Asian tiger mosquito poses low risk for Zika virus outbreaks
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The Asian tiger mosquito does not pose a major risk for Zika virus epidemics, according to a study published December 31 in the open-access journal *PLOS Pathogens* by Albin Fontaine of the Institut de Recherche Biomédicale des Armées, and colleagues.

Zika virus has triggered large outbreaks in human populations, in some cases causing congenital deformities, fetal loss, or neurological problems in adults. While the yellow fever mosquito *Aedes aegypti* is considered the primary vector of Zika virus, the Asian tiger mosquito *Aedes albopictus* has been shown experimentally to transmit the virus and was involved in several transmissions of the virus in France in 2019. Originating from Southeast Asia, *Ae. aegypti* is an aggressive biter that has invaded the world and is now present on all inhabited continents, including temperate Europe, due to its ability to endure harsh winter conditions. As the second most important vector of human viral pathogens, *Ae. albopictus* is displacing *Ae. aegypti* populations due to competitive advantages. But it is not known if *Ae. albopictus* could trigger large-scale Zika virus epidemics.

To address this question, the researchers exposed *Ae. albopictus* to Zika virus and assessed infection rates in experiments, modeled the dynamics of Zika virus infection within individual humans, and used epidemiological simulations. The highest risk of transmission occurred during the pre-symptomatic stage of the disease. At this dose, mosquito infection probability was estimated to be 20%, and 21 days were required to reach median systemic infection rates. Despite these unfavorable characteristics for transmission, *Ae. albopictus* was still able to trigger large outbreaks in a simulated environment in the presence of sufficiently high mosquito densities and biting rates. According to the authors, active surveillance and eradication programs should be implemented in territories occupied by *Ae. albopictus* to maintain the low risk of Zika virus outbreaks.

The authors conclude, "The complementary combination of dose-dependent experimental infection, modeling of intra-human viremia dynamics, and in silico epidemiological simulations confirms the low epidemic potential of *Aedes albopictus* for Zika virus."
