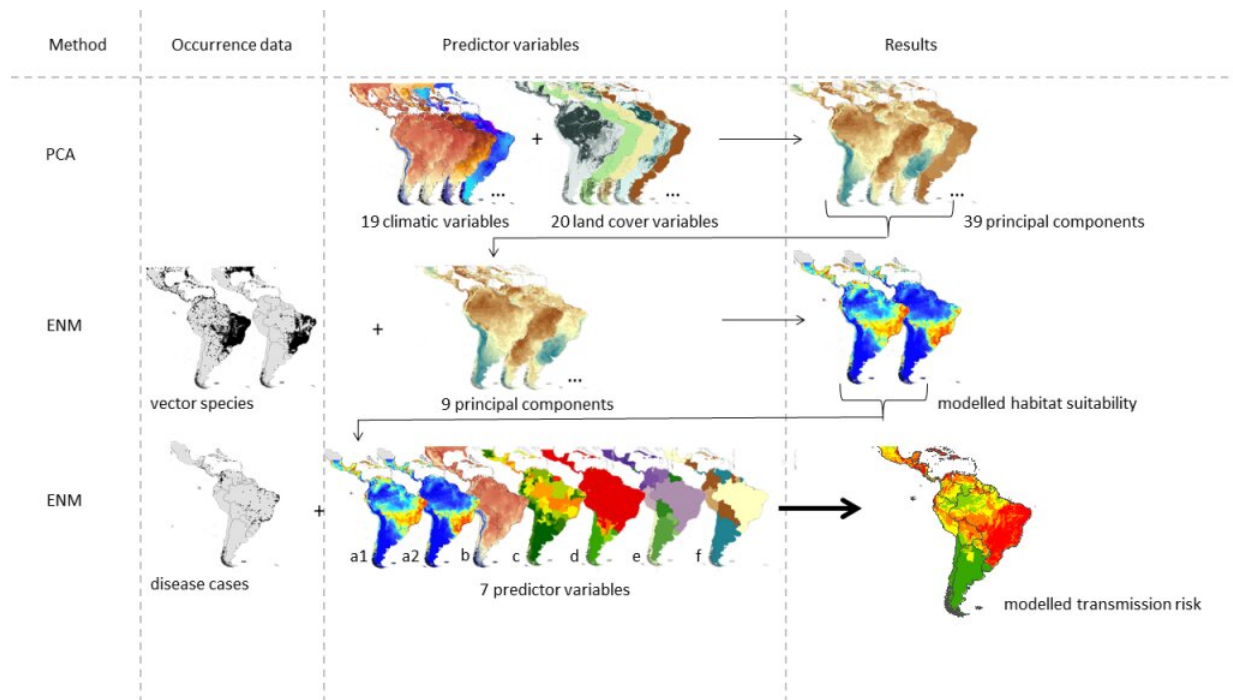


# Researchers developing maps on Zika virus infection risk

November 11 2019



Correlative modelling approach for regions with increased Zika infection risk in South America. In addition to the modelled expansion of the two main vector types (*Aedes aegypti* a1 and *Ae. albopictus* a2), the model incorporates the average temperature of the warmest quarter (b), the occurrence of the Zika virus (c) and the Dengue virus (e), population density (f) and the gross domestic product. Credit: Goethe University Frankfurt am Main

The spread of infectious diseases such as Zika depends on many

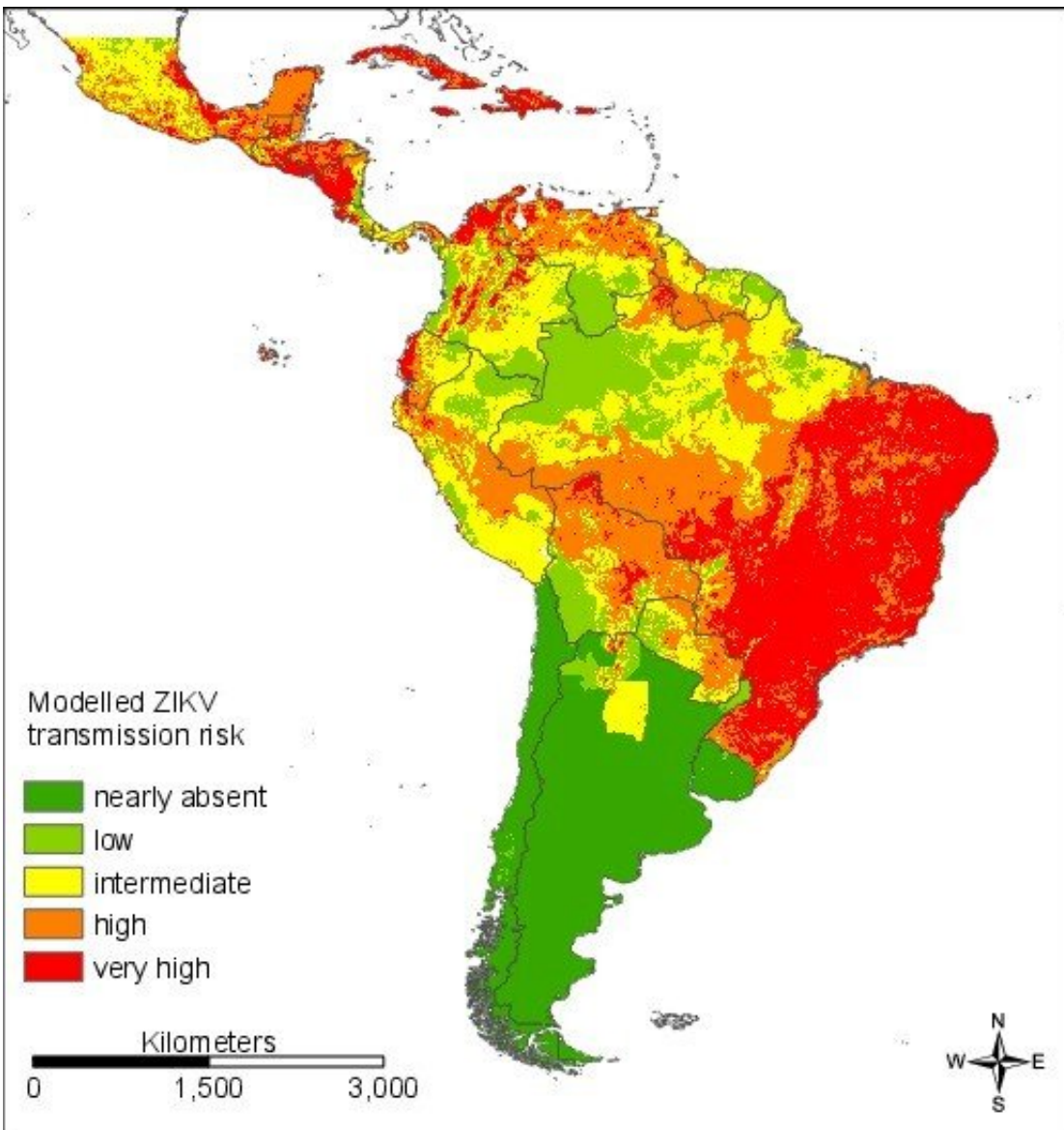
different factors. Environmental factors play a role, as do socioeconomic factors. Recently, several attempts have been made to predict the transmission risk of the Zika virus at a global and local level, but the spatial and temporal patterns of transmission are still not well understood. Researchers from Goethe University and the Senckenberg Society for Nature Research in Frankfurt were now able to generate reliable maps for the transmission risk of the Zika virus in South America. The results have been published in the scientific journal *PeerJ*. Based on the models for South America, they will use the method to determine the Zika risk for Europe as well.

In most cases, mosquitoes of the genus *Aedes* transmit the Zika virus to humans. Primary vectors are the yellow fever mosquito (*Aedes aegypti*) and the Asian tiger mosquito (*Aedes albopictus*). Both mosquito species are widespread in South America. Whereas the yellow fever mosquito (*Aedes aegypti*) is nearly absent in Europe, the Asian tiger mosquito is widespread in the Mediterranean region.

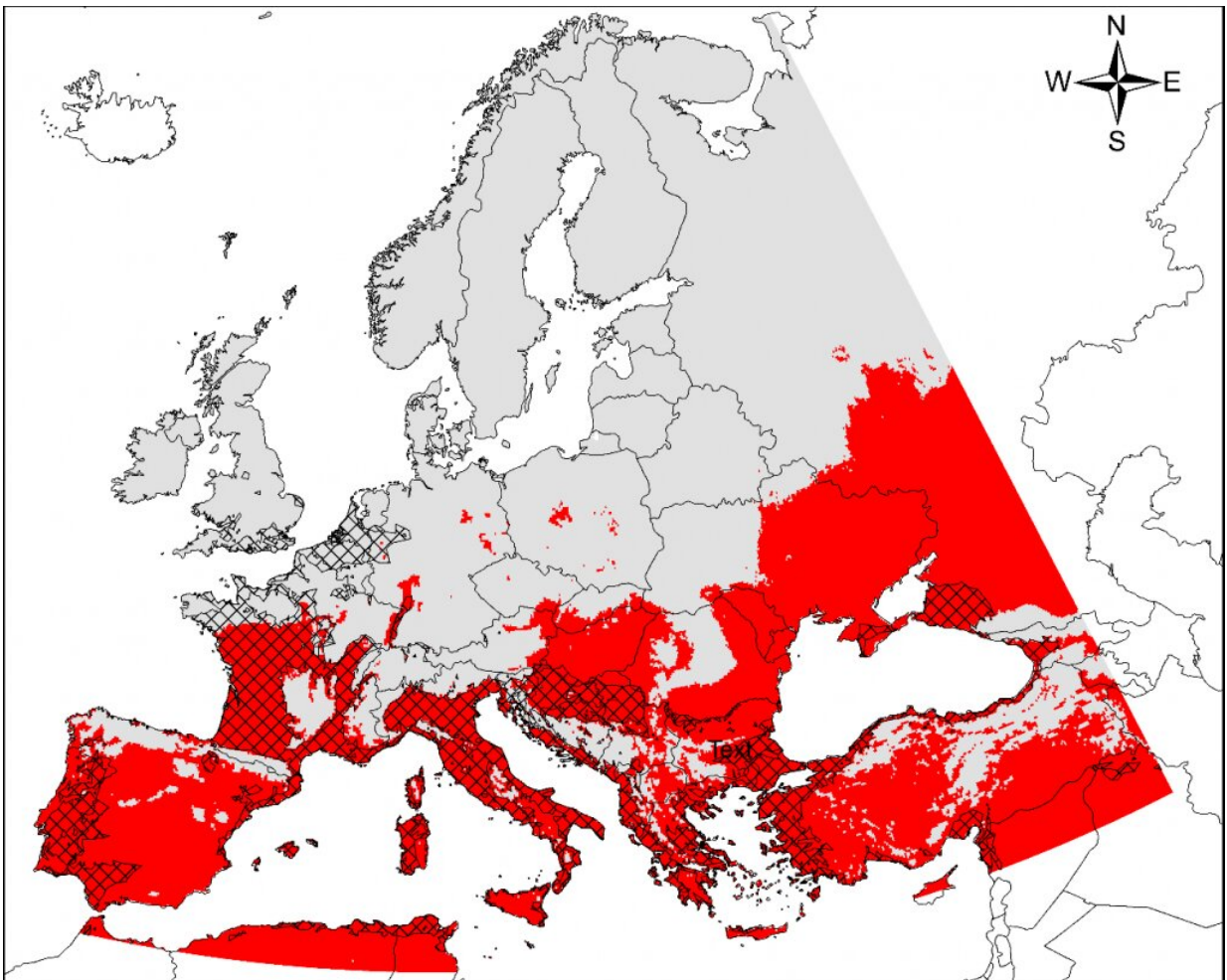
"With our new modeling approaches we can illustrate the risk areas for Zika infections in Latin America," says Sven Klimpel Professor for Parasitology and Infection Biology at Goethe University in Frankfurt and the Senckenberg Biodiversity and Climate Research Centre. "The models additionally allow us to illustrate Zika risk areas for Europe. For example, our models indicate the two autochthonous cases in southern France in Département Var (see illustration)." At the end of October, French authorities announced the first Zika case in Europe; about a week later, a second case was made public.

According to the researcher's calculations, the Zika infection risk in South America is highest along the Brazilian East Coast and in Central America. The risk is moderate in the Amazon region and lowest in the southern areas of the continent. The following countries are especially affected according to the model: Brazil, Columbia, Cuba, the Dominican

Republic, El Salvador, Guatemala, Haiti, Honduras, Jamaica, Mexico, Puerto Rico and Venezuela. In Europe, a risk of infection exists mainly in the Mediterranean region, but also in the inland regions of France and in the Rhine areas of Baden-Württemberg.




Zika infection risk modelled for South America. Credit: Goethe University Frankfurt am Main



 18°C bis 34°C

 modellierte Habitateignung für

*Aedes albopictus*

Kilometer  
  
 0 500 1,000

Regions where an autochthonous transmission of the Zika virus through the bite of an infected tiger mosquito is possible, since the temperature conditions are suitable (red areas) and the tiger mosquito is already present as vector species (hatched areas). Credit: Goethe University Frankfurt am Main

To determine the infection risk of a specific area, the researchers Dr. Sarah Cunze and Professor Sven Klimpel modeled the potential spread of the two species of mosquito, *Aedes aegypti* and *Aedes albopictus*. Since the mosquitoes can only transmit the Zika virus in regions where the virus is present in the first place, the researchers included an Evidence Consensus Map in their risk model. This map categorizes the number of reported Zika illnesses at the regional level. The average temperature of the warmest quarter was also incorporated in the model, since temperature has a significant influence on whether the virus can survive and multiply in the mosquito. Finally, the scientists added socioeconomic factors such as population density and gross domestic product to their risk model.

**More information:** Sarah Cunze et al. Vector distribution and transmission risk of the Zika virus in South and Central America, *PeerJ* (2019). [DOI: 10.7717/peerj.7920](https://doi.org/10.7717/peerj.7920)

Provided by Goethe University Frankfurt am Main

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