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Zika virus: Endemic and epidemic ranges of *Aedes* mosquito transmission



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KEYWORDS

Geographic information systems; Geographic information science; Risk mapping; Zika; Aedes modeling **Summary** As evidence linking Zika virus with serious health complications strengthens, public health officials and clinicians worldwide need to know which locations are likely to be at risk for autochthonous Zika infections. We created risk maps for epidemic and endemic *Aedes*-borne Zika virus infections globally using a predictive analysis method that draws on temperature, precipitation, elevation, land cover, and population density variables to identify locations suitable for mosquito activity seasonally or year-round. *Aedes* mosquitoes capable of transmitting Zika and other viruses are likely to live year-round across many tropical areas in the Americas, Africa, and Asia. Our map provides an enhanced global projection of where vector control initiatives may be most valuable for reducing the risk of Zika virus and other *Aedes*-borne infections.

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Introduction

Zika virus is not a new infection [1]. The first known case of Zika was identified in a monkey

in Uganda in 1947. Cases of the virus in humans were later identified in Uganda and Tanzania in 1948 [2]. Cases from many other countries in central and West Africa were reported between the 1960s and 1980s, and the virus was also found in Indonesia, Malaysia, Pakistan and Costa Rica. In 2009, an outbreak occurred on the island of Yap in the Federal States of Micronesia. In subsequent years, cases were recognized in several other Asian

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and Pacific Island nations. In 2015, a large number of cases of Zika were diagnosed in Brazil, and cases began occurring in other countries within Central and South America and the Caribbean [1].

As the evidence linking Zika virus with serious health complications such as Guillain-Barré syndrome and microcephaly strengthens, public health officials and clinicians worldwide need to know which locations are likely to be at risk for autochthonous Zika infections [2]. This kind of location intelligence informs actions such as vector control strategies, public risk communications for disease prevention, and clinician training in diagnosis and response. At present, Aedes mosquitoes are considered to be the primary vectors of transmission for the Zika virus, particularly Aedes aegypti and Aedes albopictus [3]. In this paper, we use a variety of environmental and population datasets to identify locations where the habitat is suitable for Aedes mosquitoes to be present year-round or on a seasonal basis.

Methods

The goal of this analysis was to create an efficient, robust, and high resolution $(1 \text{ km} \times 1 \text{ km})$ global analysis of Zika risk by creating a map of suitable environments for *Aedes* mosquitoes. To this, we (1) conducted a review of the dengue literature to identify the social and environmental factors most strongly associated with *Aedes*-borne infections such as Zika, chikungunya, dengue, and yellow fever [4], (2) acquired map layers for all five parameters that the literature search revealed were the best predictors of areas where mosquitoes are known to be present, and then (3) used ArcGIS Predictive Analysis Tools to map the places that are most suitable for Zika outbreaks.

For seasonal Aedes suitability, the literature review identified as the most suitable locations (1) elevation below 1800 m; (2) precipitation levels allowing mosquitoes to survive and breed; (3) annual mean temperature between 10 °C and 30 °C; (4) land cover that excludes areas that are barren, covered with permanent snow or ice, or bodies of water; and (5) a population density of greater than 0 people, since humans must be present in a place for it to be a risk zone for human Zika infections [4]. For mapping year-round suitability for Aedes mosquitoes, the minimum temperature of the coldest month had to be above $10^{\circ}C$ and the other four criteria remained unchanged. This change in temperature suitability narrowed the possible areas where Aedes mosquitoes could survive throughout the year and gave a clearer picture of which locations globally have the greatest potential to become endemic for *Aedes*-borne diseases [5].

Spatial data sources with global coverage were identified for each of the key parameters: (1) elevation from the WorldClim (publically available) database version 1.4 (release 3) [6]; (2) annual precipitation, the precipitation of the wettest and driest months, and the precipitation of the wettest, driest, and coldest quarters from WorldClim [6]; (3) annual mean temperature, the mean of the difference between the monthly maximum and minimum temperatures, the standard deviation of mean temperatures by guarter, the maximum temperature of the warmest month, the minimum temperature of the coldest month, and the mean temperature of the warmest guarter from WorldClim [6]; (4) 2009 land cover data from the ESA 2010 and UCLouvain Team (publically available) [7]; and (5) 2011 data on human population density from Oak Ridge National Laboratory's LandScan Database (available to researchers upon request) [8]. Additional details about these spatial datasets and the process for selecting key variables are available elsewhere [4].

The ArcGIS Predictive Analysis Tools Add-In (PA Tools), a free add-on to ArcGIS available from Esri (Redlands, CA) since 2014 [9], was used to identify all the locations globally where the elevation, precipitation, temperature, land cover, and population density characteristics were similar to the places where previous outbreaks of Aedes-borne infections have occurred. The Query Expression Editor within the PA Tools classified each location as having zero to five of the five key risk factors for Zika (altitude, temperature, precipitation, land cover, and population density) were present. Each location was classified on a scale from not suitable (0 parameters present) to highly suitable (all 5 parameters present). The adaptable analysis within the geographic user interface for PA Tools allowed for immediate visualization of how various inputs affect the suitability model, and the values for each parameter could be analyzed individually.

As a validation check, we compared our results with historic data (1960 through 2014) about *Aedes* locations from the Dryad digital repository [10]. This congruence provided face validity for the map we created from our literature search. While historic data on mosquito locations have been critical for past risk mapping efforts, they are not sufficient on their own to illustrate where Zika could occur today or in future years. First, there is no database of locations were *Aedes* are known to be absent, so sites not included in the database cannot necessarily be assumed to be *Aedes*-free. Also, since the range for these mosquitoes is expected

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