Performance of 2 Commercial Serologic Tests for Diagnosing Zika Virus Infection
Severine Matheus, Cheikh Talla, Bhety Labeau, Franck de Laval, Sebastien Briolant, Léna Berthelot, Muriel Vray, Dominique Rousset

To cite this version:
Severine Matheus, Cheikh Talla, Bhety Labeau, Franck de Laval, Sebastien Briolant, et al.. Performance of 2 Commercial Serologic Tests for Diagnosing Zika Virus Infection. Emerging Infectious Diseases, Centers for Disease Control and Prevention, 2019, 25 (6), pp.1153-1160. 10.3201/eid2506.180361. hal-02263773

HAL Id: hal-02263773
https://hal-amu.archives-ouvertes.fr/hal-02263773
Submitted on 3 Jun 2020

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L’archive ouverte pluridisciplinaire HAL, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d’enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Distributed under a Creative Commons Attribution 4.0 International License
Reliable serologic tests are needed for diagnosis and surveillance of Zika virus infection. We evaluated the Euroimmun and Dia.Pro serologic tests for detection of Zika virus IgM and IgG by using a panel of 199 samples from a region endemic for flaviviruses. Kinetics of Zika virus antibodies were monitored from 300 sequential specimens sampled over a period of 10 months after infection. We observed suboptimal performance; sensitivity for Zika virus IgM was low, especially in the Euroimmun assay (49%), whereas IgM could be detected for months with the Dia.pro assay. The specificity of the Zika virus IgG assays was also low, especially that of Dia.Pro (62%); findings were strongly influenced by the epidemiologic context. These results highlight the complexity of serologic diagnosis of Zika virus infection in regions endemic for flaviviruses. Accurate analysis of the performance of assays is required to adapt and interpret algorithms.

Zika virus belongs to the Flaviviridae family, genus Flavivirus, and is an arbovirus transmitted mainly by mosquitoes of the genus Aedes (Stegomyia). Initially isolated in 1947 from a sentinel monkey during yellow fever surveillance in Uganda, Zika virus was reported as causing only sporadic human infections, associated with asymptomatic or mild, self-limiting illness, until 2007 (1,2). In 2007, Zika virus spread first to Pacific islands and then throughout the Americas, resulting in large outbreaks in several regions of the world. Zika virus infection is estimated to be symptomatic in 18%–73% of cases (3–5); severe complications have been reported, including neurologic disorders, such as Guillain-Barré syndrome, and congenital Zika virus syndrome, which is characterized by severe microcephaly, brain and ocular anomalies, congenital contractures, and neurologic impairment in the fetuses and newborns of mothers infected during pregnancy (3,4,6).

The mild signs and symptoms of Zika virus infection include fever, rash, joint pain, conjunctivitis, headache, and myalgia (7). These manifestations are difficult to distinguish clinically from those caused by other arboviral infections, such as dengue or chikungunya, which are often observed in the same geographic areas. Therefore, specific, reliable diagnostic tools are needed.

Several commercial kits are available for direct viral detection by nucleic acid–based testing, which enable diagnosis during the acute phase of the disease (8,9): up to 7 days after symptom onset in serum samples, up to 20 days in urine, and even longer in semen (10–13). This virologic window, combined with the high proportion of asymptomatic forms, makes the monitoring of Zika virus infection difficult, especially in pregnant women. Therefore, serologic tools for diagnosis of Zika virus infection are urgently needed. This challenge remains because of cross-reactivity among flaviviruses, especially in a context of secondary flavivirus infection or previous immunization.

The first objective of this study was to evaluate the performance of two commercial serologic kits for the detection of Zika virus–specific IgM and IgG in serum samples from patients with an arboviral-like syndrome in a region where other arboviruses are known to circulate, including dengue and chikungunya viruses. The 2 commercial kits (4 assays) studied were the Euroimmun Zika virus IgM and IgG ELISAs (https://www.euroimmun.com) and Dia.Pro Zika virus IgM and IgG ELISAs (Diagnostic Bioprobes Srl, https://www.diapro.it). The second objective was to determine the kinetics of Zika virus IgM and IgG induced after infection, as defined by these kits.

Material and Methods

Clinical Samples and Study Design

The clinical samples used in this study were selected according to the standards for reporting diagnostic accuracy.
requirements. Two thirds came from the serum collection of the National Reference Centre (NRC) for Arboviruses in French Guiana and one third from samples collected for a descriptive prospective study of Zika virus disease in the French military community in French Guiana (ZIFAG) (14). The NRC collection comprises clinical specimens received during 2002–2017 for routine diagnosis and expertise. The protocol of the ZIFAG study received ethics approval from the Comité de Protection des Personnes Sud-Méditerranée I (ID RCB: 2016-A00394–47); all participants provided written informed consent. All the selected specimens were obtained from patients with an arboviral-like syndrome and an etiologic diagnosis confirmed by real-time reverse transcription PCR (RT-PCR) on acute-phase serum or urine samples. A commercial qualitative RT-PCR kit (Altona Diagnostics, https://www.altona-diagnostics.com) was used for Zika virus detection and NRC in-house real-time RT-PCR for dengue and chikungunya viruses.

We first evaluated the performance of the 2 commercial immunoassays against a panel of 199 serum samples collected from days 3 through 20 after the onset of symptoms (onset defined as day 0—that is, within the first 24 hours) (Table 1). We evaluated the sensitivity of the assays in a subgroup of 109 serum samples with confirmed Zika virus infection diagnosed from the end of 2015 through 2016, during the outbreak in French Guiana (Zika subgroup). We evaluated the specificity of the assays in a subgroup of 109 serum samples with a strong potential for causing flavivirus cross-reactions (non-Zika subgroup). This subgroup comprised serum samples obtained from 35 patients with confirmed dengue virus infection sampled during 2002–2013 dengue epidemics; 29 patients with confirmed chikungunya virus infection sampled during the chikungunya outbreak in French Guiana in 2014–2015, just after the dengue outbreak in 2012–2013; and 45 patients with neither dengue virus, chikungunya virus, nor Zika virus infection sampled just before the Zika virus outbreak. We performed Zika microneutralization tests on the Zika virus, dengue virus, and chikungunya virus–negative subgroup of serum samples, enabling confirmation of the absence of Zika neutralizing antibodies.

To determine the kinetics of Zika virus antibodies, we used a panel of 300 serum samples collected from 124 patients with confirmed Zika virus infection from day 0 through day 300 after the onset of symptoms (Table 1). We collected 1–8 samples from each Zika virus–infected patient. The distribution of samples according to time since onset was as follows: 76 samples collected during days 0–4, 50 during days 5–14, 55 during days 15–30, 19 during days 31–60, 32 during days 61–90, 44 during days 91–180, and 24 during days 181–300.

**Euroimmun and Dia.Pro Zika Virus IgM and IgG ELISAs**

We tested all samples with the Euroimmun ELISAs according to the manufacturer’s recommendations and calculated signal-to-cutoff (S/CO) ratios; values <0.8 were regarded as negative, ≥0.8 to <1.1 as equivocal, and ≥1.1 as positive. We tested all serum specimens with the respective Dia.Pro ELISAs for qualitative determination of IgM and IgG, according to the manufacturer’s instructions; we interpreted results as positive if the S/CO ratio was ≥1.1, negative if <0.9, and equivocal if 0.9–1.1. Recombinant Zika virus nonstructural protein 1 was the antigen in both the Euroimmun and Dia.Pro assays.

**Statistical Analysis**

Continuous variables were expressed as mean (± SD) or median with interquartile range (IQR) and discrete variables as percentages and 95% CIs. We calculated the sensitivity and specificity of the assays with 95% CIs. The differences in the S/CO ratios of the IgM and IgG assays at each time were compared with the Student t test. We performed all statistical analyses using R 3.4 statistical software (https://www.r-project.org).

### Table 1. Characteristics of panels evaluated in study of Zika virus diagnostic tests

<table>
<thead>
<tr>
<th>Panel performance evaluation 3–20 days after onset</th>
<th>Sample characterization</th>
<th>No. samples</th>
<th>Mean time to collection after onset of fever, d (± SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Zika virus subgroup</td>
<td>Positive Zika virus</td>
<td>90</td>
<td>10 ± 6</td>
</tr>
<tr>
<td>Non–Zika virus subgroup</td>
<td>Negative Zika virus with confirmed dengue virus infection</td>
<td>35</td>
<td>10 ± 3</td>
</tr>
<tr>
<td></td>
<td>Negative Zika virus with confirmed chikungunya virus infection</td>
<td>29</td>
<td>13 ± 3</td>
</tr>
<tr>
<td></td>
<td>Negative Zika virus, dengue virus, chikungunya virus</td>
<td>45</td>
<td>11 ± 5</td>
</tr>
<tr>
<td>Total panel performance</td>
<td></td>
<td>199</td>
<td>10 ± 5</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Panel IgM–IgG kinetics 0–300 days after onset</th>
<th>No. samples</th>
<th>Mean time to collection after onset of fever, d (± SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Positive Zika virus</td>
<td>300*</td>
<td>Minimum 0, maximum 300, median 20, interquartile range 4–81</td>
</tr>
</tbody>
</table>

*300 samples from 124 patients, including the 90 positive Zika virus samples in the first panel.
Results

Patient Characteristics
The 199 samples used to evaluate the performance of the serologic kits came from patients with a mean ± SD age of 36 ± 16 years (range 1–74 years; IQR 27–46 years). This panel was composed of the Zika virus–positive subgroup (n = 90), 51 (57%) female and 39 (43%) male, with a mean age ± SD of 39 ± 12 years; and the Zika virus–negative subgroup (n = 109), 62 (57%) female and 45 (43%) male, with a mean ± SD age of 34 ± 18 years (Table 1).

The 300 samples used to determine the kinetics of Zika virus IgM and IgG came from 124 patients, 82 (66%) female and 42 (34%) male. The age range of these patients was 8–74 years (mean ± SD 37 ± 11 years).

Performance of Zika Virus IgM ELISAs
We evaluated the sensitivity and specificity of the 2 Zika virus IgM tests against 90 Zika virus subgroup samples and 109 non-Zika subgroup samples (Table 2). Each test gave inconclusive results for 6 of the 199 samples: with the Euroimmun test, we obtained 6 inconclusive results from the group negative for dengue virus, chikungunya virus, and Zika virus. The sensitivity of this assay was 71% (58/82; 95% CI 92%–100%) and the specificity 70% (71/101; 95% CI 61%–79%). With the Dia.Pro IgM test, 8 samples gave inconclusive results: 5 samples from the Zika virus–positive group (2 collected before day 5 and the others on day 9 or later after the onset of illness) and 3 samples from the Zika virus–negative group (collected on days 4 to 6 after onset) (Table 2). The sensitivity of this assay was 79% (67/85; 95% CI 70%–88%) and the specificity 62% (66/106; 95% CI 53%–71%).

The false positivity rate of the 2 Zika virus IgG assays varied according to the subpanel used. These rates were 40%–58.6% for positive dengue virus or chikungunya virus sample subgroups and 11.9%–21.4% for the negative dengue virus, negative chikungunya virus, and negative Zika virus sample subgroups (Table 3).

Table 2. Performance of Zika virus IgM and IgG assays in panels of characterized samples obtained during days 3–20 after onset of symptoms*

<table>
<thead>
<tr>
<th>Results</th>
<th>Zika subgroup</th>
<th>Non–Zika subgroup</th>
<th>Sensitivity (95% CI), specificity (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Euroimmun IgM test</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Positive</td>
<td>41</td>
<td>1</td>
<td></td>
</tr>
<tr>
<td>Negative</td>
<td>43</td>
<td>108</td>
<td>35</td>
</tr>
<tr>
<td>Inconclusive</td>
<td>6</td>
<td>0</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>90</td>
<td>109</td>
<td>35</td>
</tr>
<tr>
<td><strong>Dia.Pro IgM test</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Positive</td>
<td>59</td>
<td>4</td>
<td></td>
</tr>
<tr>
<td>Negative</td>
<td>27</td>
<td>103</td>
<td>30</td>
</tr>
<tr>
<td>Inconclusive</td>
<td>4</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>90</td>
<td>109</td>
<td>35</td>
</tr>
<tr>
<td><strong>Euroimmun IgG test</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Positive</td>
<td>58</td>
<td>30</td>
<td>14</td>
</tr>
<tr>
<td>Negative</td>
<td>24</td>
<td>71</td>
<td>21</td>
</tr>
<tr>
<td>Inconclusive</td>
<td>8</td>
<td>8</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>90</td>
<td>109</td>
<td>35</td>
</tr>
<tr>
<td><strong>Dia.Pro IgG test</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Positive</td>
<td>67</td>
<td>40</td>
<td>14</td>
</tr>
<tr>
<td>Negative</td>
<td>18</td>
<td>66</td>
<td>21</td>
</tr>
<tr>
<td>Inconclusive</td>
<td>5</td>
<td>3</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>90</td>
<td>109</td>
<td>35</td>
</tr>
</tbody>
</table>

*CHIKV, chikungunya virus; DENV, dengue virus; +, positive; −, negative.
Performance of Combined IgM/IgG Assays

The sensitivity of the combined Euroimmun Zika virus IgM and IgG assays was 82% (71/87; 95% CI 74%–90%) and the specificity was 69% (70/101; 95% CI 60%–78%). The sensitivity of the combined Dia.Pro Zika virus IgM and IgG assays was 87% (75/86; 95% CI 80%–94%) and specificity was 62% (66/106; 95% CI 53%–71%).

Time-Course Analysis of Zika Virus IgM and IgG, Days 0–300 after Onset of Symptoms

We used the panel of sequential samples from patients with confirmed Zika virus infection to determine the kinetics of Zika virus IgM and IgG over 10 months after clinical onset (Figure 1). For the Euroimmun Zika virus IgM test, maximum percentage detection (71%) was 15–30 days after the onset of disease. After this time, the percentage of detectable IgM decreased rapidly, to only 21% of samples collected in days 31–60 and <9% for those collected >60 days after infection (Figure 1, panel A). The Dia.Pro Zika virus IgM test was more sensitive, with higher rates of positive samples observed over a longer time: 29% positive samples on days 0–4 after clinical onset, increasing to a maximum of 93% positivity during days 15–30. The positivity rate decreased more slowly than with the Euroimmun test, and 29% of samples were still positive for IgM during days 181–300 after infection. For Zika virus IgG, both assays detected the antibody in >40% of samples collected during the acute phase of disease (days 0–4) and in 100% of samples collected during days 31–180 after onset of disease (Figure 1, panel B). A slight decrease in the positivity rate for IgG observed with the Euroimmun assay before day 300 suggests a possible lack of sensitivity over time.

We also evaluated the evolution of the overall mean S/CO ratios by time and the test used (Figure 1, panels C and D). We observed similar kinetics for the mean S/CO ratios...
for the 2 assays: the maximum mean S/CO ratio peaked during days 15–31 for the IgM assays and during days 91–180 for the IgG assays. Nevertheless, the differences between the mean S/CO ratios for both the IgM and the IgG assays at each time after infection class were significant (all p<0.05) (Figure 1, panels C and D). The Dia.Pro assays gave higher S/CO ratios for the same threshold values, explaining the greater sensitivity of these tests. We obtained the kinetics of Zika virus IgM and IgG with both assays for patients for whom we had ≥5 sequential samples (Figure 2).

Figure 2. Individual time-course analyses of Zika virus IgM and IgG signal-to-cutoff ratios obtained by using Euroimmun and Dia.Pro kits for 18 patients for whom 5 or more sequential samples were available.
Discussion

In this assessment of the performance of Euroimmun and Dia.Pro Zika virus IgM and IgG ELISAs for diagnosis of Zika virus infection, we had a large panel of well-characterized samples from areas endemic for arboviruses, the dates of symptom onset, and infection confirmed by real-time RT-PCR. Although the Dia.Pro IgM assay was more sensitive (69%) than the Euroimmun IgM test (49%), the performance of both IgM assays was suboptimal. The sensitivity of a screening test is a major factor in determining its usefulness, because poor sensitivity of the first test used in a diagnostic algorithm could lead to false-negative results that would not be further evaluated.

The sensitivity of the Euroimmun Zika virus IgM test was significantly lower than that reported previously (15–18). The differences might be caused by differences in study design, selection criteria, and the few positive samples in the previous studies (17–18). The lower sensitivity we found for this assay might also be the result of the larger proportion of secondary flavivirus infections in the panel used, as the samples were taken in an area endemic–epidemic for dengue and with mandatory vaccination against yellow fever. A significant lower sensitivity of the Euroimmun Zika virus IgM assay has also been reported in travelers from Israel compared with travelers from Europe and Chile, possibly related to the West Nile virus background immunity of the population of Israel (19). More recent evaluations have also reported low sensitivity (39.5% and 37%) of the Euroimmun assays (20,21).

Sensitivity is essential for a frontline diagnostic test, and specificity should also be carefully evaluated. In our study, we assessed the specificity of all the tests with various non–Zika virus samples to evaluate potential cross-reactivity. A first subpanel of samples from patients with confirmed acute dengue virus infection was formed because of the high potential for flavivirus cross-reactivity; a second subpanel consisted of samples from patients with confirmed chikungunya virus infection; and a third subpanel consisted of samples from patients with no dengue virus, chikungunya virus, or Zika virus infection. The specificity of the Euroimmun Zika virus IgM assay was 99% and that of the Dia.Pro test was 96%, with cross-reactivity varying according to the subpanel. Most cross-reactions were observed in the subpanel of acute dengue samples, in which 3 of 33 samples were false positive for Zika virus IgM, whereas 1 of 74 samples collected >1 year after the dengue epidemic (chikungunya virus subpanel and dengue virus, chikungunya virus, and Zika virus negative subpanel) was false positive. Maximum cross-reactivity of IgG tests was seen in the subpanel collected in the post–dengue epidemic period in 2014–2015. The false-positivity rate was 40% (14/35) in the acute dengue subpanel for both commercial tests; the rate grew to 45.8% (11/24) for the Euroimmun Zika virus IgG test and 58.6% (17/29) for the Dia.Pro Zika virus IgG test in the subpanel of acute chikungunya virus samples collected right after the dengue epidemic period in 2014–2015. The IgG cross-reactions tended to decrease with time: 2 years after the end of the latest dengue epidemic in French Guiana, 5/42 (11.9%) samples were falsely positive by the Euroimmun IgG test and 9/42 (21.4%) samples were falsely positive with the Dia.Pro test. All samples except 2 with a false-positive result for Zika virus IgG were positive for dengue virus IgG by our in-house IgG antibody capture ELISA technique. The predictive positive value of both Zika virus IgG assays therefore largely depends on the epidemiologic situation of other flaviviruses, like dengue virus.

A study performed in Martinique during March–June 2016 during the Zika virus epidemic showed a good correlation between a high Euroimmun Zika virus IgG ratio and a positive Zika virus seroneutralization result. Ratios >4 were associated with positive seroneutralization in >95% of cases, whereas ratios >5 were associated with seroneutralization in 100% of cases (22). At the time of the study, only sporadic dengue cases were reported in Martinique, as the previous epidemic occurred in 2013–2014, >2 years earlier. In our assessment, the IgG S/C ratio value of false Zika virus IgG–positive samples varied from 1.1 to 6.7 (mean 3.9) for acute dengue samples to 1.2 to 3.7 (mean 2.0) for other non–Zika virus samples. These results underline the importance and potential efficacy of using selected panels to evaluate performance and, especially, the specificity of serologic assays. These results also indicate that when there is major cocirculation of dengue virus and Zika virus, the interpretation of serologic assays could be increasingly complex.

The overall specificity of the Euroimmun Zika virus IgG assay was 79% and that of the Dia.Pro assay was 62%. These results indicate suboptimal specificity, which is lower than that reported previously (20,21), and is a concern for serologic diagnosis of Zika virus infection. Seroneutralization is the classical reference test for confirming contact with Zika virus in cases of positive results with ELISA assays in regions where flaviviruses cocirculate; however, even seroneutralization tests could be difficult to interpret, leading the US Centers for Disease Control and Prevention to change its guidance for interpretation of Zika virus antibody test results in May 2016 (23,24). In cases of secondary flavivirus infection, a microneutralization test might not discriminate between the past and recent infecting viruses, leading to an assumption of just a recent flavivirus exposure.

As reported previously by others, when IgM and IgG results were combined, sensitivity increased to 82% for Euroimmun and 87% for Dia.Pro assays, whereas specificity decreased to 69% for Euroimmun and 62% for Dia.Pro
(20,21). However, according to this combined analysis, 42% (30/71) of samples positive by the Euroimmun assays and 21% (16/75) for the Dia.Pro assays correspond to IgM negative/IgG positive samples, for which distinction between recent and past infections is not possible. Thus, a combined interpretation is not suitable for Zika infection diagnosis in the context of endemic–epidemic circulation.

We not only assessed the performance of Euroimmun and Dia.Pro Zika virus IgM and IgG tests but also evaluated the kinetics of the antibodies through 300 days after the onset of symptoms. A major concern in serologic diagnosis of Zika virus infection is determining the date of infection, due to the high proportion of asymptomatic forms. Data on the duration of IgM persistence after Zika virus infection are still limited, but our results indicate that this antibody could persist for at least several months, as described for other arboviruses (25). Such persistence could preclude determination of the recent nature of an infection, and other assays should be evaluated. The analysis of individual Zika virus antibody kinetics revealed distinct patterns. In some patients (such as patients 4, 5, 10, 16, 53), high IgM ratios during the acute phase were associated with delayed and moderate increases in Zika virus IgG ratios, possibly reflecting a primary flavivirus or Zika virus infection; for other patients (such as patients 20, 34, 41, 58, 62), an early increase in the IgG ratio was combined with a low or even negative Zika virus IgM ratio throughout follow-up, indicating secondary flavivirus infections. These 2 types of kinetics showed contrasted signal intensities, which are not observed with our in-house ELISA assays, in which whole Zika virus is used as an antigen (data not shown).

A limitation of our study is that the Zika virus–positive samples were confirmed by RT-PCR and thus were all from symptomatic cases. If antibody levels are different in symptomatic and asymptomatic infections, as described for dengue, the performance and antibody kinetics observed in this study might be considerably different in samples from asymptomatic infections (26,27).

This study highlights the complexity of interpreting serologic assays in areas where various arboviruses co-circulate and demonstrates the importance of evaluating serologic assays with serum specimens from persons living in endemic–epidemic areas and use of parallel testing antibodies to maximize the reliability of diagnosis. Further studies are also needed to identify specific biomarkers of each flavivirus infection for diagnosis after the acute phase of disease.

This study was supported by Institut Pasteur de la Guyane, Cayenne, French Guiana. Her primary research interest is the diagnosis and pathophysiology of arboviruses.

About the Author
Dr. Matheus is a research assistant at the Institut Pasteur de la Guyane, Cayenne, French Guiana. Her primary research interest is the diagnosis and pathophysiology of arboviruses.

References


Address for correspondence: Dominique Rousset, Institut Pasteur de la Guyane, Laboratoire de Virologie, 23 avenue Pasteur, BP 6010, 97306 Cayenne CEDEX, French Guiana; email: drousset@pasteur-cayenne.fr

Zika virus, a mosquito-transmitted flavivirus, has been isolated from sentinel monkeys, mosquitoes, and sick persons in Africa and Southeast Asia. Serologic surveys indicate that Zika virus infections can be relatively common among persons in southeastern Senegal and other areas of Africa, but that Zika virus-associated disease may be underreported or misdiagnosed. In 2007, a large outbreak of Zika virus infection occurred on Yap Island in the southwestern Pacific that infected ≈70% of the island’s inhabitants, which highlighted this virus as an emerging pathogen. The purpose of this study was to investigate and report 3 unusual cases of arboviral disease that occurred in Colorado in 2008.

Clinical and serologic evidence indicate that two American scientists contracted Zika virus infections while working in Senegal in 2008. One of the scientists transmitted this arbovirus to his wife after his return home. Direct contact is implicated as the transmission route, most likely as a sexually transmitted infection.

Visit our website to listen: https://www2c.cdc.gov/podcasts/player.asp?f=7106489