Zika virus outbreak: ‘a perfect storm’

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Emerging Microbes and Infections (2016) 5, e21; doi:10.1038/emi.2016.42; published online 9 March 2016

Zika virus, a once-overlooked single-stranded ribonucleic acid (RNA) virus that causes a mosquito-borne viral disease, is causing a major ongoing pandemic worldwide. On 1 February 2016, the World Health Organization declared Zika virus-associated clusters of microcephaly and related neurological disorders a ‘Public Health Emergency of International Concern’.

Why has a largely ignored virus, which normally causes only mild symptoms, reemerged with a different pathophysiology, much like an unpredicted storm?

THE EPIDEMIC EVOLUTION FROM SPORADIC TO PANDEMIC

Zika virus was first isolated from a *Macaca* monkey in Zika, Uganda, in 1947; human transmissions were later observed. The most important vector of Zika virus is *Aedes aegypti*. The virus can also be transmitted from mother to fetus during pregnancy or birth. Other modes of transmission such as sexual contact and transfusion with contaminated blood are also possible.

Between the 1950s and 1970s, only sporadic cases of Zika virus infection were reported in western and central Africa and south-east Asia. The disease nearly disappeared over two decades until multiple outbreaks in the Pacific islands marked the eastward movement of the virus in the 2000s. In 2007, Yap in Micronesia reported the first outbreak of Zika virus. The second outbreak occurred in 2013 in French Polynesia, with nearly 30 000 people becoming infected. From there, the epidemic gradually spread to the Cook Islands, New Caledonia and Easter Island in 2014. Genetic analyses have shown that the strains in the French Polynesia outbreak were more closely related to the Asian strains, indicating that the disease may have been introduced into the Pacific islands from Asia.

The Zika virus epidemic in South America first surfaced in February 2015, and the Ministry of Health of Brazil first confirmed the outbreak of Zika virus in northeastern Brazil in May 2015. Since then, the epidemic has spread further in South America. By 17 January 2016, the Pan American Health Organization reported that the Zika virus epidemic had spread to 18 countries or territories in the Americas.

ZIKA FEVER AND ITS POSSIBLE RELATIONSHIP WITH FETAL MICROCEPHALY

For more than half a century since its initial discovery, Zika virus seemed to pose little threat to human beings. Zika virus infection is asymptomatic in ~80% of infected individuals, and in those who develop the disease, the symptoms are usually mild and rarely result in human death. Therefore, updates on the treatment strategy for Zika virus infection have been slow, and there has been a relatively small amount of research on the Zika virus compared with other mosquito-borne viral diseases.

By the end of January 2016, nearly 4000 suspected cases of microcephaly have been reported in Brazil, constituting a 20-fold increase in incidence rate compared with the period of 2010–2014. Circumstantial evidence continues to accumulate. First, the period of the sudden microcephaly outbreak overlaps with the period when the Zika virus epidemic was introduced into Brazil. In addition, almost all mothers whose babies were diagnosed with microcephaly had symptoms of Zika virus infection in their early pregnancies. In the amniotic fluid of two mothers whose fetuses were diagnosed with microcephaly, Zika virus RNA has also been detected by reverse transcription polymerase chain reaction. However, until now, the strongest evidence may yet be the report of a case with severe brain disease and microcephaly related to vertical transmission of Zika virus. The fetal autopsy showed pathological changes detected only in the brain, suggesting a strong neurotropism of the virus.

However, we are still far from drawing relevant conclusions. Currently, suspected cases are diagnosed mostly due to patients’ traveling history and serological tests. Because serological tests of Zika virus have a high cross-reactivity with other flaviviruses, the epidemic scale may be exaggerated. The Brazilian outbreak affected far more people than any other previous outbreaks (Table 1), thus possibly explaining why no microcephaly cases were previously reported. However, the actual number of fetus microcephaly cases may be overestimated due to the low specificity of the screening test. Further studies on the differential diagnosis of the suspected cases are needed before we can establish a relationship between the Zika virus epidemic and fetal microcephaly.

WHY BRAZIL?

Brazil is the center of this current outbreak, and an estimated 440 000–1 300 000 Brazilians were infected in 2015. At first glance, this outbreak of the Zika virus seemed completely random. However, several factors may have contributed to this outbreak.

First, genetic analyses have uncovered close relationships between the virus strains in the South American and Pacific Island outbreaks.
In 2015 compared with 2014, highlighting the poor control of arbovirus mainly transmitted by mosquitoes. In fact, the reported cases of Dengue fever, another mosquito-borne disease in Brazil. The lack of immunity for the Zika virus in Brazilians is perhaps the most compelling factor underlying the outbreak. In a country without a history of Zika virus disease, Brazilians’ total vulnerability to the virus would further accelerate disease transmission in this outbreak.

In addition, Brazil has a population density that is significantly higher than that of the Pacific Islands, thus making the disease much more difficult to control. Furthermore, most of Brazil has a tropical climate, and a large part of the Amazon rainforest is also located within this area, all of which could lead to a difficult battle against mosquitoes. In fact, the reported cases of Dengue fever, another arbovirus mainly transmitted by Aedes aegypti, increased by 227.12% in 2015 compared with 2014,17,18 highlighting the poor control of Aedes mosquitoes in Brazil. The lack of immunity for the Zika virus in Brazilians is perhaps the most compelling factor underlying the outbreak. In a country without a history of Zika virus disease, Brazilians’ total vulnerability to the virus would further accelerate disease transmission in this outbreak.

LESSONS AND FUTURE PERSPECTIVES

The Zika virus outbreak is currently experiencing ‘a perfect storm’, because of the widespread Aedes mosquitoes in South America, the introduction of the virus into a country with a high population density, possible viral mutations, the lack of immunity for Zika virus in Brazilians, increasing international travel and global warming all working together to create a negative dynamic with respect to human health. Currently, no country can declare itself completely safe from this outbreak, and all countries that have Aedes mosquitoes should stay on high alert. The Zika pandemic is still in progress, and the next few months may be crucial. Brazil is about to enter the rainy season, and the entire northern hemisphere will soon slowly transition to spring and summer, making the control of mosquitoes even more difficult. If high-population-density countries such as China, India, Japan or the United States experience a secondary outbreak (China and the United States have already reported imported cases), it would cause the Zika virus epidemic to become even more difficult to control. Currently, controlling Aedes mosquitoes and recommending that pregnant women avoid travel to Zika-virus-endemic regions and avoid being bitten by mosquitoes seems to be our best chance. However, in the future, we urgently need more research on the relationship between Zika virus and microcephaly, and doctors should establish differentiation in the diagnosis of suspected cases of Zika virus infection.

The Zika virus was unexpected and caught us off guard, and but it has taught us an important lesson. How many other neglected pathogens are out there, lurking in the shadows and waiting for the next ‘perfect storm?’ This is a scary question to ask. In a world with increasing globalization, international travel, urban crowding and global climate change, perhaps it is time for us to rethink our disease-control strategies, strengthen our research and public health infrastructures, and devise preventive measures against possible future disease outbreaks.

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Table 1 Past Zika virus outbreaks

| Country               | Year | Population | Reported Zika virus disease | Estimated total cases of Zika virus disease | Estimated infection rate | Reported complications |
|-----------------------|------|------------|-----------------------------|--------------------------------------------|--------------------------|------------------------|
| Yap, Micronesia⁵      | 2007 | 7391       | Confirmed: 49 Probable: 59  | 900                                        | 75%                      |                        |
| French Polynesia³,⁶   | 2013 | 268 000    | Confirmed: 396 Suspected: 8723 | 30 000                                     | 17%                      | 42 patients diagnosed with Guillain–Barré syndrome; 32 patients had other neurological or autoimmune syndromes; Unusual increase of 17 cases of central nervous system malformations in fetus and infants |
| Cook Island¹³         | 2014 | 14 974     | Confirmed: 50 Suspected: 932 | 44 000–1 300 000                            | 20%                      | Around 4000 cases of microcephaly in fetus and infants (up to December 2015); 121 neurological manifestations and Guillain–Barré syndrome (up to December 2015) |
| New Caledonia¹⁴       | 2014 | 254 000    | Confirmed: 1400              | 10 000                                      | 20%                      |                        |
| Brazil⁵,⁹,¹⁵           | 2015 | 205 338 000 |                              | 10 000                                      | 20%                      |                        |

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