Development of an on-chip detection of Zika virus and antibodies simultaneously using array of nanowells Touyana Semenova, MD, PhD¹, Alexandria Voigt, BS¹, William Donelan, PhD¹, Alek Aranyos, BS¹, Janet Yamamoto, PhD¹, Mobeen R. Rathore, MD², and Cuong Q. Nguyen, PhD^{1,3,4} ¹Department of Infectious Diseases and Immunology, College of Veterinary Medicine, ²University of Florida Center for HIV/AIDS Research, Education and Service (UF CARES), ³Department of Oral Biology, College of Dentistry, ⁴Center of Orphaned Autoimmune Diseases, University of Florida, Gainesville Florida, USA. Address correspondence: Cuong Q. Nguyen, PhD Department of Infectious Diseases and Immunology PO Box 110880, College of Veterinary Medicine University of Florida, Gainesville, Florida 32611-0880 USA Telephone: 352-294-4180, Fax: 352-392-9704

ABSTRACT

24

25

26

27

28

29

30

31

32

33

34

35

36

37

38

39

40

41

42

43

44

45

46

47

48

Zika virus (ZIKV) infections are an emerging health pandemic of significant medical importance. ZIKV appeared recently in the Americas from Africa via the South Pacific. The current outbreak has garnered attention by exhibiting unique characteristics of devastating neurodevelopmental defects in newborns of infected pregnant women. Current guidelines for ZIKV diagnostics developed by the Center of Diseases Control and Prevention (CDC) consist of nucleic acid testing, plague reduction neutralization test (PRNT), and a serologic test for IgM detection. To better accommodate and comply with these guidelines, we developed a simultaneous on-chip detection of ZIKV and anti-ZIKV antibodies using an array of nanowells. Using on-chip microengraving, we were able to detect anti-ZIKV antibodies and their immunoglobulin isotypes. In parallel, applying on-chip real-time PCR with epifluorescence microscopy, we were able to quantify ZIKV viral load as low as one copy. To test clinical samples of patients at the postconvalescent stage, we analyzed samples from 8 patients. The on-chip nanowells could effectively identify antibodies that reacted against ZIKV envelope protein and their isotypes with high sensitivity and specificity. The small sample requirement with high specificity and sensitivity and combined molecular and serological tests could potentially be very advantageous and beneficial in accurate detection of Zika infection for better disease monitoring and management. Keywords: Zika virus, microengraving, real-time PCR, on-chip nanowells, diagnoses, serology

INTRODUCTION

49

50

51

52

53

54

55

56

57

58

59

60

61

62

63

64

65

66

67

68

69

70

71

72

73

After the first reported human infection outbreak on Yap island in 2007, Zika virus (ZIKV) spread dramatically in the Pacific Ocean by a larger epidemic in French Polynesia in 2013-2014 with 32,000 estimated infections (1, 2) and subsequent outbreaks on other Pacific Islands and in the Americas (1, 3-7). Ninety-five countries have been classified by the CDC as risk areas for ZIKV transmission, and 47 countries and territories in the Americas reported ZIKV outbreaks during 2015-2016 (3, 8). In recent years, ZIKV became a serious cause for public health due to its teratogenic and neuropathic outcome in infants and neurological disorders such as Guillain-Barré syndrome in adults (1, 4, 7, 9-11). ZIKV is a single-stranded RNA arbovirus (Family Flaviviridae, genus Flavivirus) transmitted by Aedes mosquitos. The ZIKV genome contains 10,941 nucleotides encoding 3,419 amino acids with 5' and 3' non-coding regions (NCR) and one open reading frame. The open reading frame encodes a single polyprotein that is later cleaved into three structural components: capsid (C), precursor membrane (prM), and envelope protein (E); and seven nonstructural (NS) proteins: NS1, NS2A, NS2B, NS3, NS4A, NS4B, and NS5 (12). Hierarchical cluster analysis shows that ZIKV and Dengue virus (DENV) cluster at a higher hierarchical level and that ZIKV is phylogenetically most related to Spondweni virus (12, 13). This separation of the ZIKV cluster from other flavivirus clusters at a similar hierarchical level may play an important role on pathogenesis and tissue tropism despite the similarity of clinical symptoms to other flavivirus infections (13). Contrary to other flaviviruses, ZIKV can be transmitted vertically by sexual contact and intravenous transfusion. Consequently the virus can be present in human aqueous (14), seminal fluid (14, 15), urine (15), vaginal secretions (16), breast milk, amniotic fluid (5, 17), fetal cerebrospinal fluid, cord blood, infant blood at the second day of birth, and placenta (17). Additionally, ZIKV infection has recently been described in posttransplant patients of solid organs and stem cells from asymptomatic infections to meningoencephalitis (18).

75

76

77

78

79

80

81

82

83

84

85

86

87

88

89

90

91

92

93

94

95

96

97

Current guidelines of ZIKV diagnostics developed by the Center of Diseases Control and Prevention (CDC) consist of testing for ZIKV antibodies using the IgM antibody capture enzymelinked immunosorbent assay (MAC-ELISA) followed by validation using plaque reduction neutralization testing (PRNT). Additionally, nucleic acid testing (NAT) should be performed during the first 6 weeks after the onset of symptoms (19). However, final results may be misinterpreted due to ZIKV epidemiological characteristics and diagnostic limitations. In generally ZIKV is characterized by having an asymptomatic course (20, 21), short transient incubation, and viremic periods (3-14 days, median: 6.2 days and 5 days respectively) (22, 23). Viremia may fluctuate depending on samples tested (whole blood, serum, urine, semen, or amniotic fluid) (21, 24). Furthermore, detection of viral RNA can be prolonged in pregnant women and in adults with Guillain-Barré syndrome (25-29). Laboratory results for serological testing of IgM against ZIKV may sometimes be difficult to interpret, especially for pregnant women, due to possible long persistence of IgM against ZIKV (2-4 months) (19, 30). For competent management of infection, the CDC has recommended that it is necessary to concurrently obtain a patient-matched serum specimen for NAT and/or IgM serological tests. Therefore, in this study we proposed the development of a novel diagnostic method based on massively parallel on-chip detection of ZIKV using a modified fluorescent polymerase chain reaction (PCR) and isotypic anti-ZIKV antibodies by microengraving in nanowells. The results indicated that this on-chip molecular biology test exhibited significant sensitivity for detection of low viral copy number. Simultaneously, the microengraving serological test was able to identify anti-ZIKV antibodies and their isotypes. Therefore, utilization of on-chip detection using nanowells might provide a significant technological advantage which benefits the monitoring and clinical management of ZIKV infection.

MATERIAL AND METHODS

98

99

100

101

102

103

104

105

106

107

108

109

110

111

112

113

114

115

116

117

118

119

Patient materials. Serum and plasma samples were purchased from Boca Biolistics (Pompano Beach, USA). These clinical samples were tested for ZIKV RNA by real-time PCR (COBAS Z480 PCR instrument and Light Mix Modular Zika Virus PCR Real Time, Roche, Switzerland) and for IgG/IgM antibody reactive with ZIKV antigens by Euroimmun IgG and IgM EIA (Euroimmun AG, Germany). Furthermore, samples were tested for West Nile virus (WNV), Chikungunya virus (CHIKV), and Dengue virus (DENV) RNAs and virus-specific antibodies by real-time PCR and ELISA according to the manufacturer's instructions (InBios International, Inc., Seattle, WA). All procedures were reviewed and approved by the University of Florida Institutional Review Board. Fabrication of arrays of nanowells. Sylgard 184 silicone elastomer base (polydimethyl-siloxane, PDMS) and curing agent with a 10:1 weight ratio was combined and mixed vigorously. The mixture was degassed under vacuum for 2 hrs and poured into a custom-built aluminum mold containing a silicon wafer with a patterned array of posts. The mixture was set to cure for 2 hrs at 80°C and adhered directly to a 3"×1" glass slide. The pattern on the master aluminum mold was transferred to the cured PDMS in bas-relief. In this experiment, a master aluminum mold was used that contained blocks of 7x7 nanowells, 4x4 blocks and 6 columnsx18 rows with nanowell dimensions of 50 µmx50 μm×50 μm for a total of 84,672 nanowells per array. Real-time PCR amplification of ZIKV and DENV. ZIKV RNA was extracted from 140 µl of serum and plasma samples using the Qiamp Viral RNA Mini RNA kit (Qiagen, Hilden, Germany) according to the manufacturer's instructions. Primers

(Forward: 5'd CAGCTGGCATCATGAAGAAYC 3'; Reverse 1: 5'd CACCTGTCCCATCTTTTCTCC 3') and probe (5'd FAM-CYGTTGTGGATGGAATAGTGG-BHQ-1 3') were designed according to the previous study (31) and purchased from LGC Biosearch Technologies (Petaluma, CA). A mastermix contained: 5 μl iTaq universal probes reaction mix, 0.25 μl of iScript reverse transcriptase, 100 nM for each primers, 150 nM for each probes, 2.75 μl of nuclease-free water and 1 ng viral RNA. Mastermix was deposited on a 96 well PCR plate and sealed with PCR plate sealing film. Real-time PCR was performed on the CFX96 Touch real-time PCR detection system (Bio-Rad, CA). A thermal cycling protocol was as followed: reverse transcription at 50 °C for 10 min, polymerase activation and DNA denaturation at 95 °C for 3 min followed by 40 cycles of amplification: denaturation at 95 °C for 15 sec, annealing/extension with plate reading at 60 °C for 30 sec. Similar protocols were used for DENV RNA detection using the FDA-approved CDC DENV-1-4 RT-PCR assay (32) performed on the CFX96 Touch real-time PCR detection system (Bio-Rad, CA).

Plague reduction neutralization test

The presence of neutralizing antibodies was determined as previously described (33). ZIKV Puerto Rico strain PRVABC59 was used for this assay. Patients' samples, positive, and negative controls were titrated with media containing Eagle's Minimum Essential Medium (EMEM) (Corning, NY), 2.5 % fetal bovine serum (FBS, Atlanta Biologicals, GA), and gentamicin at 25µg/mL (Gybco,BRL, NY), and viral stock was added into a 96 well plate and incubated for 1 hour at 37°C with 5% CO₂. Vero cells (ATCC CCL 81, epithelial cells of African green monkey—*Cercopithecus aethiops;* Manassas, VA) were plated in a 96 well plate at 90% monolayer confluency. The growth media was removed from the 96 well plate and patients' samples with viral stock were deposited into the plate with Vero cells. After 48 hours of

146

147

148

149

150

151

152

153

154

155

156

157

158

159

160

161

162

163

164

165

166

167

168

169

incubation at 37°C with 5% CO₂, unabsorbed virus was removed, and methylcellulose overlay medium containing EMEM, 7% NAHCO₃, and gentamicin (25 µg/mL) was added to each well. After 48 hours of incubation at 37°C with 5% CO₂, the plate was stained by crystal violet solution containing crystal violet, methanol, and distilled water, and the plaques were counted. According to the CDC guidelines, (19) a titer <10 is considered as negative, and a titer ≥ 10 is considered positive. On-chip microengraving. Capture slides were coated for 1 hour with anti-mouse Abs (for mouse monoclonal antibodies: anti-ZIKV E clone ZV-2 and anti-flavivirus clone D1-4G2-4-15) or anti-human Abs (human sera and plasmas) as previously described (34, 35). Anti-ZIKV envelope protein ZV-2 was added to the nanowells (NR-50414, BEI Resources) as positive control and anti-flavivirus clone D1-4G2-4-15 served as negative control (NR-50327, BEI Resources). After incubation capture slide was blocked in 3% milk buffer solution for 1 hour, rinsed with PBST, PBS, deionized water, spun dry, and stored at 4°C (36). Ten microliters of serum or plasma sample was deposited onto the nanowells and hybridized with the treated capture slide for 2 hours in a hybridization chamber (Agilent Technologies, CA) at RT. After incubation, the capture slide was processed using Tecan Pro HS 4800 Hybridization Station (Tecan, Männedorf, Switzerland) by adding a mixture of goat anti-mouse IgM-PE, anti-mouse IgG-Alexa Fluor (AF)647, anti-mouse IgA-AF555 (mouse monoclonal antibodies) or goat anti-human IgM-PE, anti-human IgG-AF647, anti-human IgA-AF555 (human sera/plasmas) (SouthernBiotech, AL), and ZIKV envelope protein ectodomain (Protein Science Corporation, CT) conjugated with AF488 using DyLight Antibody Labeling kit (Thermo Scientific, IL). Capture slide was scanned using the Genepix 4400A scanner (Molecular Devices, CA). On-chip real-time PCR. A mastermix was prepared containing 5 µl iTag universal probes reaction mix, 0.25 µl of iScript reverse transcriptase, 100 nM for each primers, and 150 nM for each probes as presented previously, 2.75 µl of nuclease-free water, and 1 µl serum/plasma.

Zika RNA or Zika virus were used as positive controls, and healthy donor sera or no template were used as negative controls. The mastermix was deposited on the nanowell chip. The chip was sealed with the Frame-Seal™ in Situ PCR and Hybridization Slide Chambers and placed on the Eppendorf™ In Situ Block Adapter for Mastercycler™ Thermal Cycler (Eppendorf, Hamburg, Germany) to run one-step realtime-PCR. A thermal cycling protocol was as followed: reverse transcription at 50 °C for 10 min, polymerase activation and DNA denaturation at 95 °C for 3 min followed by 40 cycles of amplification: denaturation at 95 °C for 15 sec, annealing/extension with plate reading at 60 °C for 30 sec. After PCR, the microarray chip was analyzed for detection of signal and quantification of fluorescent intensity using an automated epifluorescence microscope equipped with a phase contrast, motorized stage, 405-nm and 488-nm wavelength filter sets using Nikon NIS-Elements Advanced Research image capture software (Nikon, NY).

Data and statistical analyses.

The mean fluorescent intensity (MFI) for each well with a positive signal were generated using GenePix Pro7 Software (Molecular Devices, CA). NIS-Elements Microscope Imaging Software (Nikon, NY) was used to quantify the MFI of real-time PCR results. Data was analyzed using the unpaired two-tailed Mann-Whitney test (GraphPad Prism, CA) to determine the statistical significance. In all cases, p values ≤ 0.05 were considered significant. Excel (Microsoft, WA) was used to perform regression analysis.

RESULTS

190

191

192

193

194

195

196

197

198

199

200

201

202

203

204

205

206

207

208

209

210

211

212

Microengraving for anti-ZIKV antibodies using the on-chip nanowells.

Diagnosis of ZIKV is typically based on nucleic acid amplification to enumerate the viral load or immunoassays to determine the antibody response to the virus. Nucleic acid amplification and antibody determination are routinely performed separately, since there is no technique that detects both parameters simultaneously. Therefore, we sought to determine if we could perform ZIKV amplification and detect ZIKV-specific Abs using the on-chip nanowells. As a proof-of-concept, we utilized monoclonal anti-ZIKV clone ZV-2, which has been shown to recognize ZIKV envelope protein (ZIKV E) as a positive control and monoclonal anti-flavivirus clone D1-4G2-4-15 as negative control, which has neutralizing ability against ZIKV, but does not bind to the viral envelope protein. Abs were serially diluted at 1:10, 1:50, and 1:100 and 10 µl of undiluted and serially diluted Abs were deposited onto the nanowell chip and hybridized with a capture slide coated with goat anti-human Ig and goat anti-human IgG (H+L). Detection antibodies conjugated with specific fluorochromes were added to the capture slide microarray. As presented in Figure 1A, monoclonal anti-ZIKV clone ZV-2 was able to bind to ZIKV E protein as anticipated. Additionally, the microengraving process was able to detect the IgG and IgA, but not IgM isotypes. Monoclonal anti-flavivirus clone D1-4G2-4-15 which has been shown to not react against the E protein, was negative for E protein, IqA, and IqM, but positive for IqG isotype using on-chip microengraving. Regression analysis showed that anti-ZIKV E dilution was positively correlated with fluorescent intensity (R²=0.7909) (Figure 1B). Therefore, the data indicated on-chip microengraving can be utilized to concomitantly detect ZIKV-specific antibodies and the isotypes present in the sample.

Real-time PCR for ZIKV using the nanowell chip.

214

215

216

217

218

219

220

221

222

223

224

225

226

227

228

229

230

231

232

233

234

235

236

237

Significant progress has been made in molecular detection of viruses. The optimization of detection methods using real-time PCR based assays allows for assays to be performed rapidly and produce specific, sensitive, and reproducible results for virus detection. However, the real-time PCR based assays still have limitations, particularly the sample volume and threshold of detection. To address these specific challenges, we performed the real-time PCR assay on the nanowell chip with a limited number of viral copies and volume. Undiluted ZIKV samples were serially diluted in plasma of healthy control and different viral copy numbers (1, 10, 100, or 1000) were deposited into the each individual nanowell predicted by the Poisson distribution. A mastermix containing the reverse transcriptase, polymerase, primers, and probes specific to ZIKV were added to the nanowell chip and placed on a standard laboratory PCR instrument. Using epifluorescence microscopy, the chip was imaged to examine the change in fluorescent signals based on the change in viral loads (Figure 2A). As indicated in Figure 2B, the nanowell chip was able to capture fluorescent intensity from 1 copy to 1000 copies and correlated strongly by linear regression analysis (R²=0.9631). DENV was not detected using ZIKV primer sets (data not shown). The result demonstrates the sensitivity and specificity of the nanowell chip for ZIKV detection, and it can be used to quantify the exact viral load based on fluorescent intensity.

Combining microengraving and real-time PCR for clinical samples.

We selected eight patients that originated from the Dominican Republic in the post-convalescent period of ZIKV infection. Demographic characteristics of analysed patients are shown in **Table 1**. ZIKV-infected patients were initially exanimated for developing a humoral response against ZIKV and presented high serological titer (IgG 8.54 – 20.4) using ELISA (**Table 2**). Further testing was performed to detect the presence of neutralizing antibody titer against ZIKV using PRNT. All patients had a neutralizing antibody titer, and six patients (45%) had a high score of neutralizing antibody titer against ZIKV (from 1:600 to 1:4000). All samples

were negative for ZIKV using conventional real-time PCR. To detect and characterize antibody profiles in patients, we analyzed samples using the microengraving serological assay. As illustrated in **Figure 3**, capture slides coated with anti-human Ig/IgG (H+L) were hybridized with nanowell chips containing 10 µl of sample. After hybridization, the micrographs were processed for IgG, IgM, IgA antibodies and ZIKV envelope protein. All patient samples were positive for IgG, IgA, and ZIKV E protein and negative for IgM (**Figure 2**, **Table 2**). Since the samples were negative for ZIKV using conventional real-time PCR, we spiked the serum samples with ZIKV at different dilutions (1, 10, 100, and 1000 copies per nanowells predicted by Poisson distribution). The spiked samples were subjected to real-time PCR using the nanowells, and similar to **Figure 2**, the spiked samples were positive for ZIKV with a positive correlation (data not shown). These results demonstrate the ability to simultaneously detect ZIKV-specific antibodies and RNA using the nanowell chip assay.

DISCUSSION

There are currently 5 serological assays and 14 molecular assays for ZIKV detection with FDA emergency use authorization (EUA). The serological assays measure the IgM response against either E or NS1 protein for acute infection. The molecular assays amplify the E, NS1, NS3, or prM genes of the virus to quantify the presence of ZIKV using the real-time PCR platform (37). These diagnostic assays are essential for proper understanding of the transmission and clinical disease manifestations of Zika infection. However, the current serological and molecular tests are performed separately on different technical platforms which require larger sample volumes, is labor intensive, time consuming, and it carries a high degree of technical variability. In this study, to circumvent the inherent technical challenges, we developed an on-chip method for detection of ZIKV and anti-ZIKV antibodies with specific isotypes simultaneously using an array of nanowells. Our results demonstrate that by using microengraving as a serological test and real-time PCR as a molecular test, we were able to simultaneously detect isotypic antibodies against E protein and viral load of ZIKV at low copy numbers.

The ZIKV MAC-ELISA (CDC) serological assays use the recombinant, non-infectious ZIKV-like particles as capture antigen and demonstrated 94% positive agreement with PRNT and 83-100% positive agreement with peer-reviewed studies independently assessing the performance of the assays (37). Similar to the on-chip microengraving, the ZIKV Detect IgM ELISA (InBios, Seattle, WA, USA) utilizes the E protein as capture antigen. This assay showed 100% positive agreement with PRNT and 100% positive agreement with peer-reviewed studies (37). Using fluorochrome-conjugated ZIKV E protein as detection, our on-chip microengraving demonstrated significant specificity in which only gold standard anti-ZIKV E protein was detected, whereas negative control anti-flavivirus clone D1-4G2-4-15 performed as expected. To further test the on-chip microengraving, we analyzed plasma and serum samples from 8

post-convalescent patients (≥ one-year post infection). As demonstrated, these patients exhibited positive PRNT at various titers and due to the extended duration of time post-acute infection, these samples were all positive for IgG and IgA, but negative for IgM by standard ELISA. Similarly, our result demonstrates sera were positive for anti-ZIKV E, IgA and IgG, and negative for IgM.

The challenges of molecular assays to detect ZIKV RNA are that a low limit of detection can result in a high proportion of false negative results and testing conditions (samples, assay reagents, and experimental design) can compromise the sensitivity and specificity of detection. Available real-time PCR kits have varying limits of detection from 30 to 1000 copies/mL (38-40). The Trioplex RT-PCR (CDC) assay detects the ZIKV E gene using TaqMan real-time PCR with 100% positive agreement and a limit of detection of 1.93 X 10⁴ genome copy equivalents (GCE)/ml. Using the same primers and probes with modified reporters and quenchers, the onchip assays were able to measure detectable fluorescent signals at one copy and showed positive correlation to 10⁴ copies. To examine the specificity of the test, DENV samples were also tested, and no signals were detected (data not shown). Therefore, real-time PCR on the chip in conjunction with epifluorescence microscopy exhibited remarkable sensitivity and specificity that are comparable or surpass current molecular assays in the market.

Due to the unfavorable consequences of ZIKV infection for populations living in areas endemic for ZIKV, and especially for pregnant women with a high risk of fetal abnormalities, rapid, highly specific, and sensitive diagnostic assays are urgently needed. The rapid outbreak and severe clinical manifestations launch a great urgency in the development of diagnostic tests for ZIKV. Zhang and colleagues proposed using a simultaneous serological assay on a nanostructured plasmonic gold platform for the detection of IgG, IgA, and IgG avidity against ZIKV and DENV-2 antigens in serum samples (41). Simultaneous detection of ZIKV, DENV, and

Chikungunya based on the reverse transcription-loop mediated isothermal amplification (RT-LAMP) was proposed (42-44) with possible smartphone imaging (43). This one-step nucleic acid amplification method based on the PCR diagnostic test has many advantages such as rapidity of analysis and utilizing a portable format. Our on-chip detection method is limited by the high cost of instruments and technical expertise required. Part of our future study is to develop a portable on-chip process that is affordable and applicable in the field to monitor acute Zika infection.

ACKNOWLEDGMENTS

The research was supported in part by Florida Department of Health, Biomedical Research

Program (7ZK12, CQN) and the National Institute of Health (R21AI130561, CQN). The funders
had no role in study design, data collection and interpretation, or the decision to submit the work
for publication. The authors report no conflicts of interest. The authors have no competing
financial interests regarding the subjects in this study

317

318

319

320

321

322

323

324

325

326

327

328

329

330

331

332

333

334

335

336

337

338

339

340

341

REFERENCES Cao-Lormeau VM, Roche C, Teissier A, Robin E, Berry AL, Mallet HP, Sall AA, Musso 1. D. 2014. Zika virus, French polynesia, South pacific, 2013. Emerg Infect Dis 20:1085-6. 2. Mallet HP VA MD. 2015. Bilan de L'épidémie a virus Zika en Polynésie Française, 2013-2014. Bulletin D'linformation Sanitaires, Epidemiologiques et Statistiques 13:1-5. CDC. 2016. All countries & territories with active Zika virus transmission 3. https://www.cdc.gov/zika/geo/active-countries.html 4. Demir T, Kilic S. 2016. Zika virus: a new arboviral public health problem. Folia Microbiol (Praha) 61:523-527. 5. ECfDPa C. 2015. Rapid risk assessment. Zika virus epidemic in the Americas: potential association with microcephaly and Guillain-Barré syndrome. 6. Gatherer D, Kohl A. 2016. Zika virus: a previously slow pandemic spreads rapidly through the Americas. J Gen Virol 97:269-73. 7. Musso D, Nilles EJ, Cao-Lormeau VM. 2014. Rapid spread of emerging Zika virus in the Pacific area. Clin Microbiol Infect 20:0595-6. 8. ECfDPa. C. 2016. Communicable disesases threat report week 6. http://ecdc.europa.eu/en/publications/Publications/communicable-disease-threats-report--13-feb-2016.pdf. Martines RB, Bhatnagar J, Keating MK, Silva-Flannery L, Muehlenbachs A, Gary J, 9. Goldsmith C, Hale G, Ritter J, Rollin D, Shieh WJ, Luz KG, Ramos AM, Davi HP, Kleber de Oliveria W, Lanciotti R, Lambert A, Zaki S. 2016. Notes from the Field: Evidence of Zika Virus Infection in Brain and Placental Tissues from Two Congenitally Infected Newborns and Two Fetal Losses--Brazil, 2015. MMWR Morb Mortal Wkly Rep 65:159-60. 10. Mlakar J, Korva M, Tul N, Popovic M, Poljsak-Prijatelj M, Mraz J, Kolenc M, Resman

342 Rus K, Vesnaver Vipotnik T, Fabjan Vodusek V, Vizjak A, Pizem J, Petrovec M, Avsic 343 Zupanc T. 2016. Zika Virus Associated with Microcephaly. N Engl J Med 374:951-8. 344 11. Oehler E, Watrin L, Larre P, Leparc-Goffart I, Lastere S, Valour F, Baudouin L, Mallet H, 345 Musso D, Ghawche F. 2014. Zika virus infection complicated by Guillain-Barre syndrome--case report, French Polynesia, December 2013. Euro Surveill 19. 346 347 12. Kuno G, Chang GJ. 2007. Full-length sequencing and genomic characterization of 348 Bagaza, Kedougou, and Zika viruses. Arch Virol 152:687-96. 349 13. Wang A, Thurmond S, Islas L, Hui K, Hai R. 2017. Zika virus genome biology and 350 molecular pathogenesis. Emerg Microbes Infect 6:e13. 351 14. Kodati S, Palmore TN, Spellman FA, Cunningham D, Weistrop B, Sen HN. 2017. 352 Bilateral posterior uveitis associated with Zika virus infection. Lancet 389:125-126. 353 15. Atkinson B, Thorburn F, Petridou C, Bailey D, Hewson R, Simpson AJ, Brooks TJ, 354 Aarons EJ. 2017. Presence and Persistence of Zika Virus RNA in Semen, United 355 Kingdom, 2016. Emerg Infect Dis 23:611-615. 356 16. Murray KO, Gorchakov R, Carlson AR, Berry R, Lai L, Natrajan M, Garcia MN, Correa A, 357 Patel SM, Aagaard K, Mulligan MJ. 2017. Prolonged Detection of Zika Virus in Vaginal 358 Secretions and Whole Blood. Emerg Infect Dis 23:99-101. 359 17. Calvet G, Aguiar RS, Melo ASO, Sampaio SA, de Filippis I, Fabri A, Araujo ESM, de 360 Sequeira PC, de Mendonca MCL, de Oliveira L, Tschoeke DA, Schrago CG, Thompson 361 FL, Brasil P, Dos Santos FB, Nogueira RMR, Tanuri A, de Filippis AMB. 2016. Detection 362 and sequencing of Zika virus from amniotic fluid of fetuses with microcephaly in Brazil: a 363 case study. Lancet Infect Dis 16:653-660. 364 18. Levi ME. 2017. Zika virus: a cause of concern in transplantation? Curr Opin Infect Dis 365 30:340-345. 366 Prevention UCfDCa. 2017. Guidance for US Laboratories Testing for Zika Virus Infection 19. 367 July 24, 2017.

368 20. Duffy MR, Chen TH, Hancock WT, Powers AM, Kool JL, Lanciotti RS, Pretrick M, Marfel M, Holzbauer S, Dubray C, Guillaumot L, Griggs A, Bel M, Lambert AJ, Laven J, Kosoy 369 370 O, Panella A, Biggerstaff BJ, Fischer M, Hayes EB. 2009. Zika virus outbreak on Yap 371 Island, Federated States of Micronesia. N Engl J Med 360:2536-43. 372 21. Schaub B, Vouga M, Najioullah F, Gueneret M, Monthieux A, Harte C, Muller F, Jolivet 373 E, Adenet C, Dreux S, Leparc-Goffart I, Cesaire R, Volumenie JL, Baud D. 2017. 374 Analysis of blood from Zika virus-infected fetuses: a prospective case series. Lancet 375 Infect Dis 17:520-527. 376 22. Lanciotti RS, Kosoy OL, Laven JJ, Velez JO, Lambert AJ, Johnson AJ, Stanfield SM, 377 Duffy MR. 2008. Genetic and serologic properties of Zika virus associated with an 378 epidemic, Yap State, Micronesia, 2007. Emerg Infect Dis 14:1232-9. 379 23. Krow-Lucal ER, Biggerstaff BJ, Staples JE. 2017. Estimated Incubation Period for Zika 380 Virus Disease. Emerg Infect Dis 23:841-845. 381 24. Baud D, Gubler DJ, Schaub B, Lanteri MC, Musso D. 2017. An update on Zika virus 382 infection. Lancet 390:2099-2109. 383 25. Suy A. Sulleiro E. Rodó C. Vázguez É. Bocanegra C. Molina I. Esperalba J. Sánchez-384 Seco MP, Boix H, Pumarola T, Carreras E. 2016. Prolonged Zika Virus Viremia during 385 Pregnancy. N Engl J Med 375:2611-2613. 386 26. Driggers RW, Ho CY, Korhonen EM, Kuivanen S, Jääskeläinen AJ, Smura T, Rosenberg 387 A, Hill DA, DeBiasi RL, Vezina G, Timofeev J, Rodriguez FJ, Levanov L, Razak J, 388 Ivengar P, Hennenfent A, Kennedy R, Lanciotti R, du Plessis A, Vapalahti O. 2016. Zika 389 Virus Infection with Prolonged Maternal Viremia and Fetal Brain Abnormalities. N Engl J 390 Med 374:2142-51. 391 27. Goncalves A, Peeling RW, Chu MC, Gubler DJ, de Silva AM, Harris E, Murtagh M, Chua 392 A, Rodriguez W, Kelly C, Wilder-Smith A. 2017. Innovative and new approaches to 393 laboratory diagnosis of Zika and dengue: a meeting report. J Infect Dis

394 doi:10.1093/infdis/jix678. Terzian ACB, Estofolete CF, Alves da Silva R, Vaz-Oliani DCM, Oliani AH, Brandão de 395 28. 396 Mattos CC, Carlos de Mattos L, Rahal P, Noqueira ML. 2017. Long-Term Viruria in Zika 397 Virus-Infected Pregnant Women, Brazil, 2016. Emerg Infect Dis 23:1891-1893. 398 29. Gonzalez-Escobar G VA, Adams R, Polson-Edwards K, Hinds AQJ, Misir A, et al. 2017. 399 Prolonged Zika virus viremia in a patient with Guillain-Barré syndrome in Trinidad and 400 Tobago. Rev Panam Salud Publica. 401 30. Paz-Bailey G, Rosenberg ES, Doyle K, Munoz-Jordan J, Santiago GA, Klein L, Perez-402 Padilla J, Medina FA, Waterman SH, Gubern CG, Alvarado LI, Sharp TM. 2017. 403 Persistence of Zika Virus in Body Fluids - Preliminary Report. N Engl J Med 404 doi:10.1056/NEJMoa1613108. 405 31. Waggoner JJ, Gresh L, Mohamed-Hadley A, Ballesteros G, Davila MJ, Tellez Y, Sahoo 406 MK, Balmaseda A, Harris E, Pinsky BA. 2016. Single-Reaction Multiplex Reverse 407 Transcription PCR for Detection of Zika, Chikungunya, and Dengue Viruses. Emerg 408 Infect Dis 22:1295-7. 409 32. Administration USFaD. 2012. 510(k) substantial equivalence determination decision 410 summary, K113336, p 22, U.S. Food and Drug Administration, Washington, DC ed. 411 33. Langford MP, Weigent DA, Stanton GJ, Baron S. 1981. Virus plaque-reduction assay for 412 interferon: microplaque and regular macroplaque reduction assays. Methods Enzymol 413 78:339-46. 414 34. Nguyen CQ, Ogunniyi AO, Karabiyik A, Love JC. 2013. Single-Cell Analysis Reveals 415 Isotype-Specific Autoreactive B Cell Repertoires in Sjögren's Syndrome. PLoS ONE 416 8:e58127. 417 35. Esfandiary L. VA, Ketchum J.M., Gupta N., Chan E.K., Stewart C.M., Bhattacharyya I., 418 Sukumaran S., Nguyen C.Q. Development of a highly sensitive single-cell multiplex 419 technology for early detection of Sjogren's Syndrome, p. In (ed),

420 36. Love JC, Ronan JL, Grotenbreg GM, van der Veen AG, Ploegh HL. 2006. A microengraving method for rapid selection of single cells producing antigen-specific 421 422 antibodies. Nat Biotechnol 24:703-7. 423 37. Theel ES, Hata DJ. 2018. Diagnostic Testing for Zika Virus: a Postoutbreak Update. J 424 Clin Microbiol 56. 425 38. L'Huillier AG, Lombos E, Tang E, Perusini S, Eshaghi A, Nagra S, Frantz C, Olsha R, 426 Kristjanson E, Dimitrova K, Safronetz D, Drebot M, Gubbay JB. 2017. Evaluation of 427 Altona Diagnostics RealStar Zika Virus Reverse Transcription-PCR Test Kit for Zika 428 Virus PCR Testing. J Clin Microbiol 55:1576-1584. 429 39. Frankel MB, Pandya K, Gersch J, Siddiqui S, Schneider GJ. 2017. Development of the 430 Abbott RealTime ZIKA assay for the qualitative detection of Zika virus RNA from serum, 431 plasma, urine, and whole blood specimens using the m2000 system. J Virol Methods 432 246:117-124. 433 40. Theel ES, Hata DJ. 2018. Diagnostic Testing for Zika Virus: A Post-Outbreak Update. J 434 Clin Microbiol doi:10.1128/JCM.01972-17. 435 41. Zhang B, Pinsky BA, Ananta JS, Zhao S, Arulkumar S, Wan H, Sahoo MK, Abeynayake 436 J, Waggoner JJ, Hopes C, Tang M, Dai H. 2017. Diagnosis of Zika virus infection on a 437 nanotechnology platform. Nat Med 23:548-550. 438 42. Ganguli A, Ornob A, Yu H, Damhorst GL, Chen W, Sun F, Bhuiya A, Cunningham BT, 439 Bashir R. 2017. Hands-free smartphone-based diagnostics for simultaneous detection of 440 Zika, Chikungunya, and Dengue at point-of-care. Biomed Microdevices 19:73. 441 43. Priye A, Bird SW, Light YK, Ball CS, Negrete OA, Meagher RJ. 2017. A smartphone-442 based diagnostic platform for rapid detection of Zika, chikungunya, and dengue viruses. 443 Sci Rep 7:44778. 444 Lamb LE, Bartolone SN, Tree MO, Conway MJ, Rossignol J, Smith CP, Chancellor MB. 44.

2018. Rapid Detection of Zika Virus in Urine Samples and Infected Mosquitos by

445

Reverse Transcription-Loop-Mediated Isothermal Amplification. Sci Rep 8:3803.

447

FIGURE LEGENDS

449

450

451

452

453

454

455

456

457

458

459

460

461

462

463

464

465

466

467

468

469

470

471

472

Figure 1. Detection of anti-ZIKV using the on-chip microengraving process. A. Samples were deposited in the nanowells. Microengraving was used to determine the reactivity for ZIKV E protein and Iq isotypes using fluorochrome-labeled proteins (ZIKV E: AF488, IqM: PE, IqG: AF488, and IgA: AF555). Representative microarray micrograph of capture slides showed the reactivity of the positive control (monoclonal anti-ZIKV clone ZV-2) and negative control (monoclonal anti-flavivirus antigen clone D1-4G2-4-15 and detection secondary proteins alone). Healthy human donor sera were also used as negative control for ZIKV E protein. **B**. Monoclonal anti-ZIKV clone ZV-2 was diluted at 1/10, 1/50, and 1/100 and subjected to microengraving. MFI for each nanowells at each dilution were determined using GenePix Pro7 Software and presented. The experiments were performed at least five times for consistency. Figure 2. Detection of ZIKV using on-chip real-time PCR. A. Healthy human donor sera (n=5) were used. The actual number of viral copies that deposited in each nanowell can not be controlled, but the proportion of nanowells deposited with different numbers of viral copies of starting viral load can be predicted using the Poisson distribution. The optimal viral copy concentration and real-time PCR reagent volume were determined empirically to obtain the highest percentage of nanowells containing different copy numbers of ZIKV (1, 10, 100, and 1000 copies). Samples were deposited in the nanowells and a mixture of primers, probe, and PCR reagents were added. PCR was performed using a customized Eppendorf thermal cycler and nanowells were imaged using an automated epifluorescence microscope. Representative images of nanowells at different dilutions of virus is presented. **B.** MFIs of each nanowell on the chip were analyzed by Nikon NIS-Elements Advanced Research image capture software

(Nikon, NY) and presented as viral copy number versus MFI values. The experiments were repeated five times for consistency.

Figure 3. Schematic of testing patient sera using the combined on-chip microengraving and real-time PCR. Serum samples (n=8) were deposited into the nanowells. Microengraving was performed to detect anti-ZIKV E antibody, and IgG, IgM, and IgA isotypes. On the same nanowell chip, real-time PCR was conducted to quantify the viral loads. For this study, the serum samples were post-convalescent, therefore negative for ZIKV using conventional and on-chip real-time PCR. As a proof-of-concept, serum samples that were positive for anti-ZIKV E antibody (n=8) were spiked with different dilutions of ZIKV as presented in Figure 2 and on-chip real-time PCR was performed. Both experiments were repeated five times for consistency.

Table 1. Patients' demographic profile

Subject	Gender	Age (y)	Ethnicity	Origin Country	Suspected infection
1 (plasma)	F	35	hispanic	Dominican Republic	ZIKV
2 (plasma)	F	54	hispanic	Dominican Republic	ZIKV
3 (plasma)	F	23	hispanic	Dominican Republic	ZIKV
4 (plasma)	М	62	hispanic	Dominican Republic	ZIKV
5 (plasma)	F	47	hispanic	Dominican Republic	ZIKV
6 (serum)	F	26	hispanic	Dominican Republic	ZIKV
7 (serum)	F	35	hispanic	Dominican Republic	ZIKV
8 (serum)	F	40	hispanic	Dominican Republic	ZIKV

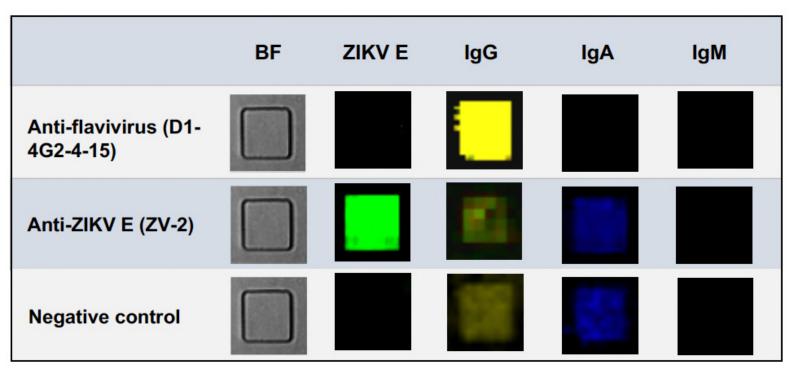
Abbreviations: M= Male; F= Female.; y= years; ZIKV= Zika virus; DENV= Dengue virus.

Table 2. Patients laboratory diagnostic profile

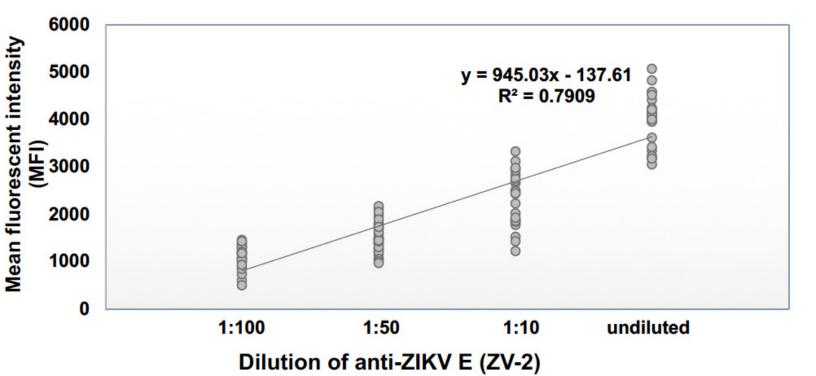
Subject	DiaPro Zika IgG (ISR)	PRNT ZIKV	Microengraving				RT-PCR	RT-PCR
			IgG	IgM	IgA	ZIKVE	ZIKV	DENV
1 (plasma)	8.67	63	+	-	+	+	-	-
2 (plasma)	8.54	100	+	-	+	+	-	-
3 (plasma)	10.83	1500	+	-	+	+	-	-
4 (plasma)	8.97	600	+	-	+	+	-	-
5 (plasma)	10.83	200	+	-	+	+	-	-
6 (serum)	20.4	30	+	-	+	+	-	-
7 (serum)	9.7	4000	+	-	+	+	-	-
8 (serum)	12.4	1200	+	-	+	+	-	-

Abbreviations: ISR= immune status ratio; IgG = immunoglobulin G; IgM = immunoglobulin M; IgA= immunoglobulin A; PRNT= plaque-reduction neutralization test; RT-PCR = realtime-polymerase chain reaction; ZIKV= Zika virus; DENV= Dengue virus; N/A= not applicable.

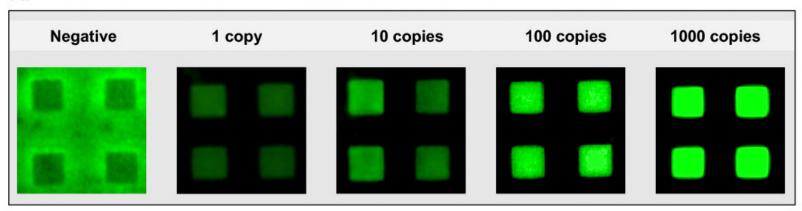
A.



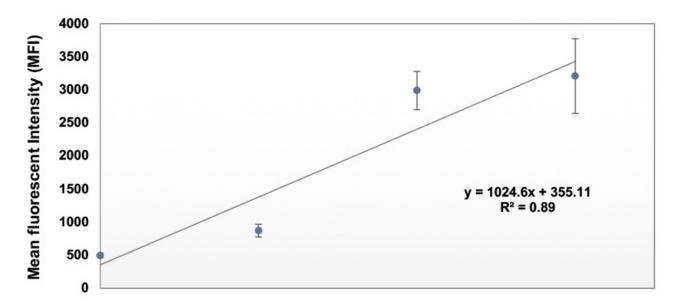




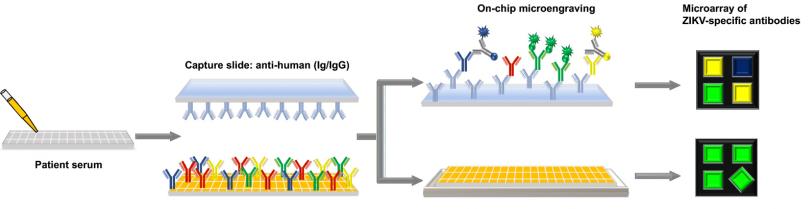




В.



Viral copy number



On-chip real-time PCR

Microarray of detected ZIKV

