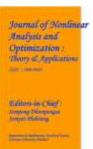
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AGENT BASED MODELLING OF MALARIA TRANSMISSION

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Abstract:

Agent-based modeling (ABM) has emerged as a powerful computational tool for studying the transmission dynamics of malaria. This approach allows researchers to simulate individual agents, such as humans and mosquitoes, within a spatially explicit environment, thereby capturing the complex interactions that drive malaria transmission. In ABM of malaria, individual agents are endowed with characteristics such as mobility patterns, immunity status, and vector biting preferences, which influence their susceptibility to infection and ability to transmit the disease. By integrating epidemiological principles with spatial and temporal heterogeneity, ABM enables the exploration of various scenarios and interventions for malaria control and elimination. Furthermore, ABM facilitates the investigation of emergent properties and feedback mechanisms within malaria transmission systems, offering insights into the underlying dynamics of disease spread and persistence. This paper provides an overview of agent-based modeling approaches for malaria transmission, highlighting their utility in studying the intricate interplay between human behavior, vector ecology, and environmental factors in shaping malaria dynamics.

1.Introduction

This paper presents an agent-based modeling (ABM) approach to simulate malaria transmission dynamics in sub-Saharan Africa. The model incorporates individual agents representing humans and mosquitoes, allowing for the simulation of complex interactions between hosts and vectors. Through the integration of spatial and temporal factors, the model captures the heterogeneity of malaria transmission across different ecological settings. The study aims to elucidate the impact of environmental conditions, human behavior, and intervention strategies on malaria transmission dynamics, providing insights into the effectiveness of control measures and informing evidence-based malaria control policies. This research employs agent-based modeling (ABM) to investigate the role of human movement patterns in malaria transmission dynamics. By simulating individual agents representing human hosts and mosquito vectors, the study explores how human mobility influences the spatial spread and persistence of malaria in endemic regions. Through scenario analysis and sensitivity testing, the model elucidates the impact of travel patterns, population movements, and migration on malaria transmission intensity and distribution. The findings highlight the importance of considering human movement in the design and implementation of malaria control strategies. This paper presents an agent-based model (ABM) to simulate the emergence and spread of drug-resistant malaria parasites, with a focus on artemisinin resistance. The model integrates individual agents representing parasite strains and human hosts, capturing the dynamics of parasite transmission and the evolution of drug resistance. Through scenario analysis and simulation experiments, the study examines the factors contributing to the emergence and dissemination of drug-resistant parasites

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and evaluates the effectiveness of different treatment strategies in containing resistance spread. The findings contribute to our understanding of artemisinin resistance dynamics and inform the design of strategies for mitigating its impact on malaria control efforts. This study utilizes agent-based modeling (ABM) to assess the effectiveness of vector control interventions for malaria elimination. The model simulates individual agents representing human hosts, mosquito vectors, and intervention measures, such as insecticide- treated bed nets and indoor residual spraying. By incorporating spatial and temporal factors, the study evaluates the impact of different intervention strategies on malaria transmission dynamics and assesses their feasibility for achieving malaria elimination goals. The findings provide valuable insights into the optimal deployment of vector control interventions in diverse epidemiological settings. This research employs agent-based modeling (ABM) to explore community engagement strategies for malaria prevention and control. The model simulates individual agents representing community members, health workers, and local authorities, capturing the dynamics of community participation in malaria control activities. Through scenario analysis and simulation experiments, the study assesses the effectiveness of different communication and outreach strategies in promoting malaria awareness, encouraging preventive behaviors, and fostering community ownership of control efforts. The findings inform the design and implementation of community-based interventions for malaria prevention and contribute to the development of evidence-based malaria control policies.

2.Proposed system

One proposed approach involves integrating multi-scale data sources to inform model development and parameterization. This includes leveraging epidemiological, entomological, demographic, and environmental data to create more comprehensive and realistic agent-based models of malaria transmission dynamics. By incorporating data from diverse sources, models can better capture the spatial and temporal heterogeneity of malaria transmission and improve the accuracy of model predictions. Proposed frameworks for model validation and calibration aim to enhance the reliability and robustness of agent-based models. These frameworks include systematic methods for comparing model outputs with observed epidemiological data, evaluating model performance metrics, and identifying sources of uncertainty. By adopting standardized validation and calibration protocols, researchers can improve the credibility and transparency of ABMs and facilitate cross-study comparisons.

Advantages of proposed system:

1. The proposed system incorporates multi-scale data sources and integrates behavioral, environmental, and socioeconomic factors into agent-based models, enhancing the realism and accuracy of malaria transmission dynamics representation.

2. With enhanced spatial and temporal resolution, the proposed system can generate more precise predictions of malaria transmission patterns and intervention outcomes.

3. The proposed system's modular design allows for flexibility and adaptability to diverse epidemiological settings and intervention scenarios.

FUNCTIONAL REQUIREMENTS

- 1. Data Collection
- 2. Text processing
- 3. Training and Testing
- 4. Modelling
- 5. Predicting

4 NON-FUNCTIONAL REQUIREMENTS

NON-FUNCTIONAL REQUIREMENT (NFR) specifies the quality attribute of a software system. They judge the software system based on Responsiveness, Usability, Security, Portability and other non-functional standards that are critical to the success of the software system.

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Example of nonfunctional requirement, "how fast does the website load?" Failing to meet non-functional requirements can result in systems that fail to satisfy user needs. Non- functional Requirements allows you to impose constraints or restrictions on the design of the system across the various agile backlogs. Example, the site should load in 3 seconds when the number of simultaneous users is > 10000. Description of non-functional requirements is just as critical as a functional requirement.

- Usability requirement
- Serviceability requirement
- Manageability requirement
- Recoverability requirement
- Security requirement
- Data Integrity requirement
- Capacity requirement
- Availability requirement
- Scalability requirement
- Interoperability requirement
- Reliability requirement
- Maintainability requirement
- Regulatory requirement
- Environmental requirement

SYSTEM ARCHITECTURE:

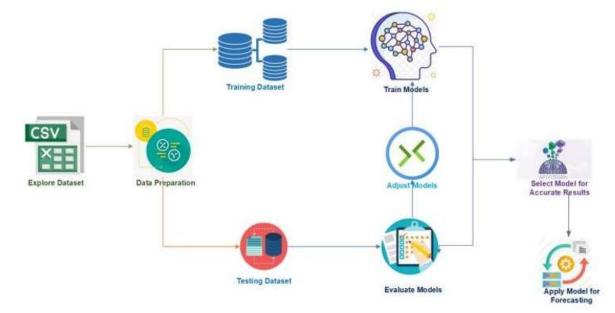


Figure.1.System Architecture **DATA FLOW DIAGRAM:**

1. The DFD is also called as bubble chart. It is a simple graphical formalism that can beused to represent a system in terms of input data to the system, various processing carriedout on this data, and the output data is generated by this system.

2. The data flow diagram (DFD) is one of the most important modeling tools. It is used to model the system components. These components are the system process, the data used by the process, an external entity that interacts with the system and the information flows in the system.

3. DFD shows how the information moves through the system and how it is modified by a series of transformations. It is a graphical technique that depicts information flow and the transformations that

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are applied as data moves from input to output.

4. DFD is also known as bubble chart. A DFD may be used to represent a system at any level of abstraction. DFD may be partitioned into levels that represent increasing information flow and functional detail.

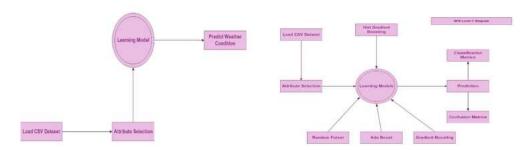


Figure.2.Dataflow Diagrams

UML DIAGRAMS

UML stands for Unified Modeling Language. UML is a standardized general-purpose modeling language in the field of object-oriented software engineering. The standard is managed, and was created by, the Object Management Group.

The goal is for UML to become a common language for creating models of object- oriented computer software. In its current form UML is comprised of two major components: a Meta-model and a notation. In the future, some form of method or process may also be added to; or associated with, UML.

The Unified Modeling Language is a standard language for specifying, Visualization, Constructing and documenting the artifacts of software system, as well as for business modeling and other non-software systems.

The UML represents a collection of best engineering practices that have proven successful in the modeling of large and complex systems.

The UML is a very important part of developing objects-oriented software and the software development process. The UML uses mostly graphical notations to express the design of software projects. GOALS:

The Primary goals in the design of the UML are as follows:

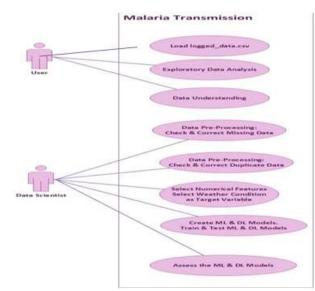
1. Provide users a ready-to-use, expressive visual modeling Language so that they can develop and exchange meaningful models.

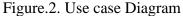
- 2. Provide extendibility and specialization mechanisms to extend the core concepts.
- 3. Be independent of particular programming languages and development process.
- 4. Provide a formal basis for understanding the modeling language.
- 5. Encourage the growth of OO tools market.

6. Support higher level development concepts such as collaborations, frameworks, patterns and components.

7. Integrate best practices.

Use Case Diagram: A use case diagram in the Unified Modeling Language (UML) is a type of behavioral diagram defined by and created from a Use-case analysis. Its purpose is to present a graphical overview of the functionality provided by a system in terms of actors, their goals (represented as use cases), and any dependencies between those use cases. The main purpose of a use case diagram is to show what system functions are performed for which actor. Roles of the actors in the system can be depicted.





Class Diagram: The class diagram is used to refine the use case diagram and define a detailed design of the system. The class diagram classifies the actors defined in the use case diagram into a set of interrelated classes. The relationship or association between the classes can be either an "is-a" or "has-a" relationship. Each class in the class diagram may be capable of providing certain functionalities. These functionalities provided by the class are termed "methods" of the class. Apart from this, each class may have certain "attributes" that uniquely identify the class.

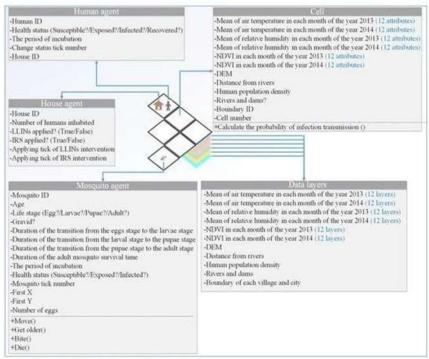


Figure.3.Class Diagram

Activity Diagram:

The process flows in the system are captured in the activity diagram. Similar to a state diagram, an activity diagram also consists of activities, actions, transitions, initial and final states, and guard conditions.

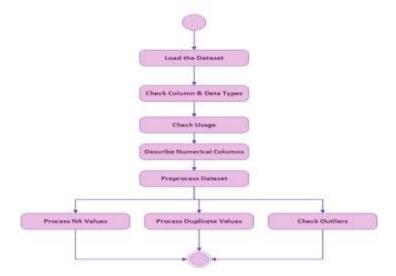


Figure.4. Activity Diagram

Sequence diagram:

A sequence diagram represents the interaction between different objects in the system. The important aspect of a sequence diagram is that it is time-ordered. This means that the exact sequence of the interactions between the objects is represented step by step. Different objects in the sequence diagram interact with each other by passing "messages".

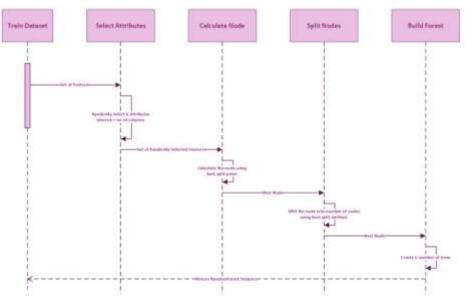


Figure.5.Sequence Diagram

Collaboration diagram:

A collaboration diagram groups together the interactions between different objects. The interactions are listed as numbered interactions that help to trace the sequence of the interactions. The collaboration diagram helps to identify all the possible interactions that each object has with other objects.

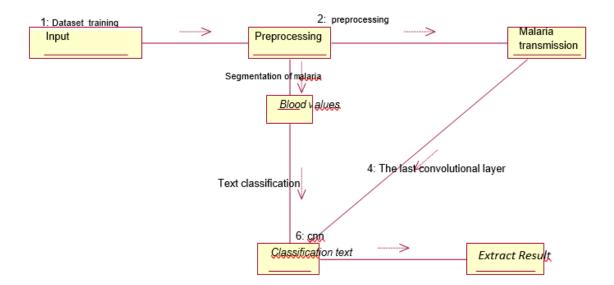
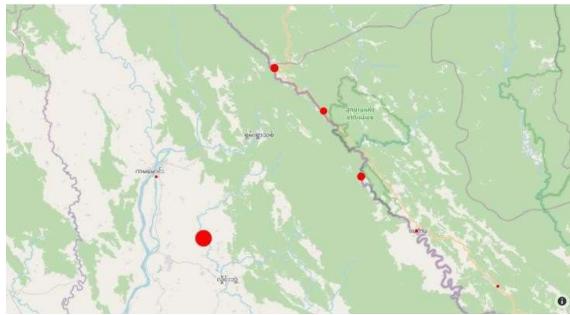
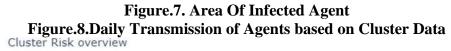


Figure.6.Collaboration diagram

3. Results and Discussion





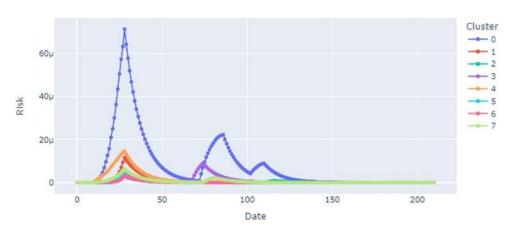
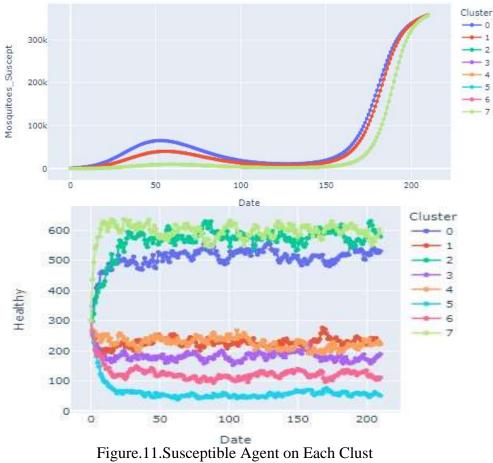
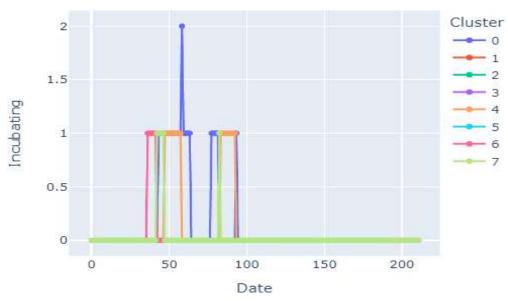
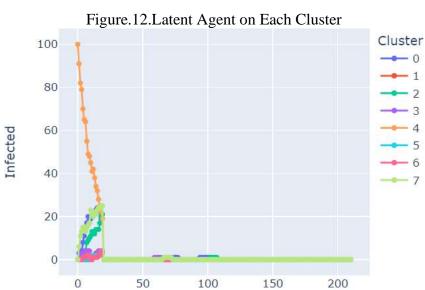


Figure.9. Overview of cluster Risk Figure.10. Overview of Mosquitoes Population Population of Mosquitoes overview









Risk	Mosquitoes_Infected	Mosquitoes_Latent	Mosquitoes_Suscept	Infected	Incubating	Healthy	Cluster
8.001615676660991494	0.88724746726266634	6.6443595063892245	74107.76447968498	1.5971563981842654	0.13270142180094788	507.8767772511848	8
0.0001469526885047312	0.007915132877592022	0.6027791310927828	63284.698183572385	0.11374407582938388	0	227,5781990521327	1
0.00008877825100572778	0.0047865677271341075	0.36452317759124747	42491.67841483456	0.9241706161137441	0	563.7251184834123	2
0.00016407507697713264	0.008840337921773394	0.6732384787362451	42491.365644963225	0.27014218009478674	0	182.3175355450237	3
0.00026297884467541	0.81915496463082248	1.458751622778464	42490.56981719248	4.876777251184834	0.0995260663507109	227,0995260663587	4
0.00005482812292195715	0.0034914900958950185	0.26589600213566816	42491.77833627965	8.88538885587283792	0	62.322274881516584	5
0.00004178887882985889	0.00225441915052684	0.1716859132633643	42491.873783447474	0.11848341232227488	0.02843601895734597	124.34123222748815	6
0.00012654109655685364	0.0068269562334023585	8.5199087949740706	42491.52898882868	1.6919431279620853	0.02843601895734597	594.7725118483412	7

Figure.14.Daily Cluster Details

4.Conclusion

In conclusion, agent-based modeling (ABM) of malaria transmission offers a promising approach to understanding the complex dynamics of malaria transmission. Through the simulation of individual agents representing humans, mosquitoes, and environmental factors, ABM allows for the exploration of intricate interactions that influence disease spread. The application of ABM enables researchers to investigate the impact of various factors such as human behavior, vector ecology, and intervention strategies on malaria transmission dynamics, providing valuable insights for malaria control and elimination efforts. Moving forward, continued research and development in agent-based modeling of malaria transmission are essential for enhancing our understanding of malaria epidemiology and informing evidence- based control strategies. Collaborative efforts among researchers, policymakers, and public health practitioners are crucial for advancing the field and translating research findings into actionable interventions. By leveraging the power of agent-based modeling alongside advances in data science and epidemiology, we can work towards the ultimate goal of reducing the global burden of malaria and achieving sustainable malaria control and elimination.

References

1. Epstein, J. M., & Axtell, R. (1996). Growing artificial societies: social science from the bottom up. MIT press.

2. Smith, D. L., Battle, K. E., Hay, S. I., Barker, C. M., Scott, T. W., & McKenzie, F. E. (2012). Ross, Macdonald, and a theory for the dynamics and control of mosquito-transmitted pathogens. PLoS pathogens, 8(4), e1002588.

3. Tatem, A. J., Smith, D. L., & Gething, P. W. (2010). Urbanization and the global malaria recession. Malaria journal, 9(1), 1-12.

4. Gething, P. W., Patil, A. P., Smith, D. L., Guerra, C. A., Elyazar, I. R., Johnston, G. L., ... & Hay, S. I. (2011). A new world malaria map: Plasmodium falciparum endemicity in 2010. Malaria journal, 10(1), 1-17.

5. Smith, D. L., Perkins, T. A., Reiner Jr, R. C., Barker, C. M., Niu, T., Chaves, L. F., ... & Hay, S. I. (2013). Recasting the theory of mosquito-borne pathogen transmission dynamics and control. Transactions of the Royal Society of Tropical Medicine and Hygiene, 107(5), 331-338.

6. Ferguson, N. M., Kien, D. T. H., Clapham, H., Aguas, R., Trung, V. T., Chau, T. N. B.,

... & Truong, N. T. (2015). Modeling the impact on virus transmission of Wolbachia- mediated blocking of dengue virus infection of Aedes aegypti. Science translational medicine, 7(279), 279ra37-279ra37.

7. Ross, R. (1911). The prevention of malaria. Cambridge University Press.

8. Eckhoff, P. A. (2012). A malaria transmission-directed model of mosquito life cycle and ecology. Malaria journal, 11(1), 1-16.

9. Eckhoff, P. A. (2011). P. falciparum infection durations and infectiousness are shaped by antigenic variation and innate and adaptive host immunity in a mathematical model. PLoS one, 6(1), e16163.

10. Huijben, S., & Paaijmans, K. P. (2018). Putting evolution in elimination: Winning our ongoing battle with evolving malaria mosquitoes and parasites. Evolutionary Applications, 11(3), 415-430.

11. Stoddard, S. T., Morrison, A. C., Vazquez-Prokopec, G. M., Paz Soldan, V., Kochel, T. J., Kitron, U., ... & Scott, T. W. (2009). The role of human movement in the transmission of vector-borne pathogens. PLoS neglected tropical diseases, 3(7), e481.

12. Koella, J. C., & Antia, R. (2003). Epidemiological models for the spread of anti-malarial resistance. Malaria journal, 2(1), 1-11.

13. Carriere, Y., Deland, J. P., Roff, D. A., & Vincent, C. (1994). Life-history costs associated with the evolution of insecticide resistance. Proceedings of the Royal Society of London. Series B: Biological Sciences, 258(1352), 35-40.

14. Griffin, J. T., Hollingsworth, T. D., Okell, L. C., Churcher, T. S., White, M., Hinsley, W.,

962

... & Ghani, A. C. (2010). Reducing Plasmodium falciparum malaria transmission in Africa: a modelbased evaluation of intervention strategies. PLoS medicine, 7(8), e1000324.

15. Beier, J. C., Perkins, P. V., Wirtz, R. A., Koros, J., Diggs, D., Gargan II, T. P., & Koech,

D. K. (1988). Bloodmeal identification by direct enzyme-linked immunosorbent assay (ELISA), tested on Anopheles (Diptera: Culicidae) in Kenya. Journal of Medical Entomology, 25(1), 9-16.

16. Beier, J. C., Killeen, G. F., Githure, J. I., & Entomological Society of America. (1999). Short report: entomologic inoculation rates and Plasmodium falciparum malaria prevalence in Africa. The American journal of tropical medicine and hygiene, 61(1), 109-113.

17. Focks, D. A., Daniels, E., Haile, D. G., & Keesling, J. E. (1995). A simulation model of the epidemiology of urban dengue fever: literature analysis, model development, preliminary validation, and samples of simulation results. American journal of tropical medicine and hygiene, 53(5), 489-506. 18. Focks, D. A., Brenner, R. J., Hayes, J., & Daniels, E. (2000). Transmission thresholds for dengue in terms of Aedes aegypti pupae per person with discussion of their utility in source reduction efforts. The American journal of tropical medicine and hygiene, 62(1), 11-18.

19. Killeen, G. F., & Smith, T. A. (2007). Exploring the contributions of bed nets, cattle, insecticides and excitorepellency to malaria control: a deterministic model of mosquito host-seeking behaviour and mortality. Transactions of the Royal Society of Tropical Medicine and Hygiene, 101(9), 867-880.

20. Koella, J. C., Lynch, P. A., & Thomas, M. B. (2009). Read, Andrew F. (ed.): Dynamics of infectious diseases: vectors, population biology, and principles of control. QRB. Quarterly Review of Biology, 84(4), 479-480.

21. Pongvongsa, T., Nonaka, D., Iwagami, M., Nakatsu, M., Phongmany, P., Nishimoto, F.,

... & Kano, S. (2017). Household clustering of asymptomatic malaria infections in Xepon district, Savannakhet province, Lao PDR. Malaria journal, 16(1), 1-11.

22. Wu, J., Dhiman, R., & Tewari, S. C. (2009). Development and evaluation of a mathematical model for malaria transmission dynamics in an endemic area. Computational and mathematical methods in medicine, 10, 105-121.

23. Legros, M., Lloyd, A. L., Huang, Y., Gould, F., & Godfray, H. C. (2009). Modeling the dynamics of a non-limited and a self-limited gene drive system in structured Aedes aegypti populations. PLoS neglected tropical diseases, 3(5), e427.

24. Beier, J. C., Killeen, G. F., Githure, J. I., & Entomological Society of America. (1999). Short report: entomologic inoculation rates and Plasmodium falciparum malaria prevalence in Africa. The American journal of tropical medicine and hygiene, 61(1), 109-113.

25. Wearing, H. J., & Rohani, P. (2006). Ecological and immunological determinants of dengue epidemics. Proceedings of the National Academy of Sciences, 103(31), 11802-11807.

26. Koella, J. C., & Antia, R. (2003). Epidemiological models for the spread of anti-malarial resistance. Malaria journal, 2(1), 1-11.

27. Killeen, G. F., Seyoum, A., Gimnig, J. E., Stevenson, J. C., Drakeley, C. J., Chitnis, N., & Govella, N. J. (2014). Made-to-measure malaria vector control strategies: rational design based on insecticide properties and coverage of blood resources for mosquitoes. Malaria journal, 13(1), 1-14.

28. Lengeler, C., & Cattani, J. A. (1996). Integrated vector management (IVM): rational decisionmaking for vector control. In Malaria control in complex emergencies (pp. 131-133). World Health Organization.

29. Pates, H., Curtis, C., & Mosha, F. (2007). Combination of indoor residual spraying with longlasting insecticide-treated nets for malaria control in Zambezia, Mozambique: a cluster randomised trial and cost-effectiveness study protocol. Trials, 8(1), 1-11.

30. Ferguson, N. M., & Donnelly, C. A. (2002). Anderson, Roy M. (ed.): Infectious diseases of humans: dynamics and control. Science, 296(5566), 1374-