

UNIVERSIDAD EAFIT

BACHELOR PUBLICATION

**Application of Sensitivity- and
Uncertainty-based techniques for the
assessment of epidemiological models in
real-life study cases**

Author:
Daniel Rojas Díaz

Supervisor:
Carlos M. Vélez Sánchez, Eng.Dr.
Co-Supervisor:
María E. Puerta Yepes, Ph.D.

*Submitted in partial fulfillment of the requirements
for the degree of Bachelor of science in biology*

Research group: Modelado matemático
Departamento de Ciencias Biológicas

October 31, 2019

“We sail within a vast sphere, ever drifting in uncertainty, driven from end to end. When we think to attach ourselves to any point and to fasten to it, it wavers and leaves us; and if we follow it, it eludes our grasp, slips past us, and vanishes forever.”

Blaise Pascal

UNIVERSIDAD EAFIT

Abstract

Escuela de Ciencias
Departamento de Ciencias Biológicas

Bachelor of science in biology

Application of Sensitivity- and Uncertainty-based techniques for the assessment of epidemiological models in real-life study cases

by Daniel Rojas Díaz

Uncertainty analysis (UA) and sensitivity analysis (SA) are tools to assess and to quantify the uncertainty spread from the input factors (parameters and initial states) to the model output, taking into account the effect of the interactions among those factors. Throughout the following works, I treat UA as a graphical assessment of uncertainty propagation based on Monte Carlo simulation, which makes it possible to state a range for the model output in cases where it is considered relevant. On the other hand, I privilege the global approach for SA instead of the local one, since the first attempts to quantify the uncertainty contribution of the model factors in their entire distribution range while the second one is only informative for a single locus in the distribution. In this way, when applying global UA/SA on a model, it is possible to identify those factors that mostly determine the model behavior. Furthermore, I have noticed that the concepts and principles of UA/SA are associated with other main tasks in modeling, as factors estimation and confidence intervals achievement: Briefly, those non-identifiable factors in a model (factors whose value can not be estimated uniquely from some information about output data) should belong to the categories of non-sensible or sensitive but correlated from SA; and, the sub-space of the space of factors where the factors may jointly exist producing a model output that fits, in some extent, to a given output data, could be approximately estimated with UA-based approaches, constituting a new kind of confidence interval. Thus, in this compendium, I present five works related to the applications of UA/SA techniques as well as its relevance. The objective of those applications evolves from the most logically immediate to some derived and more complex ones, though still preserving the model pertinence as a central topic.

Acknowledgements

First and foremost, I am in debt to the circumstances. I was in the right place at the perfect time and took the appropriate decisions, even the wrong ones, to enroll myself in research. The different papers that compose this compendium are the result of joint work and could not have been concluded without the help of my family, friends, mentors, and colleagues, though the distinctions among those categories have been the fuzziest.

I am much in debt with my mother, Nora, and my aunt, Melba, whose efforts were even higher than my ones to set me able to achieve my goals. Next, I am deeply grateful to my mentors, Carlos M. Vélez, María E. Puerta, and Carlos A. Cadavid, to whom I owe my ideas, knowledge, and enthusiasm. I also want to thank my colleagues, Alexandra Cataño and Diana P. Lizarralde, with whom I shared long hours of work, laughter, and learning.

Most of the research I did for this work was performed during my professional practice at Universidad EAFIT, for which I thank the Department of professional practices by their management, as well as my friend and professor, Daniel I. Velásquez, without him, I could not arrive at time to my workplace nor enjoy a lot of nice trips. I am also thankful to the whole family Villegas-Ruíz for the enormous support I received from them throughout my academic internship at EAFIT.

Finally, I would like to thank everyone who feels that they contribute to these works and has not been explicitly mentioned. All of their contributions helped to bring these works to light.

Contents

Abstract	v
Acknowledgements	vii
1 An alternative model to explain the vectorial capacity using as example <i>Aedes aegypti</i> case in dengue transmission	1
1.1 Authors	1
1.2 Abstract	1
2 A novel algorithm for confidence sub-contour box estimation: An alternative to traditional confidence intervals	7
2.1 Authors	7
2.2 Abstract	7
3 Global Sensitivity and Uncertainty Analyses - Confidence Subcontour Box (GSUA-CSB) Toolbox	11
3.1 Authors	11
3.2 Abstract	11
4 Influence of pulse-type inputs on parameter estimation and chemical control assessment in a dengue deterministic model	13
4.1 Authors	13
4.2 Abstract	13
5 Sensitivity, uncertainty and identifiability analyses to define a dengue transmission model with real data of an endemic municipality of Colombia	19
5.1 Authors	19
5.2 Abstract	19

To Melba

*Both her outright advice and unconditional support have
forged and will continue to shape the path I started walking,
and where I put this work as the first step.*

Chapter 1

An alternative model to explain the vectorial capacity using as example *Aedes aegypti* case in dengue transmission

Submitted to	Status	DOI (if available)
Heliyon	Published	10.1016/j.heliyon.2019.e02577

1.1 Authors

Alexandra Catano-Lopez¹, Daniel Rojas-Díaz², Henry Laniado¹, Sair Arboleda-Sánchez³, María Eugenia Puerta-Yepes¹, Diana Paola Lizarralde-Bejarano¹

1 Dept. Ciencias Matemáticas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

2 Dept. Ciencias Biológicas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

3 Grupo de Biología y Control de Enfermedades Infecciosas-BCEI, Universidad de Antioquia, Medellín, Antioquia, Colombia

1.2 Abstract

Vectorial capacity (VC), as a concept that describes the potential of a vector to transmit a pathogen, has had historical problems related to lacks in dimensional significance and high error propagation from parameters that take part in the model to output. Hence, values estimated with those equations are not sufficiently reliable to consider in control strategies or vector population study. In this paper, we propose a new VC model consistent at dimensional level, i.e., the definition and the equation of VC have same and consistent units, with a parameter estimation method and mathematical structure that reduces the uncertainty in model output, using as a case of study an *Aedes aegypti* population of the municipality of Bello, Colombia. After a literature review, we selected one VC equation following biological, measurability and dimensional criteria, then we rendered a local and global sensitivity analysis,

identifying the mortality rate of mosquitoes as a target component of the equation. Thus, we studied the Weibull and Exponential distributions as probabilistic models that represent the expectation of mosquitoes infective life, intending to include the best distribution in a selected VC structure. The proposed mortality rate estimation method includes a new parameter that represents an increase or decrease in vector mortality, as it may apply. We noticed that its estimation reduces the uncertainty associated with the expectation of mosquitoes' infective life expression, which also reduces the output range and variance in almost a half.

Bibliography

- [1] A. P. G. Almeida, S. S. S. G. Baptista, C. A. G. C. C. Sousa, M. T. L. M. Novo, H. C. Ramos, N. A. Panella, M. Godsey, M. J. Simões, M. L. Anselmo, N. Komar, C. J. Mitchell, and H. Ribeiro. Bioecology and Vectorial Capacity of *Aedes albopictus* (Diptera: Culicidae) in Macao, China, in Relation to Dengue Virus Transmission. *Journal of Medical Entomology*, 42(3):419–428, 2005.
- [2] R. M. Anderson and R. M. May. *Infectious diseases of humans: Dynamics and control*. Oxford University press, 1991.
- [3] S. Arboleda, N. Jaramillo-O., and A. T. Peterson. Mapping environmental dimensions of dengue fever transmission risk in the aburrá valley, colombia. *International Journal of Environmental Research and Public Health*, 6(12):3040–3055, Dec. 2009. doi: 10.3390/ijerph6123040. URL <https://doi.org/10.3390/ijerph6123040>.
- [4] G. Archer, A. Saltelli, and I. Sobol. Sensitivity measures, anova-like techniques and the use of bootstrap. *Journal of statistical computation and simulation*, 58:99–120, 1997.
- [5] D. R. Barnard, K. H. Posey, D. Smith, and C. E. Schreck. Mosquito density, biting rate and cage size effects on repellent tests. *Medical and Veterinary Entomology*, 12(1):39–45, Jan. 1998. doi: 10.1046/j.1365-2915.1998.00078.x. URL <https://doi.org/10.1046/j.1365-2915.1998.00078.x>.
- [6] O. J. Brady, H. C. J. Godfray, A. J. Tatem, P. W. Gething, J. M. Cohen, F. E. McKenzie, T. A. Perkins, R. C. Reiner, L. S. Tusting, M. E. Sinka, C. L. Moyes, P. A. Eckhoff, T. W. Scott, S. W. Lindsay, S. I. Hay, and D. L. Smith. Vectorial capacity and vector control: reconsidering sensitivity to parameters for malaria elimination. *Transactions of The Royal Society of Tropical Medicine and Hygiene*, 110(2):107–117, 01 2016. ISSN 0035-9203. doi: 10.1093/trstmh/trv113. URL <https://doi.org/10.1093/trstmh/trv113>.
- [7] D. G. Cacuci. *Sensitivity & Uncertainty Analysis, Volume 1: Theory*. Chapman and Hall/CRC, Boca Raton, 2003.
- [8] J. Cariboni, D. Gatelli, R. Liska, and A. Saltelli. The role of sensitivity analysis in ecological modelling. *Ecological Modelling*, 203(1-2):167–182, 2007.
- [9] J. M. Carrasco, E. M. Ortega, and G. M. Cordeiro. A generalized modified weibull distribution for lifetime modeling. *Computational Statistics & Data Analysis*, 53(2):450–462, dec 2008. doi: 10.1016/j.csda.2008.08.023. URL <https://doi.org/10.1016/j.csda.2008.08.023>.

- [10] C. Christiansen-Jucht, P. E. Parham, A. Saddler, J. C. Koella, and M.-G. Basáñez. Temperature during larval development and adult maintenance influences the survival of *Anopheles gambiae* s.s. *Parasites & Vectors*, 7(1), nov 2014. doi: 10.1186/s13071-014-0489-3. URL <https://doi.org/10.1186/s13071-014-0489-3>.
- [11] R. C. Christofferson and C. N. Mores. Estimating the magnitude and direction of altered arbovirus transmission due to viral phenotype. *Plos one*, 6:e16298, 2011.
- [12] P. Damos and P. Soulopoulou. Do insect populations die at constant rates as they become older? contrasting demographic failure kinetics with respect to temperature according to the weibull model. *PLOS ONE*, 10(8):e0127328, aug 2015. doi: 10.1371/journal.pone.0127328. URL <https://doi.org/10.1371/journal.pone.0127328>.
- [13] E. F. de Oliveira, E. T. Oshiro, W. de Souza Fernandes, P. G. Murat, M. J. de Medeiros, A. I. Souza, A. G. de Oliveira, and E. A. B. Galati. Experimental infection and transmission of leishmania by *Lutzomyia cruzi* (Diptera: Psychodidae): Aspects of the ecology of parasite-vector interactions. *PLOS Neglected Tropical Diseases*, 11(2):e0005401, feb 2017. doi: 10.1371/journal.pntd.0005401. URL <https://doi.org/10.1371/journal.pntd.0005401>.
- [14] K. Dietz, L. Molineaux, and A. Thomas. A malaria model tested in the African savannah. *Bull*, 50:347–357, 1974.
- [15] C. Dye. Vectorial capacity: Must we measure all its components? *Parasitology Today*, 2(8):203–209, 1986.
- [16] I. Fernández-Salas and M. A. Flores-Leal. El papel del vector *Aedes aegypti* en la epidemiología del dengue en México. *Salud Pública de México*, 37:1–52, 1995. ISSN 0036-3634.
- [17] D. N. Gujarati. *Basic econometrics*. Gary Burke, 2003.
- [18] Instituto de Hidrología, Meteorología y Estudios Ambientales (IDEAM). Promedios climatológicos, 2010. URL <http://www.ideam.gov.co/web/tiempo-y-clima/clima?inheritRedirect=true>.
- [19] L. Lambrechts, C. Chevillon, R. G. Albright, B. Thaisomboonsuk, J. H. Richardson, R. G. Jarman, and T. W. Scott. Genetic specificity and potential for local adaptation between dengue viruses and mosquito vectors. *BMC Evolutionary Biology*, 9:160–171, 2009.
- [20] J. Liu-Helmersson, H. Stenlund, A. Wilder-Smith, and J. Rocklöv. Vectorial capacity of *Aedes aegypti*: Effects of temperature and implications for global dengue epidemic potential. *PLoS ONE*, 9(3), 2014. ISSN 19326203.
- [21] D. P. Lizarralde-Bejarano, S. Arboleda-Sánchez, and M. E. Puerta-Yepes. Understanding epidemics from mathematical models: Details of the 2010 dengue epidemic in Bello (Antioquia, Colombia). *Applied Mathematical Modelling*, 43: 566–578, 2017. ISSN 0307904X. doi: 10.1016/j.apm.2016.11.022.
- [22] M. Loza-Murguía and F. Noireau. Vectorial capacity of *Triatoma guasayana* (Wygodzinsky & Abalos) (Hemiptera: Reduviidae) compared with two other

- species of epidemic importance. *Neotropical Entomology*, 39(5):799–809, oct 2010. doi: 10.1590/s1519-566x2010000500020. URL <https://doi.org/10.1590/s1519-566x2010000500020>.
- [23] G. Macdonald. The theory of the eradication of malaria. 1956. URL <http://apps.who.int/iris/handle/10665/64524>.
- [24] E. Massad and F. A. B. Coutinho. Vectorial capacity, basic reproduction number, force of infection and all that: Formal notation to complete and adjust their classical concepts and equations. *Memorias do Instituto Oswaldo Cruz*, 107(4): 564–567, 2012.
- [25] E. Massad, S. Ma, M. N. Burattini, Y. Tun, F. A. B. Coutinho, and L. W. Ang. The risk of chikungunya fever in a dengue-endemic area. *Journal of Travel Medicine*, 15(3):147–155, 2008.
- [26] H. McCallum. *Population parameters: estimation for ecological models*. 2000. ISBN 9780865427402. doi: 10.1002/9780470757468.
- [27] L. A. Moreira, I. Iturbe-Ormaetxe, J. A. Jeffery, G. Lu, A. T. Pyke, L. M. Hedges, B. C. Rocha, S. Hall-Mendelin, A. Day, M. Riegler, L. E. Hugo, K. N. Johnson, B. H. Kay, E. A. McGraw, A. F. van den Hurk, P. A. Ryan, and S. L. O'Neill. A wolbachia symbiont in aedes aegypti limits infection with dengue, chikungunya, and plasmodium. *Cell*, 139(7):1268–1278, Dec. 2009. doi: 10.1016/j.cell.2009.11.042. URL <https://doi.org/10.1016/j.cell.2009.11.042>.
- [28] P. Nouvellet, E. Dumonteil, and S. Gourbière. The Improbable Transmission of Trypanosoma cruzi to Human: The Missing Link in the Dynamics and Control of Chagas Disease. *PLoS Neglected Tropical Diseases*, 7(11):e2505, 2013.
- [29] V. N. Novoseltsev, A. I. Michalski, J. A. Novoseltseva, A. I. Yashin, J. M. Carey, and A. M. Ellis. An age-structured extension to the vectorial capacity model. *Plos one*, 7:e39479, 2012.
- [30] K. P. Paaajmans, L. J. Cator, and M. B. Thomas. Temperature-dependent pre-bloodmeal period and temperature-driven asynchrony between parasite development and mosquito biting rate reduce malaria transmission intensity. *Plos one*, 8:e55777, 2013.
- [31] A. Pandey, A. Mubayi, and J. Medlock. Comparing vector–host and SIR models for dengue transmission. *Mathematical Biosciences*, 246(2):252–259, Dec. 2013. doi: 10.1016/j.mbs.2013.10.007. URL <https://doi.org/10.1016/j.mbs.2013.10.007>.
- [32] G. F. Paz, M. F. B. Ribeiro, É. M. Michalsky, A. C. V. M. da Rocha Lima, J. C. França-Silva, R. A. Barata, C. L. Fortes-Dias, and E. S. Dias. Evaluation of the vectorial capacity of rhipicephalus sanguineus (acari: Ixodidae) in the transmission of canine visceral leishmaniasis. *Parasitology Research*, 106(2):523–528, dec 2009. doi: 10.1007/s00436-009-1697-1. URL <https://doi.org/10.1007/s00436-009-1697-1>.
- [33] O. S. Raigosa, E. A. Bermúdez, and A. M. Loaiza. A simulation model for the chikungunya with vectorial capacity. *Applied Mathematical Sciences*, 9 140:6953 – 6960, 2015.

- [34] R. C. Reiner, T. A. Perkins, C. M. Barker, T. Niu, L. F. Chaves, A. M. Ellis, D. B. George, A. L. Menach, J. R. C. Pulliam, D. Bisanzio, C. Buckee, C. Chiyaka, D. A. T. Cummings, A. J. Garcia, M. L. Gatton, P. W. Gething, D. M. Hartley, G. Johnston, E. Y. Klein, E. Michael, S. W. Lindsay, A. L. Lloyd, D. M. Pigott, W. K. Reisen, N. Ruktanonchai, B. K. Singh, A. J. Tatem, U. Kitron, S. I. Hay, T. W. Scott, and D. L. Smith. A systematic review of mathematical models of mosquito-borne pathogen transmission: 1970-2010. *Journal of The Royal Society Interface*, 10(81):20120921–20120921, 2013. ISSN 1742-5689. doi: 10.1098/rsif.2012.0921.
- [35] R. Ross. Some a priori pathometric equations. *The British Medical Journal*, pages 546–547, 1915. ISSN 0959-8138. doi: 10.1136/bmj.1.2830.546.
- [36] M. I. Salazar, J. H. Richardson, I. Sánchez-Vargas, K. E. Olson, and B. J. Beaty. *BMC Microbiology*, 7(1):9, 2007. doi: 10.1186/1471-2180-7-9. URL <https://doi.org/10.1186/1471-2180-7-9>.
- [37] A. Saltelli. Making best use of model evaluations to compute sensitivity indices. *Computer physics communications*, 145:280–297, 2002.
- [38] A. Saltelli, M. Ratto, T. Andres, F. Campolongo, J. Cariboni, D. Gatelli, M. Saisana, and S. Tarantola. *Global sensitivity analysis: The primer*. John Wiley & Sons, Chichester, England, 2008.
- [39] A. Saltelli, P. Annoni, I. Azzini, F. Campolongo, M. Ratto, and S. Tarantola. Variance based sensitivity analysis of model output. Design and estimator for the total sensitivity index. *Computer Physics Communications*, 181(2):259–270, 2010.
- [40] C. Shidrawi and G. R. Garret-Jones. Malaria Vectorial Capacity of a Population of *Anopheles gambiae* An Exercise in Epidemiological Entomology. *Bull. Wld Hith Org*, 40:531–545, 1969.
- [41] D. L. Smith and F. E. McKenzie. Statics and dynamics of malaria infection in *Anopheles* mosquitoes. *Malaria Journal*, 3, 2004.
- [42] D. L. Smith, K. E. Battle, S. I. Hay, C. M. Barker, T. W. Scott, and F. E. McKenzie. Ross, Macdonald, and a theory for the dynamics and control of mosquito-transmitted pathogens. *PLoS Pathogens*, 8(4), 2012.
- [43] I. M. Sobol. On sensitivity estimation for nonlinear mathematical models. *Matematicheskoe modelirovanie*, 2:112–118, 1990.
- [44] L. M. Styer, J. R. Carey, and T. W. Scott. Mosquitoes do senesce: Departure from the paradigm of constant mortality. *Am J Trop Med Hyg*, 76:111–117, 2007.
- [45] L. T. Takahashi, N. A. Maidana, W. C. Ferreira, P. Pulino, and H. M. Yang. Mathematical models for the *Aedes aegypti* dispersal dynamics: Travelling waves by wing and wind. *Bulletin of Mathematical Biology*, 2005. ISSN 00928240. doi: 10.1016/j.bulm.2004.08.005.
- [46] J. R. Taylor. *An introduction to error analysis: the study of uncertainties in physical measurements*. John Wiley & Sons, Sausalito, California, 1997.
- [47] C. M. Vélez. Global sensitivity and uncertainty analysis (gsua) of dynamical systems using variance-based methods, 2015.

-
- [48] H. M. Yang, M. d. L. d. G. Macoris, K. C. Galvani, and M. T. M. Andrighetti. Follow up estimation of *Aedes aegypti* entomological parameters and mathematical modellings. *BioSystems*, 2011. ISSN 03032647. doi: 10.1016/j.biosystems.2010.11.002.
- [49] H. M. Yang, J. L. Boldrini, A. C. Fassoni, L. F. S. Freitas, M. C. Gomez, K. K. B. de Lima, V. R. Andrade, and A. R. R. Freitas. Fitting the incidence data from the city of campinas, brazil, based on dengue transmission modellings considering time-dependent entomological parameters. *PLOS ONE*, 11(3):e0152186, Mar. 2016. doi: 10.1371/journal.pone.0152186. URL <https://doi.org/10.1371/journal.pone.0152186>.
- [50] M. Zeller and J. C. Koella. Effects of food variability on growth and reproduction of *Aedes aegypti*. *Ecology and Evolution*, 6(2):552–559, jan 2016. doi: 10.1002/ece3.1888. URL <https://doi.org/10.1002/ece3.1888>.

Chapter 2

A novel algorithm for confidence sub-contour box estimation: An alternative to traditional confidence intervals

Submitted to	Status	DOI (if available)
Arxiv	Published	arXiv:1909.09603v3

2.1 Authors

Daniel Rojas-Díaz¹, Alexandra Catano-Lopez², Carlos M. Vélez-Sánchez²

¹ Dept. Ciencias Biológicas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

² Dept. Ciencias Matemáticas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

2.2 Abstract

The factor estimation process is a really challenging task for non-linear models. Even whether researchers manage to successfully estimate model factors, they still must estimate their confidence intervals, which could require a high computational cost to turn them into informative measures. Some methods in the literature attempt to estimate regions within the estimation search space where factors may jointly exist and fit the real data (confidence contours), however, its estimation process raises several issues as the number of factors increases. Hence, in this paper, we focus on the estimation of a subregion within the confidence contour that we called as Confidence Sub-contour Box (CSB). We proposed two main algorithms for CSB estimation, as well as its interpretation and validation. Given the way we estimated CSB, we expected and validated some useful properties of this new kind of confidence interval: a user-defined uncertainty level, asymmetrical intervals, sensitivity assessment related to the interval length for each factor, and the identification of true-influential factors.

Bibliography

- [1] G. E. B. Archer, A. Saltelli, and I. M. Sobol. Sensitivity measures, anova-like techniques and the use of bootstrap. *Statist. Comput. Simul.*, 58:99–120, 1997.
- [2] A. C. Babbie, P. Kirk, and M. P. H. Stumpf. Topological sensitivity analysis for systems biology. *Proceedings of the National Academy of Sciences*, 111(52):18507–18512, 2014. ISSN 0027-8424. doi: 10.1073/pnas.1414026112.
- [3] J. Barrios, A. Piétrus, G. Joya, A. Marrero, and H. de Arazoza. A differential inclusion approach for modeling and analysis of dynamical systems under uncertainty: Application to dengue disease transmission. *Soft Computing*, 17(2): 239–253, 2013. ISSN 14327643 14337479. doi: 10.1007/s00500-012-0889-2.
- [4] B. Cantó, C. Coll, and E. Sánchez. Estimation of parameters in a structured SIR model. *Advances in Difference Equations*, 2017(1), Jan. 2017. doi: 10.1186/s13662-017-1078-5. URL <https://doi.org/10.1186/s13662-017-1078-5>.
- [5] N. R. Draper and H. Smith. *Applied Regression Analysis*. Wiley and Sons, 1998. ISBN 0471170828. doi: 10.1198/tech.2005.s303.
- [6] V. Duong, L. Lambrechts, R. E. Paul, S. Ly, R. S. Lay, K. C. Long, R. Huy, A. Tarantola, T. W. Scott, A. Sakuntabhai, and P. Buchy. Asymptomatic humans transmit dengue virus to mosquitoes. *Proceedings of the National Academy of Sciences*, 112(47):14688–14693, Nov. 2015. doi: 10.1073/pnas.1508114112. URL <https://doi.org/10.1073/pnas.1508114112>.
- [7] L. Geris and D. Gomez-Cabrero. *Uncertainty in biology a computational modeling approach*. Springer international publishing Switzerland, 2016. ISBN 978-3-319-21295-1.
- [8] Instituto Nacional de Salud. Manual del usuario del software SIVIGILA, 2011. URL <https://goo.gl/j1DpX4>.
- [9] H. Miao, X. Xia, A. S. Perelson, and H. Wu. On identifiability of nonlinear ODE models and applications in viral dynamics. *SIAM Rev Soc Ind Appl Math.*, 53(1): 3–39, 2011.
- [10] A. Olsson, G. Sandberg, and O. Dahlblom. On latin hypercube sampling for structural reliability analysis. *Structural Safety*, 25(1):47–68, Jan. 2003. doi: 10.1016/s0167-4730(02)00039-5. URL [https://doi.org/10.1016/s0167-4730\(02\)00039-5](https://doi.org/10.1016/s0167-4730(02)00039-5).
- [11] Rojas-Díaz, Daniel and Vélez-Sánchez, Carlos Mario. drojasd/gsua-csb: Gsua-csb v1.0, 2019. URL <https://zenodo.org/record/3383316>.
- [12] A. Saltelli, M. Ratto, T. Andres, F. Campolongo, J. Cariboni, D. Gatelli, M. Saisana, and S. Tarantola. *Global sensitivity analysis: The primer*. John Wiley & Sons, Chichester, England, 2008. URL <http://www.wiley.com/WileyCDA/WileyTitle/productCd-0470059974.html><http://www.amazon.es/Global-Sensitivity-Analysis-The-Primer/dp/0470059974><https://docs.google.com/file/d/0BwXJdRHn-Rd6TS1LUDhnQjJtble/edit><https://www.researchgate.net/publication/25332810>.

- [13] A. Saltelli, P. Annoni, I. Azzini, F. Campolongo, M. Ratto, and S. Tarantola. Variance based sensitivity analysis of model output. Design and estimator for the total sensitivity index. *Computer physics communications.*, 181:259–270, 2010.
- [14] G. A. F. Seber and C. J. Wild. *Nonlinear regression*. John Wiley & Sons, New Jersey, USA, 2003.
- [15] L. F. Shampine and M. W. Reichelt. The MATLAB ODE suite. *SIAM Journal on Scientific Computing*, 18(1):1–22, Jan. 1997. doi: 10.1137/s1064827594276424. URL <https://doi.org/10.1137/s1064827594276424>.
- [16] I. M. Sobol. On sensitivity estimation for nonlinear mathematical models. *Matem. Mod.*, 2:112–118, 1990.
- [17] I. M. Sobol. Global sensitivity indices for nonlinear mathematical models and their Monte Carlo estimates. *Mathematics and computers in simulation*, 55:271–280, 2001.
- [18] L. T. Takahashi, N. A. Maidana, W. C. Ferreira, P. Pulino, and H. M. Yang. Mathematical models for the *Aedes aegypti* dispersal dynamics: Travelling waves by wing and wind. *Bulletin of Mathematical Biology*, 2005. ISSN 00928240. doi: 10.1016/j.bulm.2004.08.005.
- [19] N. T. Toan, S. Rossi, G. Prisco, N. Nante, and S. Viviani. Dengue epidemiology in selected endemic countries: factors influencing expansion factors as estimates of underreporting. *Tropical Medicine & International Health*, 20(7):840–863, mar 2015.
- [20] N. Tuncer, M. Marctheva, B. LaBarre, and S. Payout. Structural and practical identifiability analysis of zika epidemiological models. *Bulletin of Mathematical Biology*, 80(8):2209–2241, jun 2018. doi: 10.1007/s11538-018-0453-z. URL <https://doi.org/10.1007/s11538-018-0453-z>.
- [21] H. U. Voss, J. Timmer, and J. Kurths. Nonlinear dynamical system identification from uncertain and indirect measurements. *International Journal of Bifurcation and Chaos*, 14(6):1905–1933, 2004.
- [22] K. W. Vugrin, L. P. Swiler, R. M. Roberts, N. J. Stucky-Mack, and S. P. Sullivan. Confidence region estimation techniques for nonlinear regression in groundwater flow: Three case studies. *Water Resources Research*, 43(3), Mar. 2007. doi: 10.1029/2005wr004804. URL <https://doi.org/10.1029/2005wr004804>.
- [23] S. Xiao, Z. Lu, and P. Wang. Multivariate global sensitivity analysis based on distance components decomposition. *Risk Analysis*, 38(12):2703–2721, July 2018. doi: 10.1111/risa.13133. URL <https://doi.org/10.1111/risa.13133>.
- [24] H. M. Yang, M. d. L. d. G. Macoris, K. C. Galvani, and M. T. M. Andrighetti. Follow up estimation of *Aedes aegypti* entomological parameters and mathematical modellings. *BioSystems*, 2011. ISSN 03032647. doi: 10.1016/j.biosystems.2010.11.002.

Chapter 3

Global Sensitivity and Uncertainty Analyses - Confidence Subcontour Box (GSUA-CSB) Toolbox

Submitted to	Status	DOI (if available)
Matlab file exchange	Published	10.5755281/zenodo.3383316

3.1 Authors

Daniel Rojas-Díaz¹, Carlos M. Vélez-Sánchez²

¹ Dept. Ciencias Biológicas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

² Dept. Ciencias Matemáticas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

3.2 Abstract

Global Sensitivity and Uncertainty Analyses - Confidence Subcontour Box (GSUA-CSB) Toolbox is a product developed by Universidad EAFIT for command-line mathematical model validation in both of Simulink or Symbolic Math Toolbox environment. At present, the toolbox allows performing the following functions: To apply and visualize several variance-based sensitivity (SA) and uncertainty (UA) analysis, to estimate model parameters (PE) and to estimate the confidence sub-contour box (CSB) of previously estimated parameters. This toolbox has its basis on the previous work of Carlos Mario Vélez: GSUA of dynamical systems using variance-based methods, published in this [mathworks file exchange link](#).

Bibliography

- [1] A. Saltelli, P. Annoni, I. Azzini, F. Campolongo, M. Ratto, and S. Tarantola. Variance based sensitivity analysis of model output. Design and estimator for the total sensitivity index. *Computer physics communications.*, 181:259–270, 2010.

- [2] S. Xiao, Z. Lu, and P. Wang. Multivariate global sensitivity analysis based on distance components decomposition. *Risk Analysis*, 38(12):2703–2721, July 2018. doi: 10.1111/risa.13133. URL <https://doi.org/10.1111/risa.13133>.

Chapter 4

Influence of pulse-type inputs on parameter estimation and chemical control assessment in a dengue deterministic model

Submitted to	Status	DOI (if available)
Plos Computational Biology	Under revision	Not yet

4.1 Authors

Alexandra Catano-López¹, Daniel Rojas-Díaz², Carlos M. Vélez Sánchez¹

¹ Dept. Ciencias Matemáticas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

² Dept. Ciencias Biológicas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

4.2 Abstract

Many deterministic mathematical models of dengue spread usually use theoretical or measured-in-lab parameters, without worrying about the uncertainties in factors, such as quality and confidence of data, or changes in control inputs. Other models consider environmental conditions and apply parameter estimations to obtain experimental parameters. This is more in accordance with real situations and may even introduce climate changes as periodical continuous-time inputs; however, they do not include the estimations of the effects of aperiodic changes in inputs, such as chemical control (fumigation) of vectors. This study estimates parameters (including chemical control parameters) and confidence contours (a type of confidence interval) under uncertainty conditions using Matlab tools and data from the municipality of Bello (Colombia) during 2010–2014. Our study shows that introducing aperiodic pulse-type inputs into the mathematical model allows us to (i) estimate feasible parameters into confidence biological intervals, (ii) highlight the importance of chemical control as a method to control disease propagation, and (iii) reproduce

the endemic behavior. We obtained a model with new and verifiable biological parameters, described a methodology and a novel Matlab toolbox for parameter and confidence interval estimation under uncertainties, and performed reliable simulations showing the behavior of dengue spread in different interesting scenarios.

Bibliography

- [1] Mudin RN. Dengue incidence and the prevention and control program in Malaysia. *International Medical Journal Malaysia*. 2015;14(1):05–10.
- [2] Gubler DJ. Resurgent vector-borne diseases as a global health problem. *Emerging Infectious Diseases*. 1998;4(3):442–450. doi:10.3201/eid0403.980326.
- [3] Kraemer MUG, Sinka ME, Duda KA, Mylne A, Shearer FM, Brady OJ, et al. The global compendium of *Aedes aegypti* and *Ae. albopictus* occurrence. *Scientific Data*. 2015;doi:10.1038/sdata.2015.35.
- [4] Wilder-Smith A, Chen LH, Massad E, Wilson ME. Threat of dengue to blood safety in dengue-endemic countries; 2009.
- [5] World Health Organization. Dengue control; 2019. Available from: <https://www.who.int/denguecontrol/disease/en/> [cited 2019-09].
- [6] Burattini MN, Chen M, Chow A, Coutinho FAB, Goh KT, Lopez LF, et al. Modelling the control strategies against dengue in Singapore. *Epidemiology and Infection*. 2007;136(3):309–319. doi:10.1017/s0950268807008667.
- [7] Chávez JP, Götz T, Siegmund S, Wijaya KP. An SIR-Dengue transmission model with seasonal effects and impulsive control. *Mathematical Biosciences*. 2017;289:29–39. doi:10.1016/j.mbs.2017.04.005.
- [8] Bustamam A, Aldila D, Yuwanda A. Understanding Dengue Control for Short- and Long-Term Intervention with a Mathematical Model Approach. *Journal of Applied Mathematics*. 2018;2018:1–13. doi:10.1155/2018/9674138.
- [9] Wickramaarachchi WPTM, Perera SSN, Jayasinghe S. Modelling and analysis of dengue disease transmission in urban Colombo: A wavelets and cross wavelets approach. *Journal of the National Science Foundation of Sri Lanka*. 2016;43(4):337–345. doi:10.4038/jnsfsr.v43i4.7968.
- [10] Andraud M, Hens N, Marais C, Beutels P. Dynamic epidemiological models for dengue transmission: a systematic review of structural approaches. *PloS one*. 2012;43(4). doi:10.1371/journal.pone.0049085.
- [11] Braselton IBJ. A Survey of Mathematical Models of Dengue Fever. *Journal of Computer Science & Systems Biology*. 2015;08(05). doi:10.4172/jcsb.1000198.
- [12] Reiner RC, Perkins TA, Barker CM, Niu T, Chaves LF, Ellis AM, et al. A systematic review of mathematical models of mosquito-borne pathogen transmission: 1970-2010. *Journal of The Royal Society Interface*. 2013;10(81):921. doi:10.1098/rsif.2012.0921.
- [13] Eisenberg JNS, Brookhart MA, Rice G, Brown M, Colford JM. Disease transmission models for public health decision making: Analysis of epidemic and endemic conditions caused by waterborne pathogens. *Environmental Health Perspectives*. 2002;110(8):783–790. doi:10.1289/ehp.02110783.

- [14] World Health Organization. Dengue guidelines for diagnosis, treatment, prevention and control. World Health Organization; 2009.
- [15] Ellis AM, Morrison AC, Garcia AJ, Scott TW, Focks DA. Parameterization and Sensitivity Analysis of a Complex Simulation Model for Mosquito Population Dynamics, Dengue Transmission, and Their Control. *The American Journal of Tropical Medicine and Hygiene*. 2011;85(2):257–264. doi:10.4269/ajtmh.2011.10-0516.
- [16] Cailly P, Tran A, Balenghien T, L'Ambert G, Toty C, Ezanno P. A climate-driven abundance model to assess mosquito control strategies. *Ecological Modelling*. 2012;227(2012):7–17. doi:10.1016/j.ecolmodel.2011.10.027.
- [17] Lana RM, Carneiro TGS, Honório NA, Codeço CT. Seasonal and nonseasonal dynamics of *Aedes aegypti* in Rio de Janeiro, Brazil: Fitting mathematical models to trap data. *Acta Tropica*. 2014;129:25–32.
- [18] Vugrin KW, Swiler LP, Roberts RM, Stucky-Mack NJ, Sullivan SP. Confidence region estimation techniques for nonlinear regression in groundwater flow: Three case studies. *Water Resources Research*. 2007;43(3). doi:10.1029/2005wr004804.
- [19] Draper NR, Smith H. *Applied Regression Analysis*. Wiley-Interscience Publication; 1998.
- [20] Saltelli A, Aleksankina K, Becker W, Fennell P, Ferretti F, Holst N, et al. Why so many published sensitivity analyses are false: A systematic review of sensitivity analysis practices. *Environmental Modelling and Software*. 2019;114:29–39. doi:10.1016/j.envsoft.2019.01.012.
- [21] IDEAM. Promedios Climatológicos 1981–2010; 2016. Available from: <https://bit.ly/2kyN8p0> [cited 2019-09].
- [22] Departamento Administrativo Nacional de Estadística (DANE). Boletín: Censo General 2005; 2011. Available from: <https://goo.gl/JSWVRr>.
- [23] Instituto Nacional de Salud. Manual del usuario del software SIVIGILA; 2010. Available from: <https://goo.gl/j1DpX4> [cited 2019-07].
- [24] World Health Organization. Indoor residual spraying: An operational manual for IRS for malaria transmission, control and elimination. 2nd ed. World Health Organization; 2015.
- [25] James Braselton IB. A Survey of Mathematical Models of Dengue Fever. *Journal of Computer Science & Systems Biology*. 2015;8:255–267. doi:10.4172/jcsb.1000198.
- [26] Yang HM, Ferreira CP. Assessing the effects of vector control on dengue transmission. *Applied Mathematics and Computation*. 2008;198(1):401–413. doi:10.1016/j.amc.2007.08.046.
- [27] Joshi V, Mourya DT, Sharma RC. Persistence of dengue-3 virus through transovarial transmission passage in successive generations of *Aedes aegypti* mosquitoes. *American Journal of Tropical Medicine and Hygiene*. 2002;67(2):158–161. doi:10.4269/ajtmh.2002.67.158.

- [28] Maciel-de Freitas R, Koella JC, Lourenço-de Oliveira R. Lower survival rate, longevity and fecundity of *Aedes aegypti* (Diptera: Culicidae) females orally challenged with dengue virus serotype 2. *Transactions of the Royal Society of Tropical Medicine and Hygiene*. 2011;105(8):452. doi:10.1016/j.trstmh.2011.05.006.
- [29] Tun-Lin W, Burkot TR, Kay BH. Effects of temperature and larval diet on development rates and survival of the dengue vector *Aedes aegypti* in north Queensland, Australia. *Medical and Veterinary Entomology*. 2000;14(1). doi:10.1046/j.1365-2915.2000.00207.x.
- [30] Taylor JR. *An Introduction to Error Analysis: The Study of Uncertainties in Physical Measurements*. 2nd ed. Sausalito, California: University Science Books; 1997.
- [31] Ljung L. *System Identification: Theory for the User*. 2nd ed. Prentice Hall; 1999.
- [32] Mathworks. *MATLAB Optimization Toolbox*; 2018b. Available from: <https://la.mathworks.com/products/optimization.html> [cited 2019-07].
- [33] Rousseeuw PJ, Hubert M. Robust statistics for outlier detection. *Wiley Interdisciplinary Reviews: Data Mining and Knowledge Discovery*. 2011;1:73–79. doi:10.1002/widm.2.
- [34] David HA, Nagaraja HN. *Order statistics*. Wiley Online Library. 1970;doi:10.1002/0471667196.ess6023.pub2.
- [35] Rojas-Díaz, Daniel and Vélez-Sánchez, Carlos Mario. drojasd/GSUA-CSB: GSUA-CSB v1.0; 2019. Available from: <https://zenodo.org/record/3383316> [cited 2019-09].
- [36] Rojas-Diaz D, Catano-Lopez A, Velez-Sanchez CM. Novel algorithm for confidence sub-contour box estimation: an alternative to traditional confidence intervals; 2019.
- [37] Saltelli A, Annoni P, Azzini I, Campolongo F, Ratto M, Tarantola S. Variance based sensitivity analysis of model output. Design and estimator for the total sensitivity index. *Computer Physics Communications*. 2010;181(2):259–270. doi:10.1016/j.cpc.2009.09.018.
- [38] Xiao S, Lu Z, Wang P. Multivariate Global Sensitivity Analysis Based on Distance Components Decomposition. *Risk Analysis*. 2018;38(12):2703–2721. doi:10.1111/risa.13133.
- [39] Yang HM, Macoris MdLdG, Galvani KC, Andrighetti MTM. Follow up estimation of *Aedes aegypti* entomological parameters and mathematical modellings. *BioSystems*. 2011;103(3):360–371. doi:10.1016/j.biosystems.2010.11.002.
- [40] Marinho RA, Beserra EB, Bezerra-Gusmão MA, Porto VdS, Olinda RA, dos Santos CAC. Effects of temperature on the life cycle, expansion, and dispersion of *Aedes aegypti* (Diptera: Culicidae) in three cities in Paraíba, Brazil. *Journal of Vector Ecology*. 2016;41(1):1–10. doi:10.1111/jvec.12187.
- [41] Lizarralde-Bejarano DP, Arboleda-Sánchez S, Puerta-Yepes ME. Understanding epidemics from mathematical models: Details of the 2010 dengue epidemic in Bello (Antioquia, Colombia). *Applied Mathematical Modelling*. 2017;43:566–578. doi:10.1016/j.apm.2016.11.022.

-
- [42] Ljung L. Characterization of the Concept of 'Persistently Exciting' in the Frequency Domain. Department of Automatic Control, Lund Institute of Technology (LTH); 1971.
- [43] Hickey WA, Craig GB. Genetic distortion of sex ratio in a mosquito, *Aedes aegypti*. *Genetics*. 1966;53(6):1177–1196.
- [44] Wood RJ. Between-family variation in sex ratio in the Trinidad (T-30) strain of *Aedes aegypti* (L.) indicating differences in sensitivity to the meiotic drive gene MD. *Genetica*. 1976;46(3):345–361. doi:10.1007/BF00055477.
- [45] Luz PM, Struchiner CJ, Galvani AP. Modeling Transmission Dynamics and Control of Vector-Borne Neglected Tropical Diseases. *PLoS Neglected Tropical Diseases*. 2010;4(10):e761. doi:10.1371/journal.pntd.0000761.
- [46] Red de biología matemática. *Epidemiología matemática*; 2018. Available from: <http://epidemiologia.eafit.edu.co/> [cited 2019-09].

Chapter 5

Sensitivity, uncertainty and identifiability analyses to define a dengue transmission model with real data of an endemic municipality of Colombia

Submitted to	Status	DOI (if available)
Plos Biology	Under revision	Not yet

5.1 Authors

Diana Paola Lizarralde-Bejarano¹, Daniel Rojas-Díaz², Sair Arboleda-Sánchez³, María Eugenia Puerta-Yepes¹

¹ Dept. Ciencias Matemáticas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

² Dept. Ciencias Biológicas/Escuela de ciencias, Universidad EAFIT, Medellín, Antioquia, Colombia

³ Grupo de Biología y Control de Enfermedades Infecciosas-BCEI, Universidad de Antioquia, Medellín, Antioquia, Colombia

5.2 Abstract

Dengue disease is a major problem of public health surveillance entities in tropical and subtropical regions which makes a significant impact not only epidemiological but social and economical. There are many factors involved in the dengue transmission process. We can evaluate the importance of these factors through the formulation of mathematical models. However, the majority of these models in the literature tend to be overparameterized, with considerable uncertainty levels and over complex formulations. We aim to evaluate the structure, complexity, trustworthiness, and suitability of three models, which simulate the transmission of dengue disease, through different strategies. To achieve this goal, we perform structural and practical identifiability, sensitivity and uncertainty analyses to these models. The

results showed that the most simple model was more appropriate and reliable than the other two to simulate the dengue transmission disease when the only available information to fit them is the cumulative number of reported dengue cases in an endemic municipality of Colombia.

Bibliography

- [1] OMS. Dengue y dengue grave; 2016. Available from: <http://www.who.int/mediacentre/factsheets/fs117/es/>.
- [2] Organization WH. Dengue; 2009.
- [3] Hernández LM, Durán DF, Buitrago DA, Garnica CA, Gómez LF, Bados DM, et al. Epidemiology and geo-referencing of the dengue fever in a hospital of second level in Colombia, 2010–2014. *Journal of infection and public health*. 2018;11(4):558–565.
- [4] Lee JS, Lim JK, Dang DA, Nguyen THA, Farlow A. Dengue vaccine supplies under endemic and epidemic conditions in three dengue-endemic countries: Colombia, Thailand, and Vietnam. *Vaccine*. 2017;35(50):6957–6966.
- [5] Hethcote HW. The basic epidemiology models: models, expressions for R_0 , parameter estimation, and applications. In: *Mathematical understanding of infectious disease dynamics*. World Scientific; 2009. p. 1–61.
- [6] Esteva L, Vargas C. A model for dengue disease with variable human population. *Journal of Mathematical Biology*. 1999;38(3):220–240.
- [7] Yang HM, Ferreira CP. Assessing the effects of vector control on dengue transmission. *Applied Mathematics and Computation*. 2008;198(1):401–413.
- [8] Esteva L, Vargas C. Coexistence of different serotypes of dengue virus. *Journal of Mathematical Biology*. 2003;46(1):31–47.
- [9] Andraud M, Hens N, Marais C, Beutels P. Dynamic epidemiological models for dengue transmission: a systematic review of structural approaches. *PloS one*. 2012;7:e49085.
- [10] Reiner RC, Perkins TA, Barker CM, Niu T, Chaves LF, Ellis AM, et al. A systematic review of mathematical models of mosquito-borne pathogen transmission: 1970–2010. *Journal of The Royal Society Interface*. 2013;10(81):20120921.
- [11] Heesterbeek J, Dietz K. The concept of R_0 in epidemic theory. *Statistica Neerlandica*. 1996;50(1):89–110.
- [12] Tuncer N, Marctheva M, LaBarre B, Payoute S. Structural and practical identifiability analysis of Zika epidemiological models. *Bulletin of mathematical biology*. 2018;80(8):2209–2241.
- [13] Miao H, Xia X, Perelson AS, Wu H. On identifiability of nonlinear ODE models and applications in viral dynamics. *SIAM review*. 2011;53(1):3–39.
- [14] Chis OT, Banga JR, Balsa-Canto E. Structural identifiability of systems biology models: a critical comparison of methods. *PloS one*. 2011;6(11):e27755.

- [15] Bellu G, Saccomani MP, Audoly S, D'Angiò L. DAISY: A new software tool to test global identifiability of biological and physiological systems. *Computer Methods and Programs in Biomedicine*. 2007;88(1):52–61. doi:10.1016/j.cmpb.2007.07.002.
- [16] Chiş O, Banga JR, Balsa-Canto E. GenSSI: A software toolbox for structural identifiability analysis of biological models. *Bioinformatics*. 2011;27(18):2610–2611. doi:10.1093/bioinformatics/btr431.
- [17] Karlsson J, Anguelova M, Jirstrand M. An efficient method for structural identifiability analysis of large dynamic systems. *IFAC Proceedings Volumes*. 2012;45(16):941–946.
- [18] Meshkat N, Er-zhen Kuo C, DiStefano J. On finding and using identifiable parameter combinations in nonlinear dynamic systems biology models and combos: A novel web implementation. *PLoS ONE*. 2014;9(10). doi:10.1371/journal.pone.0110261.
- [19] Villaverde AF, Barreiro A, Papachristodoulou A. Structural Identifiability of Dynamic Systems Biology Models. *PLoS Computational Biology*. 2016;12(10):1–22. doi:10.1371/journal.pcbi.1005153.
- [20] Saltelli A, Ratto M, Andres T, Campolongo F, Cariboni J, Gatelli D, et al. *Global sensitivity analysis: The primer*. Chichester, England: John Wiley & Sons; 2008.
- [21] Cacuci DG. *Sensitivity & Uncertainty Analysis, Volume 1: Theory*. Boca Raton: Chapman and Hall/CRC; 2003.
- [22] Turányi T, Tomlin AS. In: *Sensitivity and Uncertainty Analyses*. Springer Berlin Heidelberg; 2014. p. 61–144.
- [23] Babbie AC, Kirk P, Stumpf MPH. Topological sensitivity analysis for systems biology. *Proceedings of the National Academy of Sciences*. 2014;111(52):18507–18512. doi:10.1073/pnas.1414026112.
- [24] Saltelli A, Annoni P, Azzini I, Campolongo F, Ratto M, Tarantola S. Variance based sensitivity analysis of model output. Design and estimator for the total sensitivity index. *Computer physics communications*. 2010;181:259–270.
- [25] Esteva L, Vargas C. Analysis of a dengue disease transmission model. *Mathematical biosciences*. 1998;150(2):131–151.
- [26] Heesterbeek JAP. A brief history of R_0 and a recipe for its calculation. *Acta biotheoretica*. 2002;50(3):189–204.
- [27] Diekmann O, Heesterbeek J, Roberts MG. The construction of next-generation matrices for compartmental epidemic models. *Journal of the Royal Society Interface*. 2009;7(47):873–885.
- [28] Diekmann O, Heesterbeek JAP, Metz JA. On the definition and the computation of the basic reproduction ratio R_0 in models for infectious diseases in heterogeneous populations. *Journal of mathematical biology*. 1990;28(4):365–382.
- [29] Pérez-Restrepo LS, Triana-Chávez O, Mejía-Jaramillo AM, Arboleda-Sánchez SO. Vector competence analysis of two *Aedes aegypti* lineages from Bello, Colombia, reveals that they are affected similarly by dengue-2 virus infection. *Archives of virology*. 2019;164(1):149–158.

- [30] Lizarralde-Bejarano DP, Arboleda-Sánchez S, Puerta-Yepes ME. Understanding epidemics from mathematical models: Details of the 2010 dengue epidemic in Bello (Antioquia, Colombia). *Applied Mathematical Modelling*. 2017;43:566–578.
- [31] MATLAB Symbolic Math Toolbox; 2019.
- [32] Rojas-Díaz, Daniel and Vélez-Sánchez, Carlos Mario. drojasd/GSUA-CSB: GSUA-CSB v1.0; 2019. Available from: <https://zenodo.org/record/3383316>.
- [33] Shampine LF, Reichelt MW. The MATLAB ODE Suite. *SIAM Journal on Scientific Computing*. 1997;18(1):1–22. doi:10.1137/s1064827594276424.
- [34] Draper NR, Smith H. *Applied Regression Analysis*. Wiley and Sons; 1998.
- [35] Olsson A, Sandberg G, Dahlblom O. On Latin hypercube sampling for structural reliability analysis. *Structural Safety*. 2003;25(1):47–68. doi:10.1016/s0167-4730(02)00039-5.
- [36] Archer GEB, Saltelli A, Sobol IM. Sensitivity measures, anova-like techniques and the use of bootstrap. *Statist Comput Simul*. 1997;58:99–120.
- [37] Xiao S, Lu Z, Wang P. Multivariate Global Sensitivity Analysis Based on Distance Components Decomposition. *Risk Analysis*. 2018;38(12):2703–2721. doi:10.1111/risa.13133.
- [38] Sobol IM. On sensitivity estimation for nonlinear mathematical models. *Matem Mod*. 1990;2:112–118.
- [39] Sobol IM. Global sensitivity indices for nonlinear mathematical models and their Monte Carlo estimates. *Mathematics and computers in simulation*. 2001;55:271–280.
- [40] Gábor A, Villaverde AF, Banga JR. Parameter identifiability analysis and visualization in large-scale kinetic models of biosystems. *BMC Systems Biology*. 2017;11(1). doi:10.1186/s12918-017-0428-y.
- [41] Raue A, Kreutz C, Maiwald T, Bachmann J, Schilling M, Klingmüller U, et al. Structural and practical identifiability analysis of partially observed dynamical models by exploiting the profile likelihood. *Bioinformatics*. 2009;25(15):1923–1929. doi:10.1093/bioinformatics/btp358.
- [42] Tuncer N, Le TT. Structural and practical identifiability analysis of outbreak models. *Mathematical Biosciences*. 2018;299:1–18. doi:10.1016/j.mbs.2018.02.004.
- [43] Rousseeuw PJ, Hubert M. *Robust statistics for outlier detection*. Wiley Interdisciplinary Reviews: Data Mining and Knowledge Discovery. 2011;1(1):73–79.
- [44] Johansson MA, Hombach J, Cummings DAT. Models of the impact of dengue vaccines: A review of current research and potential approaches. *Vaccine*. 2011;29(35):5860–5868. doi:10.1016/j.vaccine.2011.06.042.
- [45] Tuncer N, Gulbudak H, Cannataro VL, Martcheva M. Structural and practical identifiability issues of immuno-epidemiological vector–host models with application to rift valley fever. *Bulletin of mathematical biology*. 2016;78(9):1796–1827.

-
- [46] Yashima K, Sasaki A. Spotting Epidemic Keystones by R0 Sensitivity Analysis: High-Risk Stations in the Tokyo Metropolitan Area. PLOS ONE. 2016;11(9):1–19. doi:10.1371/journal.pone.0162406.
- [47] Wang L, Zhao H, Oliva SM, Zhu H. Modeling the transmission and control of Zika in Brazil. Scientific reports. 2017;7(1):7721.
- [48] Martcheva M. An Introduction to Mathematical Epidemiology. Springer (US); 2015. Available from: <https://doi.org/10.1007/978-1-4899-7612-3>.
- [49] Chastaing G, Gamboa F, Prieur C. Generalized Hoeffding-Sobol decomposition for dependent variables - application to sensitivity analysis. Electronic Journal of Statistics. 2012;6(0):2420–2448. doi:10.1214/12-ejs749.