<u>Electronic supplementary material</u>

Mosquito-borne transmission in urban landscapes: the missing link between vector abundance and human density

Victoria Romeo-Aznar¹, Richard Paul^{2,3}, Olivier Telle⁴ and Mercedes Pascual¹

¹Department of Ecology and Evolution, University of Chicago, Chicago IL, USA 60637.

²Institut Pasteur, Functional Genetics of Infectious Diseases Unit, 75724 Paris Cedex 15, France.

³Centre National de la Recherche Scientifique (CNRS), Génomique évolutive, modélisation et santé

UMR 2000, 75724 Paris Cedex 15, France.

⁴Centre National de la Recherche Scientifique (CNRS), CSH, Delhi, Inde.

doi: 10.1098/rspb.2018.0826

Parameter	Value	Description
$\lambda_{ m N}$	1/(70*365) days ⁻¹	Humans' birth rate
$\mu_{ m N}$	1/(70*365) days ⁻¹	Humans' mortality rate
γ	1/5 days-1	Recovery rate
δ	0.75	virus transmission probability given a
		mosquito bite
α	0.23 days-1	Biting rate
Without temperature		
λ_{M}	0.09 days ⁻¹	Mosquitoes' birth rate
μ_{M}	0.09 days ⁻¹	Mosquitoes' mortality rate
With temperature (T)		
$\lambda_{M}(T)$	0.003 T days ⁻¹ C ⁰⁻¹	Mosquitoes' birth rate
μ _M (T)	(0.08326+e ^{-T/6 °C}) days ⁻¹	Mosquitoes' mortality rate
Spatial		

F(V(N)) (0.059+0.342822071 e^{-0.003777299 V(N)}) days⁻¹

Mosquitoes' spatial-dispersion rate

Table S1. Model parameters values (see main text). Parameter values are the same for all models. The mortality and birth rates of mosquitoes are the two parameters that vary periodically with temperature. In the spatial model, F is the emigration rate from a spatial unit per mosquito. The values are adapted from Otero & Solari, 2010 and Romeo Aznar et al., 2013.



Fig. S1. Diagram illustrating the steps to compute the probability *P* for a given typology.



Figure S2. Rescaled force of infection for (A) linear and (B) quadratic V(N) for different values of N and biting rates. The change of scale is: (FOI-min(FOI))/max(FOI-min(FOI)), where minimum and maximum values are with respect to N (for a given biting rate). Regardless of biting rate, we observe a decreasing and increasing behavior with N for the linear and quadratic cases respectively.



N Figure S3. R_0 as a function of *N*.



Figure S4. Temporal evolution of incidence for linear V(N) in the non-spatial model with temperature forcing (see Fig. 2). Lines correspond to the mean values of incidence and shadow areas, to the 90% confidence intervals. Only simulations with total cumulated cases representing more than 1% of the total host population were included.



Figure S5. Number of infected hosts as a function of time In the spatial model, without (A) and with (B) temperature forcing (see Fig. 3 in main text). Lines correspond to the mean values of incidence and shadow areas, to the 90% confidence intervals. Only simulations with total cumulated of infected cases representing more than 1% of the total host population were included.



Figure S6. Probability P (see main text) as a function of N for dengue Delhi data for, Dep. LD (black), Dep. MD (red), Rich (blue) and Planned (green) typologies. A) For years 2008 and 2009 together. Per year for B) Dep. MD and C) Rich typologies. (The Dep. LD data is not shown for the two separate years because with only three points, it does not exhibit enough points to address the existence or lack of a trend).



Figure S7. The natural logarithm of *V* is plotted as a function of the natural logarithm of *N* for Dep. MD typology. The solid line corresponds to the best linear fit and the dashed line to the best fit with a fixed slope of one. In this case, contrary to Fig 4C, the assumed outlier point (first point with the smallest *N*) is taken into account. A slope value of k= 1.45 +/- 0.17 was obtained.