Discrepancies in Infectivity of Flavivirus and SARS-CoV-2 Clinical Samples: An Improved Assay for Infectious Virus Shedding and Viremia Assessment

Mizuki Fukuta 1, Co Thach Nguyen 2, Thi Thu Thuy Nguyen 2, Thi Thanh Ngan Nguyen 1,2, Thi Bich Hau Vu 2, Taichiro Takemura 1,3, Le Khanh Hang Nguyen 2, Shingo Inoue 1,3, Kouichi Morita 1,3, Thi Quynh Mai Le 2, Futoshi Hasebe 1,3 and Meng Ling Moi 1,3,4,*

1. Introduction

Virus shedding during acute viral infection has important implications for the transmission and control of infectious disease [1]. In addition to control measures, high blood viral load levels have been associated with greater disease severity, and an understanding of the detailed patterns of virus shedding and viremia is important for determining disease pathogenesis [2–5]. While quantitative real-time polymerase chain reaction (RT-PCR) is the gold standard for virus particle titration, the method detects viral nucleic acid but not the infectious capability of virus particles [6–8]. The infectivity of a virus is typically determined using permissive cells on which the virus has a cytopathic effect (CPE) or leads to plaque formation on virus infection [9,10]. Measuring the viral titer is further challenged if the clinical virus strain does not have a CPE or lead to plaque formation; hence, measurement of infectivity is limited to the detection of CPE forming viruses. As a result of the reliance on detection of a CPE, the infectivity may not be accurately reflected.
During the course of SARS-CoV-2 infection, individuals may remain PCR-positive for several weeks [11]. Early diagnosis is important for managing cases and controlling disease spread. While PCR positivity alone is not associated with infectiousness, accurate reflection of virus infectivity in association with virus shedding, as determined by RT-PCR, is an important consideration of test-based clearance and to accurately determine infectiousness [12]. To determine virus infectivity, conventional methods rely on CPE and plaque formation, and it is possible that these assessment methods may lead to discrepancies in measuring infectious virus titers because they have lower sensitivity than RT-PCR [13]. An ideal detection method should retain the simplicity and practical attributes of the standard virus detection method and demonstrate improved efficiency, reproducibility, shorter assay time, and adaptability to an array of viral infectious diseases.

In the present study, we used flavivirus and SARS-CoV-2 clinical isolates and patient samples to determine the utility of an improved virus infectivity assessment method for virus titration using an improved overlay medium [14–16]. The novel method led to improved infectious virus detection and a higher virus titer compared to the conventional plaque assay. Infectious dengue virus (DENV) and SARS-CoV-2 were detected in some samples, despite negative results using conventional methods. Overall, the modified infectious virus particle detection system using Avicel microcrystalline cellulose [17] is simple and retains the practical features of conventional methods for the detection of infectious virus particles.

2. Materials and Methods

2.1. Cell Lines

Baby hamster kidney cells (BHK-21, Japan Health Science Research Resource Bank), African green monkey kidney Vero cells (Vero 9013, Japan Health Science Research Resource Bank), and FcγRIIA-expressing BHK-21 [18] were used in this study. BHK-21 and Vero cells were cultured in Eagle’s Minimum Essential Medium (E-MEM; Sigma, St. Louis, MO, USA) supplemented with heat-inactivated 10% fetal bovine serum (FBS, Sigma) without antibiotics at 37 °C in 5% CO₂. FcγRIIA-expressing BHK-21 cell lines were cultured in E-MEM supplemented with heat-inactivated 10% FBS and 0.5 mg/mL neomycin (G418; FAA Laboratories GmbH, Pasching, Austria) at 37 °C in 5% CO₂ [19].

2.2. Virus

The following viruses and strains were used in this study: DENV-1 (01-44-1 HuNIID and 99St12A strains); DENV-2 (DHF0663, TLC-30, 08-77, and 00St22A strains); DENV-3 (CH53469 strain); DENV4 (SLMC318 and TVP360 strains); Zika virus (ZIKV; MR766, PRV-ABC9, MRS.OPY_Martinique_PaRi_2015, and H/PF/2013 strains); Japanese encephalitis virus (JEV; OH0566 strain); yellow fever virus (YFV; 17D vaccine strain); SARS-CoV-2 (2019-nCoV/Japan/TY/WK-521/2020, NGS1B, hCoV-19/Japan/QK002/2020, hCoV-19/Japan/TY7-501/2021, and hCoV-19/Japan/TY8-612/2021 strains). Laboratory virus strains were propagated on Vero and BHK cells at 37 °C in 5% CO₂ for 5–7 days or until CPE formation was observed. Cell culture supernatant was then collected, clarified by centrifugation, aliquoted and, stored at −80 °C until use.

2.3. Patient Samples

A total of 27 serum samples obtained from DENV-1 patients during a DENV outbreak in Northern Vietnam in 2017 were used in this study. All patients were confirmed to be infected with DENV-1 by RT-PCR [20] (Supplementary Information). DENV primary and secondary infections were distinguished by DENV immunoglobulin M (IgM) and immunoglobulin G (IgG) enzyme-linked immunosorbent assay (ELISA) (Vircell, Granada, Spain), and RT-PCR. A DENV E/M protein-specific IgM/IgG ratio was used to define primary and secondary infections. In primary infection, IgM levels increase the first day after virus infection, and IgG levels increase 4–5 days after infection. In the secondary infection, because of the presence of pre-existing immunity, IgG levels are higher during the
early acute phase [21]. Primary infection was defined as samples with an IgM/IgG optical density (OD) ratio of $\geq 1.2$, or negative for both DENV IgM and DENV IgG. Secondary infection was defined as samples with an IgM/IgG OD ratio of $<1.2$ or, negative for DENV IgM, and positive for DENV IgG [22,23]. For SARS-CoV-2 patient samples, 53 nasopharyngeal swab samples were used. All samples was confirmed as positive for SARS-CoV-2 by RT-PCR, as described previously [24] (Supplementary Information). The study protocol was approved by the Institutional Review Board at the Institute of Tropical Medicine, Nagasaki University (approval No.: 08061924–7) and the National Institute of Hygiene and Epidemiology (approval No.: IRB-VN01057–45/2016).

2.4. Virus Overlay Medium

Three types of overlay media to facilitate plaque formation for infectious virus detection were used in this study: (1) Avicel medium (Av), (2) Avicel and methylcellulose medium (AvMc), and (3) methylcellulose medium (Mc). To prepare the Av medium, a total of 12 g of Avicel (Asahi Kasei, Tokyo, Japan) and 9.4 g of E-MEM powder (Nissui Pharmaceutical Co., Ltd., Tokyo, Japan) were dissolved in 1 L of double distilled water (DDW) and stirred with a magnetic bar until all reagents were dissolved. To prepare the AvMc medium, a total of 5 g of Mc (Wako Pure Chemical Industries Ltd., Osaka, Japan), 6 g of Avicel, and 9.4 g of E-MEM powder were dissolved in 1 L of DDW. The Mc medium was prepared with 10 g of Mc and 9.4 g of E-MEM powder in 1 L of DDW and, the reagents were then autoclaved, cooled, and stored at 4 °C until use. For complete overlay medium, a total of 10 mL of 100 × L-glutamine (Gibco, Gaithersburg, MD, USA), 20 mL of heat-inactivated FBS (Sigma), and 31.5 mL of 7.5% sodium bicarbonate (Gibco, Gaithersburg, MD, USA) were supplemented. Each mixture was stirred for consistency prior to use.

2.5. Infectious Virus Particle Titration from Laboratory-Established Strains and Clinical Samples

Virus titration was performed using a conventional plaque assay [25]. BHK-21, FcγRIIA-expressing BHK-21, and Vero 9013 cells were used for flavivirus titration, and Vero 9013 cells were used for SARS-CoV-2 virus titration. The cells were seeded in 12-well plates (approximately $5 \times 10^5$ cells per well) with 1 mL growth medium E-MEM supplemented with 10% FBS, and incubated at 37 °C in 5% CO$_2$ overnight to allow the formation of a cell monolayer. Infectious cell culture fluids and patient samples were serially diluted in E-MEM from 1:10 to 1:10$^6$ in 10-fold dilutions. A total of 100 μL of serially diluted sample was added to each well. At least two replicates of each sample were inoculated. The plates were then incubated at 37 °C in 5% CO$_2$ for 60 min. After incubation, 2 mL of overlay medium was added to each well. For titrations of virus stock, the three types of overlay media (Av, AvMc, and Mc) were used. After 4 to 5 days of incubation, the cells were fixed with 4% paraformaldehyde for 1 h at room temperature and stained with 0.25% crystal violet (Wako Pure Chemical Industries). The number of plaques was counted with the naked eye, and plaques were captured using an Automatic Colony Counter PSF-1000 (SK-Electronics Co., Ltd., Kyoto, Japan). Virus titer was defined as the number of plaque-forming units per mL (PFU/mL). The size (mm) of 10 randomly selected plaques was measured using a Fluorescence Microscope BZ–X700 (KEYENCE, United States). Briefly, 10 plaques were randomly selected from one to three wells under the microscope with a magnifying power of 40. Each well was divided into four sections, and 2–3 plaques were randomly chosen from each section for measurement. The diameter line was placed at the major axis and measured according to manufacturer’s instructions. Flavivirus infection experiment was performed in a biosafety level 2 (BSL-2) facility, whereas SARS-CoV-2 infection experiment was performed in the biosafety level 3 (BSL-3) facility at the Institute of Tropical Medicine, Nagasaki University, according to local biosafety guidelines and regulations.

2.6. Data Analysis

The mean and standard deviation (SD) for virus titers and plaque size for each overlay medium in each cell line was determined by using GraphPad (GraphPad Software, La Jolla,
CA, USA). Virus titer results were transformed to base-10 logarithm values for analysis. For comparison of each overlay medium, the p-value was calculated using two-tailed t tests for the viral titer and Mann–Whitney U tests for the plaque size. Statistical analysis was performed using GraphPad Prism version 6.07 (GraphPad Software, La Jolla, CA, USA).

3. Results

3.1. Infectious Virus Titration Using Laboratory-Established Flavivirus Strains and SARS-CoV-2 Strains

Of the 15 flavivirus strains used and among the three cell lines tested, up to one log higher virus titer was detected in 10 (66.7%), 4 (28.5%), and 14 (93.3%) flavivirus strains for BHK, Vero, and FcγR-BHK cells, respectively, when Av medium was used compared to Mc medium (Table 1, Figure 1). The log<sub>10</sub> viral titer (PFU/mL) was significantly different between cultures grown in Av and Mc (p Av vs. Mc = 0.016, p AvMc vs. Mc = 0.0057, Figure 1b). Similarly, plaque size was up to five times larger when Av-containing medium was used (Table 2), and there were significant differences in plaque size when Av-containing medium was used (p Av vs. Mc < 0.0001, p AvMc vs. Mc < 0.0001, Figure 1c). Notably, while the JEV strain (OH0566) did not demonstrate higher virus titers between cell lines and overlay medium (Table 1), the plaque size was larger in Av-containing medium than in Mc medium (Table 2), demonstrating that plaque formation was larger and clearer when an Av overlay medium was used (Figure 1a). In a series of virus titrations using five types of SARS-CoV-2 strains, plaques were not detected in the two strains when Mc was used as the overlay medium (Table 3). Consistent with the results for flaviviruses, SARS-CoV-2 plaque sizes were up to two times larger when Av-containing medium was used compared to Mc medium (Table 3, p AvMc vs. Mc < 0.05).

Table 1. Virus titers of 15 flavivirus strains as determined in BHK-21, Vero, and FcγR-expressing BHK cell lines.

| Cell Lines | BHK-21 | Vero | FcγR-Expressing BHK-21 |
|------------|--------|------|------------------------|
| Viral Strain | Av<sup>a</sup> | AvMc<sup>b</sup> | Mc<sup>c</sup> | Av | AvMc | Mc | Av | AvMc | Mc |
| DENV-1 01-44-1 HuNIID | 6.4 | 6.6 | 4.1 | 3.3 | 4.4 | 3.8 | 6.5 | 6.1 | 5.7 |
| DENV-1 99St12A | 4.4 | 5.5 | 2.5 | 4.5 | 4.7 | 4.8 | 4.9 | 6.1 | 4.6 |
| DENV-2 DF0663 | 5.4 | 5.1 | 4.4 | 5.4 | 5.6 | 5.0 | 5.9 | 5.8 | 5.2 |
| DENV-2 TLC-30 | 5.0 | 4.8 | 4.5 | 5.0 | 4.7 | 5.0 | 4.5 | 4.8 | 4.0 |
| DENV-2 08-77 | 4.9 | 4.3 | 5.2 | 5.6 | 5.4 | 5.6 | 6.1 | 5.8 | 5.2 |
| DENV-2 00St22A | 4.7 | 5.6 | 5.7 | 6.3 | 6.0 | 6.0 | 6.6 | 5.6 | 5.8 |
| DENV-3 CH53469 | 5.8 | 6.7 | 5.4 | 4.6 | 4.3 | 4.1 | 5.9 | 6.9 | 5.8 |
| DENV-4 SLMC318 | 5.7 | 5.6 | 4.9 | 5.4 | 4.1 | 4.4 | 6.1 | 6.0 | 4.3 |
| DENV-4 TVP360 | 4.7 | 5.1 | 3.4 | 4.8 | 4.2 | 4.6 | 4.9 | 5.0 | 3.9 |
| ZIKV MR766 | 6.2 | 6.8 | 6.9 | 8.2 | 7.6 | 7.1 | 6.9 | 7.4 | 6.1 |
| ZIKV PRVABC59 | 5.5 | 5.9 | 5.0 | 7.1 | 6.8 | 6.7 | 5.1 | 6.1 | 4.4 |
| MRS_OPY_Martinique_PaRi_2015 | 5.8 | 6.1 | 5.6 | 7.1 | 6.9 | 6.8 | 5.9 | 6.2 | 4.9 |
| ZIKV H/PF/2013 | 6.5 | 6.7 | 5.9 | 7.4 | 7.2 | 6.2 | 6.8 | 6.7 | 5.7 |
| JEV OH0566 | 7.8 | 7.8 | 7.9 | 7.6 | 7.7 | 7.5 | 7.9 | 7.7 | 7.7 |
| YFV 17D | 6.6 | 6.5 | 5.2 | 5.8 | 6.3 | 6.5 | 6.9 | 6.6 | 6.6 |

<sup>a</sup> Av represents Avicel medium, <sup>b</sup> AvMc represents Avicel and methylcellulose mixed medium, <sup>c</sup> represents methylcellulose medium. Plaque size was determined by random selection of 10 plaques, and values are mean +/- s.d. of each plaque diameter.
Figure 1. Virus titers of laboratory flavivirus strains and clinical DENV-1 and SARS-CoV-2 as determined using plaque assay. (a) Laboratory-established DENV-2, DENV-4, and ZIKV strains were diluted from 1:10 to 1:10^6 and inoculated onto Vero cell monolayer in 12-well plates. Plaque formation differed significantly by number (b) and size (c) at the same dilution for each virus between Avicel medium (Av) and methylcellulose (Mc). (d) Infectious virus titers of six clinical samples obtained from DENV-1 and SARS-CoV-2 patients were determined using Avicel-methylcellulose medium (AvMc) and methylcellulose medium (Mc). The mean of the DENV plaque size for each overlay media (mean ± SD). Each bar shows the mean value, and the error bar represents standard deviation. Significance was determined by unpaired Student’s t test, ns indicates “not significant”, * p < 0.05, ** p < 0.01 and *** p < 0.001.
Table 2. Average plaque size (mm) of 15 flavivirus strains using BHK-21, Vero, and FcyR-expressing BHK cell lines.

| Cell Lines   | BHK-21 | Vero | FcyR-Expressing BHK |
|--------------|--------|------|--------------------|
| Virus Strain | Av      | AvMc | Mc     | Av    | AvMc | Mc    | Av    | AvMc | Mc    |
| DENV-1 01-44-1 HuNIID | 0.8 ± 0.2 | 0.8 ± 0.4 | 0.3 ± 0.1 | 0.3 ± 0.2 | 0.3 ± 0.2 | 0.3 ± 0.1 | 1.1 ± 0.3 | 0.4 ± 0.1 | 0.4 ± 0.1 |
| DENV-1 99St12A | 1.0 ± 0.3 | 1.0 ± 0.3 | 0.5 ± 0.1 | 0.5 ± 0.2 | 0.5 ± 0.2 | 0.5 ± 0.1 | 0.9 ± 0.2 | 1.0 ± 0.2 | 0.2 ± 0.0 |
| DENV-2 DHF0663 | 1.1 ± 0.3 | 1.1 ± 0.2 | 0.2 ± 0.1 | 1.1 ± 0.3 | 1.5 ± 0.3 | 0.7 ± 0.2 | 1.0 ± 0.4 | 1.1 ± 0.7 | 0.5 ± 0.1 |
| DENV-2 TLC-30 | 1.7 ± 0.5 | 1.8 ± 0.6 | 0.5 ± 0.1 | 1.2 ± 0.3 | 1.0 ± 0.5 | 0.6 ± 0.1 | 0.8 ± 0.1 | 1.3 ± 0.1 | 0.3 ± 0.0 |
| DENV-2 08-77 | 0.4 ± 0.1 | 0.6 ± 0.1 | 0.5 ± 0.1 | 1.0 ± 0.5 | 0.6 ± 0.2 | 0.4 ± 0.1 | 0.5 ± 0.1 | 1.4 ± 0.2 | 0.3 ± 0.0 |
| DENV-2 05S22A | 0.6 ± 0.1 | 0.8 ± 0.2 | 0.3 ± 0.1 | 1.2 ± 0.4 | 0.9 ± 0.4 | 0.6 ± 0.2 | 0.7 ± 0.2 | 1.5 ± 0.3 | 0.3 ± 0.1 |
| DENV-3 CH53469 | 1.5 ± 0.5 | 1.8 ± 0.7 | 0.5 ± 0.1 | 1.3 ± 0.5 | 0.6 ± 0.2 | 0.2 ± 0.0 | 1.9 ± 0.8 | 1.7 ± 0.8 | 0.4 ± 0.1 |
| DENV-4 SLMC318 | 0.9 ± 0.2 | 0.7 ± 0.4 | 0.5 ± 0.1 | 0.6 ± 0.1 | 0.7 ± 0.3 | 0.4 ± 0.1 | 0.7 ± 0.1 | 0.4 ± 0.3 | 0.4 ± 0.1 |
| DENV-4 TVP360 | 0.9 ± 0.2 | 1.8 ± 0.6 | 0.4 ± 0.1 | 0.9 ± 0.6 | 0.8 ± 0.2 | 0.8 ± 0.3 | 1.0 ± 0.3 | 1.6 ± 0.3 | 0.3 ± 0.1 |
| ZIKV MR766 | 0.8 ± 0.2 | 1.1 ± 0.6 | 0.4 ± 0.1 | 2.2 ± 0.0 | 1.7 ± 0.4 | 1.1 ± 0.2 | 1.2 ± 0.4 | 1.5 ± 0.6 | 0.3 ± 0.0 |
| ZIKV PRVABC59 | 0.4 ± 0.1 | 0.6 ± 0.2 | 0.5 ± 0.1 | 1.2 ± 0.5 | 1.3 ± 0.3 | 0.5 ± 0.1 | 0.4 ± 0.1 | 0.5 ± 0.1 | 0.4 ± 0.2 |
| MRSOPY_Martineque_PaRi_2015 | 0.6 ± 0.1 | 0.8 ± 0.2 | 0.4 ± 0.1 | 1.3 ± 0.6 | 1.2 ± 0.4 | 0.5 ± 0.2 | 0.5 ± 0.2 | 0.6 ± 0.1 | 0.2 ± 0.1 |
| ZIKV H/PF/2013 | 0.4 ± 0.1 | 0.8 ± 0.3 | 0.4 ± 0.1 | 1.4 ± 0.9 | 1.4 ± 0.6 | 0.9 ± 0.3 | 0.6 ± 0.2 | 0.8 ± 0.3 | 0.2 ± 0.0 |
| JEV OH566 | 0.2 ± 0.0 | 0.6 ± 0.1 | 1.0 ± 0.3 | 1.4 ± 0.4 | 1.5 ± 0.4 | 0.8 ± 0.1 | 1.3 ± 0.9 | 0.7 ± 0.3 | 0.3 ± 0.1 |
| YFV 17D | 1.2 ± 0.5 | 0.6 ± 0.1 | 0.3 ± 0.1 | 0.3 ± 0.1 | 0.7 ± 0.2 | 0.3 ± 0.1 | 1.4 ± 0.2 | 0.8 ± 0.3 | 0.4 ± 0.1 |

a Av represents Avicel medium, b AvMc represents Avicel and methylcellulose mixed medium, c represents methylcellulose medium. Plaque size was determined by random selection of 10 plaques, and values are mean +/- s.d. of each plaque diameter.

Table 3. Virus titers and plaque size of four SARS-CoV-2 strains using Vero cells.

| SARS-CoV-2 Strain | Virus Titer (Log_{10} PFU/mL) Vero | Plaque Size |
|-------------------|----------------------------------|-------------|
|                   | AvMc a                           | Mc b        | AvMc a | Mc b |
| 2019-nCoV/Japan/TY/TK521/2020 | 4.9 | 4.8 | 1.4 ± 0.5 | 0.6 ± 0.1 |
| NGS1B             | 5.5 | 4.6 | 1.3 ± 0.5 | 0.6 ± 0.1 |
| hCoV-19/Japan/QRK002/2020 (Alpha) d | 3.7 | ND c | 0.7 ± 0.2 | ND |
| hCoV-19/Japan/TYM-501/2021(Gamma) | 3.6 | ND | 0.8 ± 0.1 | ND |
| hCoV-19/Japan/TYM-612/2021 (Beta) | 3.5 | 3.2 | 0.7 ± 0.1 | 0.3 ± 0.1 |

a AvMc indicates Avicel and methylcellulose mixed medium, b Mc indicates methylcellulose medium, c indicates that plaques were not detected, d indicates the WHO label for SARS-CoV-2 variants of concern [26]. Plaque size was determined by random selection of 10 plaques and values are mean +/- s.d. of each plaque diameter.

3.2. Dengue Virus Titers in Serum Samples from Dengue Patients and SARS-CoV-2 Titers in Nasopharyngeal Swab Samples from COVID-19 Patients

A total of 27 samples were obtained from patients with DENV-1 infection, and virus titers in each clinical sample were determined using three different cell lines and overlay media (Table 4). In these patient samples, clear plaques were observed in 20 (74%) samples using Av or AvMc medium. In contrast, when Mc overlay medium was used, plaque formation was observed in only two samples (7%) (Table 4, Figure 1d). Using a total of 53 COVID-19 patient nasopharyngeal swab samples, clear plaques were detected in 11 (20.7%) samples using Av overlay medium. In comparison, nine (16.9%) COVID-19 samples exhibited plaques using Mc medium (Table 5, Figure 1d). The log_{10} virus titer (PFU/mL) differences between Av-containing media and Mc medium were statistically significant (p AvMc vs. Mc < 0.0001), the detection limit of virus titer by plaque assay was log_{10} 2.0 PFU/mL. While a total of 42 COVID-19 samples were positive for the SARS-CoV-2 genome by RT-PCR, the samples were negative for plaque formation (Supplementary Table S1). These results were consistent with the findings of other investigators that virus genome levels do not reflect infectious virus particle levels [27]. In samples obtained from DENV-1 and COVID-19 patients, plaque sizes were consistently larger and viral titer was higher when Av-containing overlay medium was used compared to Mc overlay medium (Tables 4 and 5). In addition, the SARS-CoV-2 plaque size cultured in samples
from COVID-19 patients was significantly larger when Av-containing overlay medium was used compared to Mc overlay medium \((p < 0.05)\).

Table 4. Virus titers in primary and secondary DENV-1 infection as determined by plaque assay using Avicel and methylcellulose as the overlay medium.

| Sample Code | Days \(^a\) | Log\(_{10}\) Genome Copies/mL | Dengue ELISA | Virus Titer (Log\(_{10}\) PFU/mL) |
|-------------|-------------|-------------------------------|--------------|---------------------------------|
|             |             |                               | OD \(^c\)-IgG | OD-IgM | IgM/IgG OD Ratio | AvMc \(^c\) | Mc \(^d\) |
| Primary Infection | | | | | | | |
| 01-TN-145 | 0 | 9.5 | 0.8 | 0.1 | 0.2 | 6.1 ± 0.03 | ND |
| 01-TN-087 | 1 | 10.2 | 0.5 | 0.1 | 0.2 | 5.4 ± 0.04 | ND |
| 29-HD-011 | 1 | 9.5 | 0.1 | 0.1 | 1.0 | 5.1 ± 0.04 | ND |
| 01-TN-067 | 2 | 7.6 | 0.5 | 0.1 | 0.2 | 6.4 ± 0.1 | 4.49 ± 0.03 |
| 01-TN-076 | 2 | 10.3 | 0.8 | 0.07 | 0.09 | 7.3 ± 0.1 | ND |
| 01-TN-078 | 2 | 9.2 | 0.9 | 0.06 | 0.06 | 6.2 ± 0.2 | ND |
| 01-TN-091 | 2 | 10.5 | 0.4 | 0.2 | 0.4 | 6.6 ± 0.02 | ND |
| 01-TN-040 | 3 | 8.1 | 0.5 | 0.2 | 0.3 | 6.0 ± 0.2 | ND |
| 01-TN-063 | 3 | 6.9 | 0.8 | 0.2 | 0.2 | 6.1 ± 0.07 | ND |
| 04-HD-010 | 7 | 5.2 | 1.8 | 3.5 | ND | ND |
| 08-HD-001 | 8 | 7.6 | 1.0 | 2.6 | 2.5 | ND | ND |
| 09-HD-014 | 8 | 5.4 | 1.6 | 3.3 | 2.0 | ND | ND |

| Secondary Infection | | | | | | | |
| 01-TN-088 | 1 | 10.3 | 1.1 | 0.13 | 8.5 | 6.3 ± 0.2 | ND |
| 01-TN-077 | 1 | 10.4 | 1.0 | 0.05 | 22.1 | 5.0 ± 0.1 | ND |
| 01-TN-106 | 3 | 10.0 | 1.5 | 0.16 | 9.9 | 5.3 ± 0.06 | ND |
| 11-HD-006 | 3 | 7.7 | 2.3 | 0.45 | 5.2 | ND | ND |
| 11-HD-012 | 3 | 8.9 | 1.0 | 0.14 | 7.2 | 5.8 ± 0.1 | ND |
| 29-HD-001 | 3 | 8.9 | 1.4 | 0.30 | 4.7 | ND | ND |
| 01-TN-056 | 3 | 7.6 | 1.5 | 0.13 | 11.5 | 6.1 ± 0.1 | ND |
| 01-TN-061 | 3 | 6.8 | 1.3 | 0.13 | 10.0 | 6.7 ± 0.03 | ND |
| 01-TN-064 | 3 | 7.5 | 1.0 | 0.14 | 6.7 | 6.4 ± 0.08 | ND |
| 11-HD-002 | 4 | 8.3 | 1.1 | 0.16 | 0.2 | 6.0 ± 0.05 | 5.5 ± 0.03 |
| 01-TN-062 | 5 | 7.2 | 1.6 | 0.13 | 12.8 | 6.1 ± 0.1 | ND |
| 01-TN-068 | 5 | 6.5 | 2.0 | 0.39 | 5.1 | 5.8 ± 0.2 | ND |
| 01-PK-119 | 6 | 5.0 | 1.4 | 0.20 | 7.0 | ND | ND |
| 11-HD-004 | 6 | 6.1 | 1.9 | 2.0 | 1.0 | ND | ND |
| 03-HD-014 | 8 | 9.5 | 1.9 | 0.79 | 2.4 | 6.5 ± 0.01 | ND |

\(^a\) Indicates days after onset of disease, \(^b\) Indicates that plaques were not detected, \(^c\) AvMc represents Avicel and methylcellulose mixed medium. \(^d\) Mc represents methylcellulose medium. \(^e\) Indicates optical density. All samples were confirmed positive for DENV by RT-PCR.

Table 5. Virus titers and average plaque size (mm) in COVID-19 patients as determined by using a conventional plaque assay with Avicel and methylcellulose overlay medium.

| Sample No | Virus Load (log\(_{10}\) Genome Copies/mL) | Virus Titer (Log\(_{10}\) PFU/mL) | Plaque Size (Mean Diameters mm ± SD) |
|-----------|-------------------------------------|----------------------------------|----------------------------------|
|           | AvMc | Mc | AvMc | Mc | AvMc | Mc |
| 1 | 10.9 | 4.3 | 4.2 | 0.87 ± 0.34 | 0.29 ± 0.1 |
| 2 | 10.2 | 3.4 | ND \(^a\) | NT | NT |
| 3 | 9.9 | 5.1 | 4.9 | 1.0 ± 0.2 | 0.4 ± 0.07 |
| 4 | 9.7 | 3.8 | 3.6 | 1.3 ± 0.2 | 0.3 ± 0.09 |
| 5 | 9.7 | 3.2 | 3.1 | 1.4 ± 0.2 | 0.38 ± 0.1 |
| 6 | 9.5 | 2.6 | ND | NT \(^b\) | NT |
| 7 | 9.4 | 4.1 | 3.94 | 0.96 ± 0.2 | 0.37 ± 0.1 |
| 8 | 9.21 | 4.0 | 3.81 | 1.1 ± 0.3 | 0.37 ± 0.1 |
| 9 | 9.03 | 3.7 | 4.11 | 1.7 ± 0.3 | 0.4 ± 0.1 |
| 10 | 8.51 | 3.3 | 3.20 | 0.7 ± 0.1 | 0.4 ± 0.1 |
| 11 | 8.44 | 2.9 | 3.23 | 1.4 ± 0.2 | 0.5 ± 0.1 |

\(^a\) Indicates that plaques were not detected, \(^b\) indicates “not tested”, plaque size was determined by random selection of 10 plaques and values are mean +/− s.d. of each plaque diameter.
4. Discussion

Detection of infectious particles remains a challenge as live virus particles decrease concurrently with decreased virus shedding or viremia. In this study, higher virus titers and clearer plaque formation were detected in flavivirus and SARS-CoV-2 laboratory-established strains and clinical samples by a conventional plaque titration method using an improved overlay medium. As high virus titers have been associated with severe clinical outcomes [2–5], assessing infectious virus titers that better reflect the infectivity of the virus is important for determining disease pathogenesis. In addition, assessing infectivity levels that better reflect the infectiousness is important for control measures, particularly in SARS-CoV-2 infection, to accurately determine transmission capability [27].

Plaque formation using Av has also been reported for enterovirus [15], Rift Valley fever virus [16], and influenza virus [14]. However, in previous studies, focus formation was detected using the immunostaining method [28–30]. In the present study, a conventional plaque assay was used for all four DENV serotypes, ZIKV, JEV, and SARS-CoV-2, at various inoculation doses and in clinical samples. In addition to clearer virus plaque formation, higher virus titers were detected in both laboratory-established strains and patient samples. In the detection of plaque formation, the overlay medium prevents convectional flows in the media, allowing viruses to attach to the bottom of wells and facilitate plaque formation [14]. In this context, Avicel may be a better medium than Mc in preventing convectional flows to allow better plaque formation and, in turn, higher virus titers and larger plaque size. In some flavivirus strains, the virus titer was equal or higher under Mc overlay compared to that of Av-containing medium. This was due to the speed of virus propagation. For the strain that has high propagation speed, such as JEV and ZIKV, the plaque formation is well facilitated by an Mc overlay that is comparable to Avicel overlay. In addition, the Av medium has an advantage in the determination of plaque formation in clinical samples, as some clinical virus strains may not form clear plaques. While plaque formation may be dependent on infectivity in different cell lines, in each of the cell lines tested, larger plaques formed when Av-containing overlay medium was used compared to Mc overlay medium. In this context, clearer plaques led to the detection of higher virus titers in clinical samples, and thus, the method may better reflect virus infectivity as compared to other conventional methods. In this study, the plaque assay was used rather than immunostaining methods because viral infectivity can be detected directly with the naked eye without the staining step; hence, the plaque assay is a widely used method because of its simplicity. Some virus strains formed clearer plaques when Av-containing medium was used, and thus the medium may be suitable for the detection of a range of viral strains using plaque assays. In terms of utility, Av medium is relatively easy to use because of its low viscosity compared to Mc medium, and it can be adapted for high-throughput assays such as the 96-well and 384-well formats. In addition to clearer plaques, the incubation time can be shortened by 1 day due to the increased size and clearer plaques when Av is used. In this study, the cells were fixed after 5 days (ZIKV, JEV, YFV) or 6 days (DENV) post-infection when Mc was used as an overlay medium. In comparison, when Av was used, plaques that were large enough to be observed with the naked eye were confirmed 4 to 5 days post-infection, demonstrating that the improved method required a shorter incubation time compared to conventional medium using methylcellulose.

5. Conclusions

Because of clearer plaque formation, plaques were consistently detected using Avicel overlay medium even when there was no plaque formation using the conventional Mc method. One of the challenges encountered in textbook descriptions of flavivirus viremia levels and SARS-CoV-2 virus shedding is that the conventional titration method uses Mc as an overlay medium. Our results demonstrated that this classical description of virus titer may have resulted in the infectious capacity of these viruses not being robustly evaluated, because of an underestimation of the infectious particle levels. In addition, clearer plaque formation allows the use of a wider range of clinical isolates for plaque-based assays,
including virological and anti-viral drug studies and neutralization antibody titration for flavivirus and SARS-CoV-2 [31–35], which in turn, can lead to better understanding and characterization of various clinical strains. Overall, the study results indicated that the improved virus detection method using Av as an overlay medium is rapid and practical and has the potential to serve as a useful tool for laboratory diagnosis of acute flavivirus and SARS-CoV-2 infection in virological studies.

**Supplementary Materials:** The following are available online at https://www.mdpi.com/article/10.3390/ijerph18189845/s1, Table S1, COVID-19 clinical samples with virus titers below detection levels.

**Author Contributions:** M.L.M. conceptualized and designed the research; M.F., C.T.N. and T.T.N.N. performed the experiments; M.F. and M.L.M. interpreted and analyzed the results; M.F. and M.L.M. wrote the original draft of the paper; C.T.N., T.T., T.B.H.V., L.K.H.N., S.I., T.T.N.N., T.Q.M.L. and F.H. assisted in the investigation and sample collection; K.M., F.H. and M.L.M. acquired funding; and M.L.M. directed the research. All authors reviewed and edited the final paper. All authors have read and agreed to the published version of the manuscript.

**Funding:** This research was supported by the Japan Agency for Medical Research and Development (AMED) under the AMED Japan Program for Infectious Diseases Research and Infrastructure (JP21wm0125006, and JP21wm0225018), Research on Emerging and Re-emerging Infectious Diseases (21fk0108109h0003, 21fk0108123h1102, 21wm0225001s0102, 21km0210086h0002), and Nagasaki University.

**Institutional Review Board Statement:** The study protocol was approved by the Institutional Review Board at the Institute of Tropical Medicine, Nagasaki University (approval No.: 08061924-7 and 200409233-3) and the National Institute of Hygiene and Epidemiology (approval No.: IRB-VN01057-45/2016).

**Informed Consent Statement:** Written informed consent was obtained from the patients or a legally authorized representative (for patients below the age of 16). Part of this research uses leftover specimens from SARS-CoV-2 laboratory tests. The use of these specimens is in accordance with the Japanese ethical guidelines and Nagasaki University IRB guidelines (IRB 200409233-3) that provide a waiver of prior informed consent for the research use of existing or stored specimens, whereby the information of the research is made public with the opportunity to opt-out from the study.

**Data Availability Statement:** The data sets generated and/or analyzed during the current study are available from the corresponding authors on reasonable request.

**Acknowledgments:** We would like to thank Minami Isaka and Kirara Iwanaga for their technical support and members of the Department of Virology, Institute of Tropical Medicine, Nagasaki University, for providing technical advice and support. The authors would like to acknowledge Makoto Takeda, Masayuki Saijo, Lim Chang-kweng, and Mutsuo Ito of the National Institute of Infectious Diseases, Japan, for providing the DENV and SARS-CoV-2 strains. We would like to thank the US CDC for providing the ZIKV PRVABC59 strain. The authors are grateful to Koichi Izumikawa, Takeyuki Tanaka, Masato Tashiro, and Ayumi Fujita from the Department of Infection Control and Education Center; Nobuhiro Kanie and Ryosaku Oshiro from the Department of General Internal Medicine and Clinical Infectious Diseases; and Takahiro Takazono, Hirofumi Imamura, and Tatsuro Hirayama from the Department of Respiratory Internal Medicine, Nagasaki University Hospital, Department of Emerging Infectious Disease, Institute of Tropical Medicine, Nagasaki University, and other healthcare staff for the support received during sample collection from hospitals and health institutions in Nagasaki, Japan. The authors are also grateful to the healthcare personnel in Vietnam for their assistance during the dengue clinical sample collection. T.T.N.N. is the recipient of a Japanese government scholarship from the Ministry of Education, Science Sport, and Culture of Japan (Monbukagakusho) through the Japan International Cooperation Agency (JICA) program. M.F. is the recipient of the Taniguchi Scholarship, BIKEN Foundation. M.F. and T.T.N.N. are grateful for the academic support from the Program for Nurturing Global Leaders in Tropical and Emerging Communicable Diseases of the Graduate School of Biomedical Sciences, Nagasaki University.

**Conflicts of Interest:** The authors declare no conflict of interest.
27. Yamada, S.; Fukushima, S.; Kinoshita, H.; Ohnishi, M.; Suzuki, T.; Fujimoto, T.; Saijo, M.; Maeda, K. Assessment of SARS-CoV-2 infectivity of upper respiratory specimens from COVID-19 patients by virus isolation using VeroE6/TMPRSS2 cells. *BMJ Open Respir. Res.* **2021**, *8*, e000830. [CrossRef]

28. Payne, A.F.; Binduga-Gajewska, I.; Kauffman, E.B.; Kramer, L.D. Quantitation of flaviviruses by fluorescent focus assay. *J. Virol. Methods* **2006**, *134*, 183–189. [CrossRef] [PubMed]

29. Okuno, Y.; Fukunaga, T.; Tadano, M.; Okamoto, Y.; Ohnishi, T.; Takagi, M. Rapid focus reduction neutralization test of Japanese encephalitis virus in microtiter system. Brief report. *Arch. Virol.* **1985**, *86*, 129–135. [CrossRef]

30. Kang, W.; Shin, E.C. Colorimetric Focus-Forming Assay with Automated Focus Counting by Image Analysis for Quantification of Infectious Hepatitis C Virions. *PLoS ONE* **2012**, *7*, e43960. [CrossRef]

31. Zhao, H.; To, K.K.W.; Lam, H.; Zhou, X.; Chan, J.F.W.; Peng, Z.; Lee, A.C.Y.; Cai, J.; Chan, W.M.; Ip, J.D.; et al. Cross-linking peptide and repurposed drugs inhibit both entry pathways of SARS-CoV-2. *Nat. Commun.* **2021**, *12*, 1517. [CrossRef]

32. Park, J.G.; Oladunni, F.S.; Chiem, K.; Ye, C.; Pipenbrink, M.; Moran, T.; Walter, M.R.; Kobie, J.; Martinez-Sobrido, L. Rapid in vitro assays for screening neutralizing antibodies and antivirals against SARS-CoV-2. *J. Virol. Methods* **2021**, *287*, 113995. [CrossRef] [PubMed]

33. Amarilla, A.A.; Modhiran, N.; Setoh, Y.X.; Peng, N.Y.G.; Sng, J.D.J.; Liang, B.; McMillan, C.L.D.; Freney, M.E.; Cheung, S.T.M.; Chappell, K.J. An Optimized High-Throughput Immuno-Plaque Assay for SARS-CoV-2. *Front. Microbiol.* **2021**, *12*, 625136. [CrossRef] [PubMed]

34. Gwon, Y.D.; Strand, M.; Lindqvist, R.; Nilsson, E.; Saleeb, M.; Elfsson, M.; Overby, A.K.; Evander, M. Antiviral Activity of Benzavir-2 against Emerging Flaviviruses. *Viruses* **2020**, *12*, 351. [CrossRef] [PubMed]

35. WHO. Update on SARS-CoV-2 Variant Nomenclature, June 2021. Available online: https://www.who.int/docs/default-source/coronaviruse/risk-comms-updates/update60_nomenclature-variants.pdf?sfvrsn=27fc6fa_4 (accessed on 15 September 2021).