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The Risk of Urban Yellow Fever Resurgence in *Aedes*-Infested American Cities

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Abstract: *Aedes aegypti*, historically known as the yellow fever mosquito, transmits a great number of other viruses like Dengue, West Nile, Chikungunya, Zika, Mayaro and perhaps Oropouche, among others. Well established in Africa and Asia, Aedes mosquitoes are now increasingly invading large parts of the American continent, and hence the risk of urban yellow fever resurgence in the American cities should be cause of great concern to public health authorities. Although no new urban cycle of yellow fever (YF) was reported in the Americas since the end of an *Aedes* eradication program in the late 1950s, the high number of non-vaccinated individuals that visit endemic areas, that is, South American jungles where the sylvatic cycle of YF is transmitted by canopy mosquitoes, and return to *Aedes*-infested urban areas, increases the risk of resurgence of the urban cycle of YF. We present a method to estimate the risk of urban yellow fever resurgence in dengue endemic cities. This method consists in: 1) to estimate the number of *Aedes* mosquitoes that explains a given dengue outbreak in a given region; 2) calculate the force of infection caused by the introduction of one infective individual per unit area in the endemic area under study. 3) using the above estimates, calculate the probability of at least one autochthonous yellow fever case per unit area produced by one single viremic traveller per unit area arriving from a yellow fever endemic or epidemic sylvatic region at the city studied. We demonstrate that, provided the vector competence (in the sense that will be explained later) of *Aedes aegypti* to transmit YF is greater than 0.7, one infected traveller can introduce urban YF in a dengue endemic area.

Keywords: Yellow Fever; Dengue; *Aedes aegypti*; mathematical models; risk.

INTRODUCTION

Introduced into the Americas by the slave trade [1], yellow fever, a highly lethal but vaccine preventable hemorrhagic fever , afflicted the Americas for three centuries. In its urban cycle yellow fever is transmitted by *Aedes aegypti*, a short-winged household mosquito that breeds in clean water collections [2]. The sylvatic cycle is kept by non-human primate reservoirs and is transmitted by two forest-living mosquitoes of the genuses Haemaghogus and Sabethes [3].

The last documented urban YF epidemic in the Americas occurred in 1928 in Rio de Janeiro, Brazil [4]. In the period between 1948 and 1954, the Brazilian National Service of Yellow Fever (SNFA) used the then new insecticide DDT in large areas of regions [5]. As a result, in March 1955 the last focus of *Aedes aegypti* was detected and eliminated from the state of Bahia in North-Eastern Brazil [6]. Over the next three decades, the mosquito was considered eradicated from Brazil.

In the 1980s, *A. aegypti* returned to the urban centers of South America [7]. Although no new urban cycle of YF occurred in South-America since the end of the *Aedes* eradication program, the high number of non-vaccinated individuals that visit jungle areas, where the sylvatic cycle of YF already exists, and return to Aedes-infested urban areas, poses a threat of resurgence of the urban cycle of YF. In recent years, an increasing number of human yellow fever cases has been reported in many Brazilian cities where forest remains with non-human primate reservoirs and the presence of the sylvatic mosquitoes are in close contact with human settlements [8].

In 2016, a yellow fever outbreak, characterized as a sylvan or jungle epizootic [9], occurred in Minas Gerais, Brazil. By mid-April 2017 there were 2,422 cases of yellow fever

reported, including 326 deaths. The case fatality rate (CFR) was 34%. After this outbreak there was an important reduction in notified cases incidence in epizootic regions [10]. Until the middle of February, 2018, the State of São Paulo reported 202 cases of yellow fever with 79 deaths. None of those recent cases reported in Brazil was attributed to the *Aedes aegypti* mosquito, the species which was responsible for all urban outbreaks in the first half of the last century. Urban yellow fever transmission, however, still occurs continuously in Africa, as demonstrated by the outbreaks in Angola in 2016 [11] and in Ivory Coast in 2017 [12]. Moreover, in October 2017, the majority of the larger parks in the city of São Paulo was closed to visitation due to the death of Allouatta monkeys infected with yellow fever (see http://www1.folha.uol.com.br/cotidiano/2017/10/1928920-macaco-e-achado-morto-com-febre-amarela-e-horto-florestal-e-fechado.shtml).

Recently, *Aedes* mosquitoes spread throughout the whole American continent (more than 50% of the USA states are endemic for both *Aedes aegypti* and *Aedes albopictus* [13]) which are present in increasing numbers in virtually all cities of tropical America [14], in particular in urban centers of the Brazilian coast [15]. The yearly increase in the number of dengue cases and more recently of chikungunya and Zika virus demonstrates the high density of *Aedes* mosquitoes in urban centers of Brazilian coastal regions. These latter urban centers are not included in areas where yellow fever vaccination is recommended. Considering that these areas comprise about 60% of the entire Brazilian population the risk of urban yellow fever resurgence is imminent [8]. The recent invasion of zika virus in the North American continent demonstrate the presence of Aedes mosquitoes and hence the risk of urban yellow fever resurgence in North American cities as well [16].

Since aedes mosquitoes are competent to transmit the YF virus [17], it is worthwhile to try to estimate the probability of at least one YF case resulting from the arrival of one infected traveller to a aedes-infested area. In addition, it is possible to estimate the expected number of YF cases and deaths after one year of its introduction.

In a recent paper, Couto-Lima et al. [17] demonstrated that the anthropophilic mosquitoes *Aedes aegypti* and *Aedes albopictus* as well as the YFV-enzootic mosquitoes *Haemagogus leucocelaenus* and *Sabethes albiprivus* from the YFV-free region of the Atlantic coast were highly susceptible to American and African YFV strains. In that paper, the authors demonstrated that the vectorial competence to transmit YFV of urban aedes mosquitoes is very similar their vector competence to transmit dengue, and this justifies using well known transmission parameters for dengue as a first approximation to the same parameters for YFV transmission. Of course the errors of such approximation must be estimated and this is one of the purposes of this paper.

Recently Massad et al. [18] proposed a method to estimate the size of population of *Aedes* mosquitoes using dengue incidence data. In that paper, the authors estimated the number of infected, latent and susceptible mosquitoes and used the total number of mosquitoes to roughly estimate the expected number YF cases resulting from the arrival of one infected traveller. Here we extend this analysis by including the estimation of the probability of at least one YF case resulting from the arrival of one infected traveller to a aedes-infested area for several values of vectorial competence of urban aedes mosquitoes.

The objective of this study is, therefore, to estimate the risk of urban yellow fever resurgence in Aedes-infested cities.

This paper is organized as follows. After this introduction we describe the method proposed in [18] to estimate the number of aedes mosquitoes from dengue incidence data. This is done step-by-step to allow the reproducibility of our results by other authors. The next section shows an illustration of the method for the case of the city of Rio de Janeiro with dengue incidence data from 2012, one of the worst years in the dengue history of that city. In this section we also comment on the limitations of the method and estimate the errors associated with the calculated outcomes. In the final section we present the conclusion of this work.

METHODS

The method used to estimate the risk of urban yellow fever resurgence in *Aedes*-infested cities consists in estimating the number of *Aedes* mosquitoes that explains a given dengue outbreak. Based on the above estimates we calculated the probability of at least one autochthonous urban yellow fever case per unit area acquired as a result of importation by a traveller arriving from a yellow fever endemic or epidemic sylvatic region in his/hers infectiousness period (first generation). Finally, we estimated the expected number of autochthonous infections per unit area produced by that infected traveller (index case) per unit area.

We begin by fitting a continuous function, denoted $Incidence_{DENV}(t)$, to the number of reported dengue cases multiplied by 4, that is, by taking into account the 4:1 asymptomatic-to-symptomatic ratio [19]. The dengue incidence data (from [20]), shown in Table 1, were fitted to the chosen function:

$$Incidence_{DENV}(t) = c_1 \exp\left[-\frac{(t-c_2)^2}{c_3}\right] + c_4 \quad (1)$$

representing the time-dependent dengue infection incidence. In equation (1) c_1 is a scale parameter that determines the maximum incidence, c_2 is the time at which the maximum incidence is reached, c_3 represents the width of the time-dependent incidence function and c_4 is just another scaling parameters. Equation (1) is intended to reproduce a "Gaussian" curve and so c_1 and c_4 are just scale parameters but c_2 represents the "mean" (and mode or maximum) time and c_3 represents the "variance" of the time distribution of cases. All parameters $c_i, i = 1,...,4$ were fitted to model (1) by the Bootstrap method [21] the results of which are shown in Table 2.

Table 1. Number of dengue infection in Rio de Janeiro, 2011-2012

Time (months)	**Number of Infections***
1	476
2	2348
3	2628
4	4212
5	6932
6	1272
7	22688
8	46180
9	104992
10	149856
11	118124
12	45956
13	12368
14	6732
15	3080
16	3032
17	2996

* Number of reported cases multiplied by 4 to take account of asymptomatic cases

Table 2. **Parameters' values (mean, lower bound and upper bound) fitted to Equation (1) by Bootstrap technique.**

Parameter	Mean	Lower Bound	Upper Bound
c_1	147144	134944	159879
c_2	10.004	9.9147	10.073
c_3	3.49035	2,2158	4.0631
c_4	1500.02	1388.89	1625.31

Figure 1 illustrates the case of the 2011-2012, the years with the outbreak with the highest incidence of dengue in the history of Rio de Janeiro, Brazil.

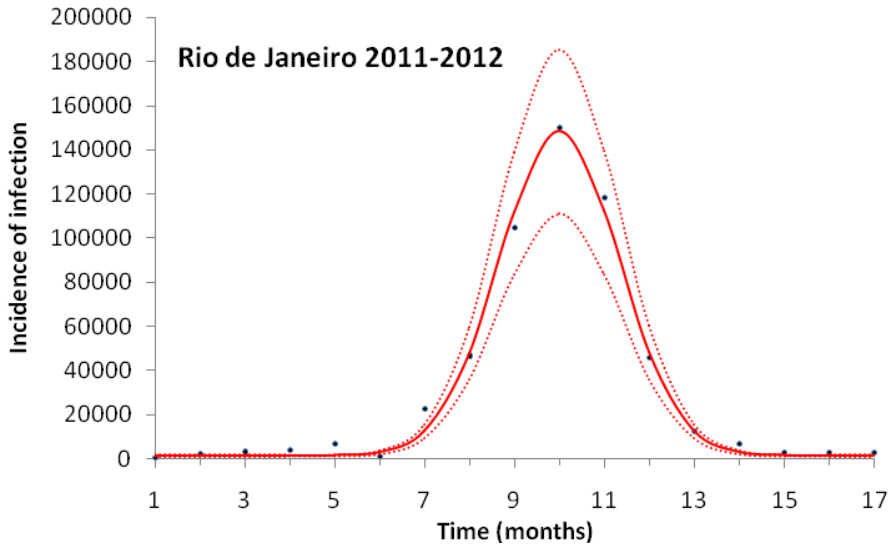

Figure 1. Fitting a function (eq. 1) to dengue incidence of infections between October 2011 and December 2012 in Rio de Janeiro. Dots represent the notified data multiplied by 4, continuous line the mean fitted incidence and dotted lines de 95%C.I.

Dengue incidence is defined as the product of the force of infection, $\lambda(t)$, times the number of susceptible individuals, $S_H(t)$. The force of infection is defined as $\lambda(t) = \frac{abI_M(t)}{N_H(t)}$, where $N_H(t)$ denotes the total human population, a is the mosquitoes biting rate, b is the

fraction of those bites produced by the infectious mosquitoes $I_M(t)$ that are infective to susceptible humans $S_H(t)$. All the parameters used in these calculations are shown in Tables 3.

Table 3. Model parameters, biological meaning, values and sources. The mean, variance and 95% CI were obtained with 10^4 Monte Carlo simulations. The dimension of rates is months^{-1}.

Parameter	Meaning	Value (Baseline)	Mean	Variance	95% CI
a	*A. aegypti* biting rate	4.104	4.12	0.652	12.76×10^{-3}
b*	Fraction of bites actually infective to humans (Dengue/YF)	0.6	0.6062	0.296	0.0337
μ_H	Human natural mortality rate	1.096×10^{-3}	1.096×10^{-3}	3.19×10^{-8}	7.05×10^{-5}
α_{YF}	YF mortality rate	2	1.98	5.72×10^{-3}	1.12×10^{-5}
$\gamma_{H\text{-}D}$	Dengue recovery rate	0.0768	0.0772	9.12×10^{-3}	3.74×10^{-3}
$\gamma_{M\text{-}D}$	Latency rate in mosquitoes for Dengue	0.0768	0.0772	9.12×10^{-3}	3.74×10^{-3}
$\gamma_{H\text{-}YF}$	YF recovery rate	0.32	0.3216	4.68×10^{-2}	9.12×10^{-4}
$\gamma_{M\text{-}YF}$	Latency rate in mosquitoes for YF	0.32	0.3216	4.68×10^{-2}	9.12×10^{-4}
μ_M	Natural mortality rate of mosquitoes	2.52	2.332	4.20×10^{-3}	1.34×10^{-3}
c*	*A. aegypti* susceptibility to (Dengue/YF)	0.54	0.5265	0.249	0.03191

* These rates for YF varied along the simulations

From the fitted incidence, $Incidence_{DENV}(t)$ we calculate the number of infective mosquitoes $I_M(t) = \dfrac{Incidence_{DENV}(t)N_H(t)}{abS_H(t)}$, where $S_H(t)$ can be estimated by the previous history of dengue outbreaks [19]. From the number of infectious mosquitoes we calculated the number of latent mosquitoes, $L_M(t)$, given by $L_M(t) = \dfrac{1}{\gamma_M}\left[\dfrac{d}{dt}I_M(t) + \mu_M I_M(t)\right]$, where $\dfrac{1}{\gamma_M}$ is the average duration of the extrinsic incubation period and μ_M is the natural mortality rate of mosquitoes. The number of susceptible mosquitoes is given by $S_M(t) = \dfrac{N_H}{acI_H(t)}\left[\dfrac{d}{dt}L_M(t) + (\mu_M + \gamma_M)L_M(t)\right]$, where c is the fraction of the mosquitoes that

bite infective humans and acquire the infection, and $I_H(t)$ is the number of humans infected with dengue. The estimated total number of mosquitoes is $N_M(t) = S_M(t) + L_M(t) + I_M(t)$. We are going to need the derivatives of these functions:

$$\frac{dN_M(t)}{dt} = \frac{dS_M(t)}{dt} + \frac{dL_M(t)}{dt} + \frac{dI_M(t)}{dt} \tag{2}$$

For details of the above calculations, see [18].

For the calculation of the probability that one infected travellers produces at least one autochthonous YF case and the total number of Y cases and deaths one year after the introduction of the infection we used a variant of the classical Ross-Macdonald model, described in details in [22] and [23].

The populations involved in the transmission are human hosts and mosquitoes. Therefore, the population densities per unit area are divided into the following compartments: susceptible humans denoted S_H; infected humans, I_H; recovered (and immune) humans, R_H; total humans, N_H; susceptible mosquitoes, S_M; infected and latent mosquitoes, L_M; infected and infectious mosquitoes, I_M. The parameters appearing in the model are defined in Table 3.

The model is defined by the following equations:

$$\frac{dS_{H-YF}}{dt} = -ab_{YF} I_{M-YF} \frac{S_{H-YF}}{N_H} + \mu_H (N_H - S_{H-YF}) + \alpha_{YH} I_{H-YF}$$

$$\frac{dI_{H-YF}}{dt} = ab_{YF} I_{M-YF} \frac{S_{H-YF}}{N_H} - (\mu_H + \gamma_{H-YF} + \alpha_{YH}) I_{H-YF}$$

$$\frac{dR_{H-YF}}{dt} = \gamma_{H-YF} I_{H-YF} - \mu_H R_{H-YF}$$

$$\frac{dS_{M-YF}}{dt} = -ac_{YF} S_{M-YF} \frac{(I_{H-YF} + I_0(t))}{N_H} + \mu_M (L_{M-YF} + I_{M-YF}) + \frac{dN_M}{dt}$$

$$\frac{dL_{M-YF}}{dt} = ac_{YF} S_{M-YF} \frac{(I_{H-YF} + I_0(t))}{N_H} - \gamma_{M-YF} L_{M-YF} - \mu_M L_{M-YF}$$

$$\frac{dI_{M-YF}}{dt} = \gamma_{M-YF} L_{M-YF} - \mu_M I_{M-YF}$$

$$\frac{dN_M}{dt} = \text{as in equation (2)}$$

$$\begin{aligned} N_H &= S_{H-YF} + I_{H-YF} + R_{H-YF} \\ N_{M-YF} &= S_{M-YF} + L_{M-YF} + I_{M-YF} \end{aligned} \tag{3}$$

Remark: Note that the expression for mosquitoes $\frac{dN_M}{dt}$ *was estimated from dengue data (see* [18] *for details).*

System (3) was solved assuming that one infected individual was introduced by unit area, $I_0(t)$, (see [23]) at $t = t_0$ and remains infective for a period of $(\mu_H + \gamma_{H-YF} + \alpha_{YH})^{-1}$ months, that is:

$$I_0(t) = \exp[-(\mu_H + \gamma_{H-YF} + \alpha_{YH})(t - t_0)]\theta(t - t_0) \tag{4}$$

The total number of YF cases after one year of its introduction ($\Delta = 12$ months), YF_{cases}, is given by:

$$YF_{cases} = \int_{t_0}^{t_0+\Delta} ab_{YF} I_{M-YF} \frac{S_{H-YF}}{N_H} dt \tag{5}$$

The total number of YF deaths after one year of its introduction, YF_{deaths}, ($\Delta = 12$ months) is given by:

$$YF_{deaths} = \alpha_{YF} \int_{t_0}^{t_0+\Delta} ab_{YF} I_{M-YF} \frac{S_{H-YF}}{N_H} dt \tag{6}$$

The risk of urban YF resurgence is defined as the probability of at least one autochthonous case per unit area produced by one single infected individual per unit area arrived at the area during his/her infectiousness period. For this, the fourth and fifth equations of system (3) take the form:

$$\begin{aligned} \frac{dS_{M-YF}}{dt} &= -ac_{YF} S_{M-YF} \frac{(I_0(t))}{N_H} + \mu_M (L_{M-YF} + I_{M-YF}) + \frac{dN_M}{dt} \\ \frac{dL_{M-YF}}{dt} &= ac_{YF} S_{M-YF} \frac{(I_0(t))}{N_H} - \gamma_{M-YF} L_{M-YF} - \mu_M L_{M-YF} \end{aligned} \tag{7}$$

From the system (3) with equations fourth and fifth replaced by equations (7) we can calculate the force of infection $\lambda_{YF}(t)$

This force of infection can now be used in a non-homogeneous simple birth process to calculate the required probabilities.

The probability generating function of such process is given by:

$$P(x,t) = \left\{ 1 + \frac{1}{\frac{e^{\rho(t)}}{x-1} - \int_0^t \lambda_{YF}(\tau) e^{\rho(\tau)} d\tau} \right\}^a \tag{8}$$

where $a = I_0(0)$, $\lambda_{YF}(t) = ab_{YF} \frac{I_{M-YF}(t)}{N_H}$ and $\rho(t) = \int_0^t -\lambda_{YF}(\tau)\theta(\tau - t_0) d\tau$.

The quantity $\rho(t)$ must be calculated numerically.

Expanding (8) in powers of x we find:

$$p_n(t) = \sum_{j=0}^{\min(n,a)} \binom{a}{j} \binom{a+n-j-1}{a-1} \alpha^{a-j} \beta^{n-j} (1-\alpha-\beta)^j \tag{9a}$$

$$p_0(t) = \alpha^a \tag{9b}$$

where $\alpha = 1 - \dfrac{1}{e^{\rho(t)} + \int_0^t \lambda(\tau)\theta(\tau - t_0)e^{\rho(\tau)}d\tau}$ and $\beta = 1 - \dfrac{e^{\rho(t)}}{e^{\rho(t)} + \int_0^t \lambda(\tau)\theta(\tau - t_0)e^{\rho(\tau)}d\tau}$

The risk of urban YF resurgence was defined as the probability of one secondary case generated by a single infected traveller ($a = I_0(0) = 1$) along his/her infectiousness period. In terms of equation (9a) it is the probability of a secondary infection, that is, $p_2(t)$, or:

$$p_2(t) = \sum_{j=0}^{1} \binom{a}{j} \binom{a+n-j-1}{a-1} \alpha^{a-j} \beta^{n-j} (1-\alpha-\beta)^j = \alpha\beta^2 + \beta(1-\alpha-\beta) \tag{10}$$

The total number of YF cases and deaths after one year and the probability of at least one autochthonous case produced by introducing in the area a density of infected individuals (equal to one) during his/her infectiousness period where calculated for various degrees of the aedes vector competence. This parameter is defined as a factor less than one that represents both the probability of infection from vector to host and from host to vector, b and c, respectively, as in Table 3.

RESULTS

Figure 1 illustrates the fitting procedure of dengue incidence with the case of the 2011-2012 dengue outbreak in Rio de Janeiro, Brazil. Figure 2 shows the calculation for the number of mosquitoes for the city of Rio de Janeiro in the same period.

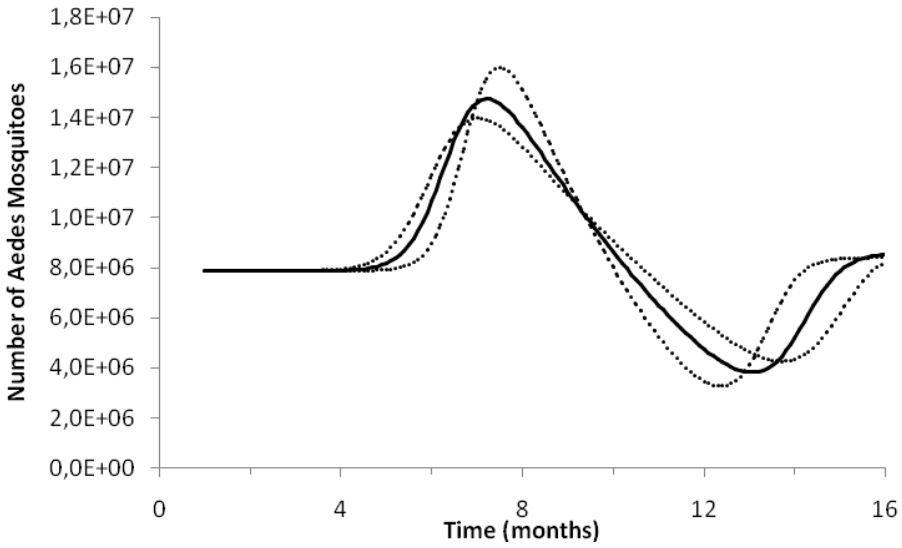

Figure 2. Estimation of the size of the *Aedes* mosquitoes' population from dengue incidence between October 2011 and December 2012 in Rio de Janeiro. Continuous line the mean number of mosquitoes and dotted lines de 95%C.I.

To test the method's accuracy for estimating the number of mosquitoes, we used the above equations to calculate the number of *Aedes* mosquitoes in the neighborhood of Olaria in Rio de Janeiro. In the 2000 census, Olaria had an estimated population of 62,509 inhabitants in an area of around 4 km^2. This neighborhood was chosen because in 2008 Maciel-de-Freitas, Eiras and Lourenço-de-Oliveira [25] estimated, through the MosquiTrap and aspirator methods, the population size of *Aedes aegypti*. The two devices were used in the same area at the same time. In the assumed 0.8 km^2 area covering the average flight range of *Aedes* mosquitoes, the authors collected 3,505 and 4,828 female mosquitoes in the MosquiTrap and aspirator,

respectively, totalizing 8,333 female mosquitoes (note that it is assumed that the two capturing techniques are equally efficient so each technique capture about half the total number of mosquitoes in the area). Using the data from dengue in the same period, we estimated a total *Aedes* population in each of the 5 areas of 0.8 km^2 of Olaria in a period of two weeks as 8,145±284 female mosquitoes, which is a good approximation to the empirical data.

To calculate the expected number of yellow fever cases generated by one infected traveller after one year, we used the Ross-Macdonald model. We introduced a term in the susceptible mosquitoes' equations of the Ross-Macdonald model, so that we reproduce the number of mosquitoes calculated before. The new equation reads $\frac{dS_M}{dt} = -acS_M \frac{I_H}{N_H} + \mu_M (L_M + I_M) + \frac{dN_M}{dt}$, where the new term $\frac{dN_M}{dt}$ is the sum of the derivatives of the equations for the susceptible, latent and infectious mosquitoes described above. The yellow fever specific parameters used were the terms related to vector-competence, c and b, the extrinsic incubation period, $1/\gamma_M$, the duration of viremia $1/\gamma_H$, and the disease-induced mortality rate α_H of infected humans. Using the Ross-Macdonald model for the areas in which the number of mosquitoes can be calculated from dengue data allows the estimation of the expected number of new cases produced by infective travellers arriving at different moments.

As an example of the above procedures, we estimated the expected number of autochthonous cases and deaths generated by introducing a density of infective individuals over the whole city arriving at different times of the year, $t = t_0$, in Rio de Janeiro. Figure 3 shows that the maximum number of autochthonous cases is reached when the imported infection arrives at around 5.6 months, with the number of yellow fever virus infections peaking between 169 and 282 and deaths peaking between 55 and 92.

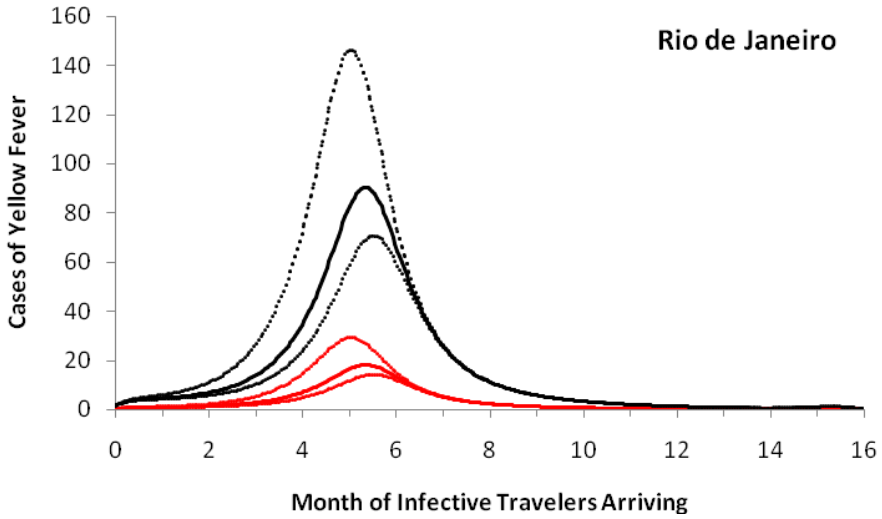

Figure 3. Cases of yellow fever estimated by the model as a function of the month of infective travellers' arrival. Estimation of the number of yellow fever infections (black lines) and mortality (red lines) after one year, if a density (equal to one) of infected individuals were introduced in the city of Rio de Janeiro (2011-2012) at different months of the year. Continuous lines represent the mean and dotted lines the 95%C.I.

The risk of urban yellow fever resurgence depends on the size of *Aedes* mosquitoes' populations and their vectorial competence, defined as the variation in the values of the parameters c and b specific for yellow fever are shown in figure 4.

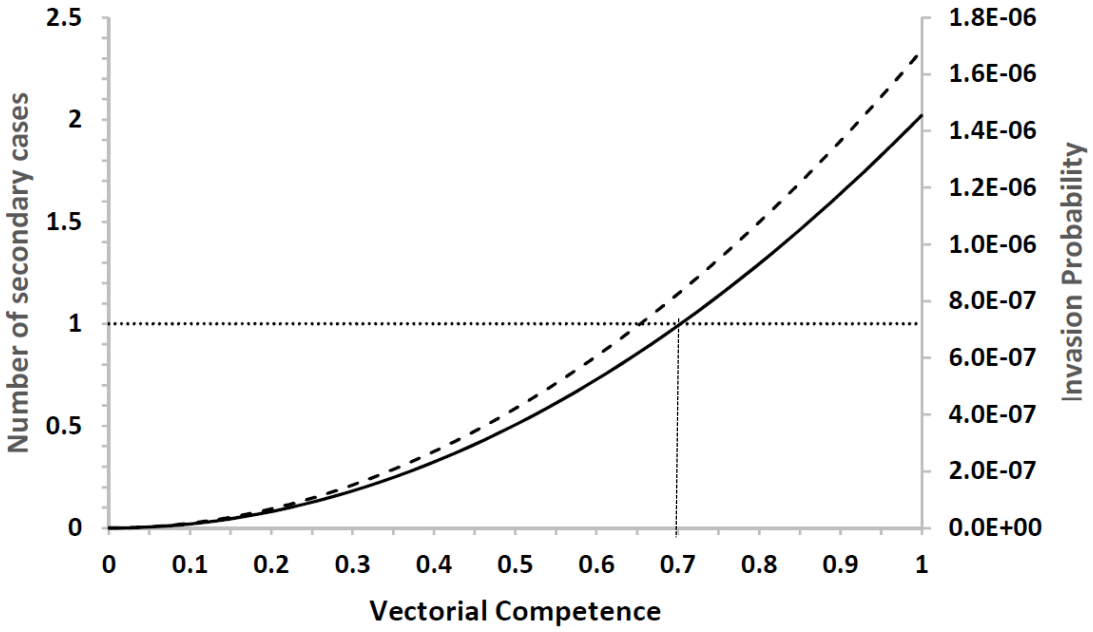

Figure 4. Estimation of the risk of yellow fever introduction in the city of Rio de Janeiro by the arrival of infective travelers (with density one) in week eleven along his/her first generation and the expected number of secondary cases as a function of the relative vector competence. The number of *Aedes* mosquitoes was calculated from dengue incidence in 2011-2012.

Note that there is a threshold in vector competence (0.7, or 70% of *Aedes* competence for YF transmission), below which the expected number of cases produced by one infected traveller along his/her infectiousness period is less than one.

DISCUSSION

In this paper we calculated the theoretical number of YF infections after one year, if one infective traveller arrived in February 2012, as well as the probability that the infected individual will produce a secondary case along his/her infectiousness period. Both estimations are presented in figure 4 varying the simulated values of the local *Aedes* vector competence and the risk incurred by the city of Rio de Janeiro of urban yellow fever re-introduction, $Risk_{UYFR}(t)$, if in

February 2012 the city of Rio de Janeiro were invaded by infected travellers with density one per area of the city.

We showed that there is a low but non-negligible risk of urban yellow fever resurgence in dengue endemic areas due to the high *Aedes* mosquitoes' densities in these areas. The actual risk will be dependent on the probability that at least one infective human arrives at the right moment of the year, that is, when the local population of *Aedes* mosquitoes is increasing in size and also on their vector competence for yellow fever transmission. Moreover, the vector competence of Aedes mosquitoes in transmitting YF is a crucial factor in determining the transmission of YF and for the risk of urban-YF resurgence. It is believed that the vector competence of *Aedes aegypti* for transmitting YF is lower than that for transmitting dengue [8] (see, however [17]). This fact, in addition to the relatively high vaccination coverage of large regions of the Brazilian territory, may explain why urban-YF has not resurged in these areas. However, the coastal areas of Brazil, including the South Eastern region currently heavily hit by YF epizooties, are not included in the routine vaccination programmes neither of the Brazilian Ministry of Health nor the States Health Authorities. This fact, associated with high density of Aedes mosquitoes and a large number of epizooties that keep recurring in these non-vaccinate areas pose a serious risk of urban-YF resurgence in these regions.

One important limitation of our approach is the fact that the Ross-Macdonald model used in the calculations assumed a homogenously mixing of humans and mosquitoes. Moreover, when we introduce an infected traveller, it is assumed that this infected individual will interact with all the mosquitoes in the area considered, a highly unlikely event in the real world. Therefore, the results of our methods would be more reliable the smaller the geographical are considered,

ideally the area covered by the mosquitoes' range of flight (the 0.8 km^2 mentioned in the Olaria example).

Finally, estimating the risk of urban yellow fever resurgence is crucial for designing an optimum vaccination strategy considering the yellow fever vaccine adverse events [26]. Moreover, since Aedes is invading large parts of the American continent (it is well established in Africa and Asia) the risk of urban yellow fever resurgence in the American cities should be cause of great concern to public health authorities. We believe that this work represents a step forward to understanding the magnitude of the problem.

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