shown in Figure 2. Of the 4,872 identified specimens, 1,282 were blood filled. The proportion of blood-filled females was highest for Cx. gelidus (n = 421, 37%) and Cx. tritaeniorhynchus (n = 609, 35%) and was lowest for Cx. quinquefasciatus (n = 41, 5%). Other Culex mosquitoes identified were Cx. fuscocephala and members of the Cx. vishnui subgroup. The remaining mosquitoes consisted of the genera Mansonia, Aedes, Anopheles, and Uranotaenia. These mosquitoes and the Culex males were not identified to species.

Analysis of risk factors associated with number of mosquitoes. Collections near human dwellings in households with and without pigs. There was no difference in the final models if the total number of pigs was used or if only the presence or non-presence of pigs were used. Therefore, only the results for the model with the number of pigs are presented (Table 3). For the total number of mosquitoes and the number of Cx. tritaeniorhynchus and all mosquitoes in the Cx. vishnui subgroup, pigs were the only risk factor with a significant effect (P = 0.021, 0.030, and 0.048, respectively). The number of Cx. gelidus collected was increased by the density of pigs in the ward (P = 0.016), but this effect was confounded by the presence of other livestock in the household and the presence of rice fields or fishponds. The number of Cx. quinquefasciatus was only significantly associated with the number of persons in the household (P = 0.035) and the density of large ruminants in the ward (P = 0.004). No factors were identified that significantly affected the number of Culex males collected or the proportion of blood-filled mosquitoes.

Collections in households with pigs. In the second analysis, the two collection sites in households with pigs were included (Table 4). The total numbers of pigs in the household and mosquito collections made close to pigs were associated with an increase in the total number of mosquitoes (P = 0.004 and P < 0.0001 respectively), as well as the number of Cx. tritaeniorhynchus (P = 0.024 and 0.0004, respectively) and all the mosquitoes in the Cx. vishnui subgroup (P = 0.035 and 0.0007, respectively). The number of Cx. gelidus was increased by collections close to pigs (P < 0.0001), number of persons in the household (P = 0.003, pig density (P = 0.010), and presence of other livestock (P = 0.010), and decreased by increased large ruminant density (P = 0.033), but again the associations were complex and confounded by presence of rice fields or fish ponds. None of the risk factors studied had

Figure 2. Mosquitoes collected in households with and without pigs in Ninh Kieu District, Can Tho City, Vietnam. Collections made close to humans are shown by thin boxes and collections made close to pigs are shown by thick boxes. A, total number of mosquitoes, B, Culex tritaeniorhynchus. C, Cx. gelidus. D, Cx. quinquefasciatus. Circles indicate outliers > 1.5 x the interquartile range (error bars) and stars indicate extreme outliers > 3 x the interquartile range.
any association with the number of Cx. quinquefasciatus, Culex males, and the proportion of blood-filled mosquitoes.

**DISCUSSION**

This study shows the association between JEV vector mosquitoes and the presence of pigs in a city. The presence of pigs per se and the number of pigs kept in the household were associated with an increase in the number of mosquitoes. Anthropophilic and zoophilic vector species were demonstrated to be associated differently with household risk factors.

Zoophilic Cx. tritaeniorhynchus was the most numerous mosquito species caught in the present study. The number of pigs per household was the only factor associated with a higher number of Cx. tritaeniorhynchus. This finding is in accordance with those of earlier studies, which have indicated that pigs are preferred feeding hosts. Rice fields were reported only in the peripheral wards of the study area, but Cx. tritaeniorhynchus could be found in all parts of the city. Our results indicate that standing water collections, which are abundant within the city even during the dry season, could also provide larval habitats also Cx. tritaeniorhynchus, as has been shown for other mosquito species. Moreover, Cx. tritaeniorhynchus may disperse hundreds of kilometers during one night, which enables mosquitoes to be found in households far away from their larval habitats. The entire Cx. vishnui subgroup and Cx. tritaeniorhynchus displayed corresponding associations with the analyzed risk factors. The Cx. vishnui subgroup is known as the most important vectors for JEV and the virus has been isolated from such a mosquito in the Can Tho City Province.

* Culex tritaeniorhynchus and Cx. gelidus comprised 60% of the mosquitoes collected. Cx. gelidus was associated with increasing overall pig densities in the ward and the presence of non-porcine livestock in the household. Because ruminants do not have clinical signs or amplify the virus, and could divert mosquitoes away from humans and pigs, they have been proposed as a zooprophylaxis. In a suburb of Bangkok, with population densities less than 3,000 persons/km², but with more rural characteristics than in our study, more than 90% of the mosquitoes were Cx. tritaeniorhynchus and Cx. gelidus. Gingrich and others isolated JEV from Cx. tritaeniorhynchus and Cx. gelidus collected in the same Bangkok suburb. Culex gelidus may breed in a variety of water collections, and it is therefore not surprising that rice fields and fish ponds did not have a direct effect on the abundance.

There was no association between the numbers of Cx. quinquefasciatus and pigs. However, the number of persons in the household was associated with a significant increase in the number of the mosquito, whereas the density of large ruminants had a negative association. Culex quinquefasciatus is known to be anthropophilic and the negative association with large ruminants could be result from the fact that large ruminant densities may be expected to be correlated to more rural, i.e., less urban, characteristics. Although anthropophilic Cx. quinquefasciatus commonly feed on humans, it may be opportunistic, and it has been demonstrated that the same individual mosquito may feed on pigs and humans. The potential of Cx. quinquefasciatus as a vector in urban areas

**Table 3**

Factors associated with collection of *Culex* mosquito vectors for Japanese encephalitis virus near human dwellings in Ninh Kieu, Can Tho City, Vietnam

<table>
<thead>
<tr>
<th>Risk factor</th>
<th>No. mosquitoes</th>
<th>Cx. vishnui subgroup</th>
<th>Cx. tritaeniorhynchus</th>
<th>Cx. gelidus</th>
<th>Cx. quinquefasciatus</th>
<th>Culex males</th>
<th>Proportion of blood-filled mosquitoes</th>
</tr>
</thead>
<tbody>
<tr>
<td>No. pigs in household</td>
<td>1.03†</td>
<td>1.05†</td>
<td></td>
<td>1.02†</td>
<td>1.05†</td>
<td></td>
<td></td>
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<tr>
<td>Pig density in ward</td>
<td></td>
<td></td>
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<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Rice field or fish pond</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Rice fields and fish ponds in ward</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Other livestock in household</td>
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<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td>Large ruminant density in ward</td>
<td>86.0</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>No. persons in household</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>1.13†</td>
<td></td>
</tr>
<tr>
<td>Population density in ward</td>
<td></td>
<td></td>
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<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Pets in household</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*Change in number of mosquitoes is the anti-logarithm of the coefficient β. Non-significant variables were confounders in the models.
† P < 0.05.
‡ P < 0.01.
§ P < 0.001.

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**Table 4**

Factors associated with number of *Culex* mosquito vectors for Japanese encephalitis virus collected at households keeping pigs in Ninh Kieu, Can Tho City, Vietnam

<table>
<thead>
<tr>
<th>Risk factor</th>
<th>No. mosquitoes</th>
<th>Cx. vishnui subgroup</th>
<th>Cx. tritaeniorhynchus</th>
<th>Cx. gelidus</th>
<th>Cx. quinquefasciatus</th>
<th>Culex males</th>
<th>Proportion of blood-filled mosquitoes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Catch near pigs</td>
<td>3.89†</td>
<td>7.77†</td>
<td>8.96†</td>
<td>17.1†</td>
<td></td>
<td></td>
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</tr>
<tr>
<td>No. pigs in household</td>
<td>1.04†</td>
<td>1.05§</td>
<td>1.05§</td>
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<tr>
<td>Pig density in ward</td>
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<tr>
<td>Rice field or fish pond</td>
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<td>Rice fields and fish ponds in ward</td>
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<tr>
<td>Other livestock in household</td>
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<td></td>
</tr>
<tr>
<td>Large ruminant density in ward</td>
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<tr>
<td>No. persons in household</td>
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<tr>
<td>Population density in ward</td>
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<td>1.53‡</td>
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</tr>
<tr>
<td>Pets in household</td>
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*Change in number of mosquitoes is the anti-logarithm of the coefficient β.
† P < 0.001.
‡ P < 0.01.
§ P < 0.05.
is further indicated by its typical breeding underground in sewers and drains and because of the fact that JEV previously has been isolated from *Cx. quinquefasciatus* in southern Vietnam.11

There were a higher number of blood-filled mosquitoes collected near pigs, in accordance with previous findings.47 However, there was no significant difference in the proportion of blood-filled mosquitoes dependent on the locality where the traps were operated. The total number of mosquitoes collected in the traps varied considerably for unknown reasons. Reisen and others recommended light traps for standardized collection of JEV vectors because of the phototactic behavior of *Cx. tritaeniorhynchus*, but multiple conflicting sources of light in a city may interfere, resulting in an increased variation of mosquitoes collected.

It is estimated that 1.9 billion persons live in rural areas where JEV is transmitted, but the number of persons considered at risk would be increased if JEV would also be considered an urban disease. Today, more than half of the world’s urban population live in Asia, and more than 60% of urban inhabitants are estimated to be poor.50,51 In low-income countries with poor infrastructure, the increasing problem of providing food for an expanding urban population at affordable prices creates a need for urban animal keeping. The urban areas of Can Tho City have a high pig density, and the presence of sows in the households indicates that there is a more permanent urban farming practice and not only a temporary keeping of pigs for slaughter.

To further elucidate the risk for human health associated to urban JEV vectors it would be important to determine the mosquitoes urban larval habitats, feeding habits, possible seasonal differences and infection rates. For instance, blood meal analysis may give some indications to which extent different mosquito species are feeding on pigs and humans, although most mosquitoes are opportunistic and may feed on hosts depending on their availability.53,54 When arboviruses circulate in mosquitoes, it is usually only a small proportion of the vectors that are infected. In a previous study in Can Tho City Province, it was possible to find one JEV-positive mosquito pool in 22,000 mosquitoes collected.37 No virus detection was performed in this study, but we showed that all components necessary for transmission of JEV can be present in a tropic urban environment because of the presence of pigs and zoophilic and anthropophilic vectors. Our results indicate that the anthropophilic *Cx. quinquefasciatus* may become more important for the transmission of JEV in areas with fewer pigs and more humans as an urban bridge vector between animals and humans, as hypothesized by Do and others.11 The recent detection of JEV RNA in *Cx. pipiens*, a sibling species of *Cx. quinquefasciatus*, in Europe also implies the importance of the *Cx. pipiens* complex, not only for urban transmission in the tropics, but also for the emergence outside Asia.55 In conclusion, our study emphasizes the increased risks for human exposure to JEV vector mosquitoes in cities with pigs, implying that JEV is not only a rural concern as has traditionally been claimed.21,56

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