

Featuring work from Professor Hiroyuki Noji's Laboratory at the University of Tokyo, which has been involved in single-molecule biophysics of ATP synthase, and the development of digital bioassay and artificial cell reactor technology. Image credit: Akiko Sato.

Mobile imaging platform for digital influenza virus counting

A Mobile Imaging Platform, 'MobiP', for digital bioassays was developed and digital influenza virus counting was demonstrated. This system will pave the way for on-site bioanalysis with single-molecule and single-particle sensitivity, and will accelerate the decentralization of diagnostic tests.

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See Hiroyuki Noji *et al.*,
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Mobile imaging platform for digital influenza virus counting†

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Droplet-based digital bioassays enable highly sensitive and quantitative analysis of biomolecules, and are thought to be suitable for point-of-care diagnosis. However, digital bioassays generally require fluorescence microscopy for detection, which is too large for point-of-care testing. Here, we developed a simple smartphone-based mobile imaging platform for digital bioassays. The size of the mobile imaging platform was $23 \times 10 \times 7$ cm (length \times width \times height). With this platform, a digital enzyme assay of bovine alkaline phosphatase was successfully completed. Digital influenza virus counting—based on a fluorogenic assay for neuraminidase activity of the virus—was also demonstrated. Distinct fluorescence spots derived from single virus particles were observed with the mobile imaging platform. The number of detected fluorescence spots showed good linearity against the virus titer, suggesting that high sensitivity and quantification were achieved, although the imaging with the mobile platform detected 60% of influenza virus particles that were identified with conventional fluorescence microscopy. The lower detection efficiency is due to its relatively lower signal-to-noise ratio than that found with conventional microscopes, and unavoidable intrinsic heterogeneity of neuraminidase activity among virus particles. Digital influenza virus counting with the mobile imaging platform still showed 100 times greater sensitivity than that with a commercial rapid influenza test kit. Virus detection of clinical samples was also successfully demonstrated, suggesting the potential to realize a highly sensitive point-of-care system for influenza virus detection with smartphones.

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1. Introduction

Digital bioassays have seen rapid growth as emerging methods of bioanalysis that enables single-molecule detection sensitivity.^{1–3} In a typical digital bioassay protocol, target molecules that have catalytic activity by themselves or when tagged with enzyme-linked binders such as antibodies are stochastically confined in micron-sized reactors. In these microreactors, enzymes hydrolyze fluorogenic substrates, accumulating fluorescent dye molecules. When the concentration of target molecules is sufficiently low, each reactor stochastically encapsulates none or one of the target molecules. The typical volume of microreactors is extremely small, in the range of femtoliters. As a result, positive reactors entrapping single target molecules accumulate fluorescent dyes in a short period of time, producing a high fluorescence signal that is distinct from the background signal. After binarization of the signal,

the number of reactors giving a signal of ‘1’, for quantification of a target molecule in the analyte, can be counted.

In the early studies of digital bioassays, microreactor systems were composed of a microfabricated polymer sheet of polydimethylsiloxane (PDMS) and a glass coverslip,^{4,5} or a fabricated glass gasket and PDMS sheet.⁶ Later, a system that displayed over a million droplets was developed⁷ and became one of the representative platform devices for microreactor arrays.^{8,9} Currently, microemulsion in fluidic systems is also widely used.^{10,11}

Successful digital bioassays include the single-molecule enzymatic assay, digital PCR,^{12–14} and digital ELISA.^{15–17} These studies achieved drastic improvements in detection sensitivity, by three orders of magnitude or more in comparison with conventional assays. Large numbers of microreactors ensure a wide range of concentrations for quantification.² Since the high sensitivity and quantification of digital bioassays have become well recognized, a wide variety of digital bioassays have been developed, such as isothermal amplification of DNA (LAMP and RCA), and electrochemical detection.^{18–20} Digital counting and analysis of membrane transport proteins, which are hard to quantify using conventional biochemical methods, were also achieved through the development of an arrayed lipid bilayer chamber system

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(ALBiCs).^{21–23} Currently, the concept of digital bioassays is expanded to microreactor-free analytical methods that include single virus counting with interferometric imaging and the tyramide deposition method.^{24–26}

Another merit of digital bioassays is compactness. The high sensitivity of digital bioassays enables analyses with small specimen volumes. Devices for femtoliter droplet array displays or emulsion generation/handling are also compact; the typical size of such devices is a few centimeters. These features of digital bioassays facilitate the realization of compact systems for on-site rapid diagnosis and point-of-care diagnosis. As the signals for detection are generally fluorescence signals, miniaturization of the fluorescence microscopy system is required for on-site diagnostic systems. To implement the requirement, a compact lens-less complementary metal-oxide-semiconductor (CMOS) image sensor directly coupled with the microreactor system has been developed for the digital ELISA, which still remains to be further elaborated.^{27,28}

Smartphones are the most ubiquitous imaging device globally that include a high quality CMOS camera. There are about 42% smartphone users in emerging and developing countries, and approximately 2.5 billion people worldwide will own a smartphone in 2019 (<https://www.statista.com>). Thus, smartphones are highly suitable as an imaging unit in personalized POC diagnostic systems, which could lead to the decentralization of health care management. In addition, when a smartphone-based diagnostic test system is spread, it would allow global health monitoring. To date, there have been several reports of smartphone-based POC diagnostic platforms^{29–31} with demonstrations of several analyses: fluorescence microscopy imaging,^{32,33} cytometry,³⁴ paper-based POC diagnostic tests,^{35,36} biochemical assays with 96 well plates,^{37,38} electrochemical biosensing,³⁹ surface plasmon resonance-based biosensing,^{40,41} and optical density sensing.⁴²

However, only a few demonstrations of digital bioassays on smartphone-based detection systems have been reported. The Nilsson and Ozcan group reported detection of digital rolling circle amplification of DNA *via* smartphone-based imaging.³³ Issadore *et al.* described an integrated fluidic system with a smartphone-based imaging unit for solution handling and detection using the digital ELISA.⁴³ However, a swift and simple digital bioassay that would meet urgent needs for early stage disease detection has not yet been developed. Previously, we reported a digital influenza virus counting (DIViC) method based on a fluorogenic assay of neuraminidase activity of the influenza virus.⁴⁴ This method does not require liquid separation procedures for free antibody washing, and thus would be suitable for on-site detection with a compact imaging unit.

In this study, we aimed to develop a simple type of mobile imaging platform (MobIP) for digital bioassays with a femtoliter reactor array device (FRAD). The MobIP was integrated with waveguide-based total internal reflection illumination and movement stages. This system enabled the digital

enzyme assay of alkaline phosphatase and also DIViC. Our system was approximately 100 times more sensitive than rapid influenza diagnostic tests (RIDTs).

2. Materials and methods

Chemicals

The influenza A virus [A/Puerto Rico/8/1934(H1N1)] was prepared as previously reported.⁴⁵ 2'-(4-Methylumbelliferyl)- α -D-N-acetylneuraminic acid (MUNANA) (Life Technologies, Carlsbad, CA, USA), Fluorinert-FC40 (3M), CYTOP (Asahi-glass, Tokyo, Japan), a highly active alkaline phosphatase recombinant (ALP, Roche, Switzerland), Fomblin Y-LVAC25/6 (Solvay, Brussels, Belgium), 4-MU phosphate (4MUP, Sigma-Aldrich, St. Louis, MI, USA), and RIDT (Quicknavi Flu; Denka Seiken, Tokyo, Japan) were purchased from the respective suppliers.

Femtoliter reactor array device (FRAD)

The fL reactor array device (FRAD) was prepared as follows. A hydrophobic carbon-fluorine polymer (CYTOP; Asahi-glass) was spin-coated on a glass coverslip (24 × 32 mm) at 2500 rpm for 30 s and baked at 80 °C for 10 min and then at 180 °C for 30 min. The thickness of the CYTOP layer was approximately 3.0 μ m. The CYTOP-coated coverslip was spin-coated with a positive photoresist (AZ-4203; AZ Electronic Materials, Luxembourg) at 1000 rpm for 30 s and baked at 100 °C for 5 min. Subsequently, photolithography was carried out with a mask structure with 3 μ m holes, which were separated by 9 μ m. The resist-patterned coverslip was dipped in a developer (AZ300MIF; Merck, Darmstadt, Germany) for 90 s in a sonic bath. The coverslip was dry-etched with O₂ plasma in a reactive ion etching system (RIE-10NR; Samco, Kyoto, Japan) to remove the exposed CYTOP. The substrate was then cleaned and rinsed with acetone and isopropanol to remove the photoresist layer remaining on the substrate. The resulting CYTOP-on-coverslip device had an array of exposed SiO₂ patterns with a diameter of about 4.5 μ m, which each held a water droplet in the digital assay for bovine alkaline phosphatase (bALP) and the influenza virus. The volume of one microreactor was 48 fL.

Smartphone-based imaging unit

The platform structure for assembling the smartphone and other optical/mechanical components was designed using AUTODESK® 123D® DESIGN and prototyped using a 3D printer (Form 2; FormLabs, Somerville, MA, USA). The following components were assembled on the platform: a power LED (365 nm at 3 W; Epileds, Taiwan), a battery-operated micro linear actuator (LA003; Seiko, Tokyo, Japan), a micrometer head (SHS2-13, Sigma), an aspherical lens (C340TMD-A; Thorlabs, Newton, NJ, USA), a long pass filter for emissions (λ = 410 nm, SC-41; Fujifilm, Tokyo, Japan), and a microcomputer system (Genuino 101; Arduino LLC, USA). The aspherical lens was set into the hole beneath which the smartphone camera was inserted. The long pass filter was placed between



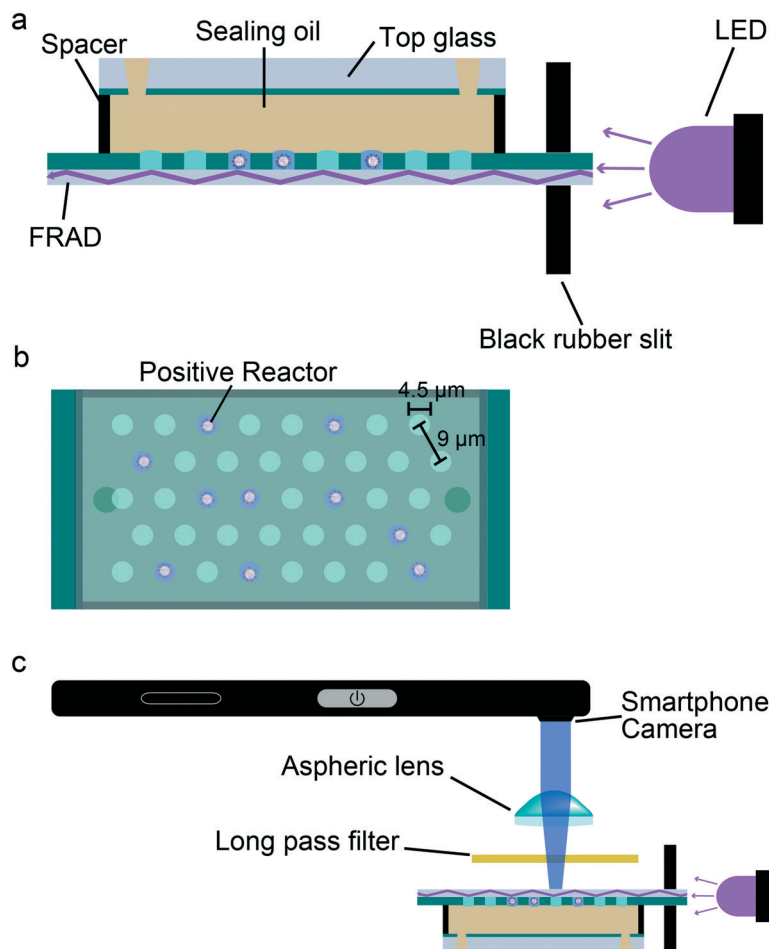


Fig. 1 Mobile imaging platform (MobIP) based on a smartphone. (a) Schematic side view of a flow cell with an integrated waveguide-based total internal reflection fluorescence imaging system. The flow cell is composed of a top glass layer, spacer, and FRAD that has a femtoliter reactor. (b) Schematic top view of the FRAD. (c) Schematic image of the MobIP. The flow cell of the FRAD was inserted below the smartphone (Xperia SO-03J, SONY, Japan). Note that the FRAD was set upside-down (in relation to Fig. 1a) to introduce fluorescence emission from the FRAD towards the camera. Emissions were collected with an aspheric lens set between the long pass filter and smartphone camera.

the aspherical lens and the FRAD (Fig. 1). The micro linear actuator enabled the lateral movement of the sample slide and enabled observation at a 1 cm distance. The micrometer head was used to move the sample slide in the *z* direction for focusing. The power LED and battery-operated linear actuator were operated *via* a Genuino 101 board, which communicated to the smartphone through a Bluetooth Low Energy (BLE) module. The size of the smartphone-based platform was approximately $23 \times 10 \times 7$ cm (Fig. S1†), and the cost was estimated to be <\$360 (USD), excluding the smartphone.

Digital bioassay with the MobIP

All images were acquired using a smartphone mounted onto our platform. The FRAD was illuminated using a power LED, activated through a smartphone app using the Genuino 101 board. The power LED was placed close to the edge of the FRAD and was used as a light source to produce waveguide-based total internal reflection excitation in the glass. The scattered excitation of the LED was blocked through the long

pass filter, creating the very efficient background that is necessary to isolate the weak fluorescence signal.

Image analysis

Photoimages obtained from the smartphone were saved as JPEG files with 8 bit data for blue, green, and red channels. After transferring the image data to a desktop computer, the total fluorescence intensity was determined for analysis, as observed in other digital bioassays. Although the fluorescence intensity of the blue channel was also available for analysis, the total intensity produced less noise and was, therefore, subjected to analysis. The image data were analyzed using ImageJ (NIH, Bethesda, MD, USA).

The signal-to-noise ratio, *S/N*, of the images is defined as: $S/N = (\mu_s - \mu_b) / \sqrt{\sigma_b^2}$, where μ_s is the mean fluorescence intensity of the positive reactor, μ_b is the mean background signal (=signal from empty reactors), and σ_b is the standard deviation of the background signal.



Digital enzyme assay for bovine alkaline phosphatase (bALP)

The flow cell was assembled with the FRAD and CYTOP-coated glass, which were bound using 80 μm double-sided tape. The bALP solution was diluted with an assay buffer (1 M diethanolamine, 2 mM MgCl_2 , 0.1% Tween-20). Next, the fluorogenic substrate 4MUP was added at 1 mM, and the reaction mixture was immediately introduced into the flow cell. Next, fluorinated oil (FC40) was introduced to flush out the excess reaction mixture and form droplets in chamber arrays. Subsequently, FC40 was replaced by Fomblin to prevent the evaporation of the water droplets. After incubation for 20 min at room temperature, fluorescence images were obtained using the smartphone on the MobIP or a conventional inverted fluorescence microscope (Olympus IX83, Olympus Co., Tokyo, Japan) equipped with an sCMOS camera (Andor Neo, Andor Technology, Belfast, Northern Ireland) and objective lens ($\times 20$, N.A. = 0.75).

DIViC procedure in the FRAD

The indicated concentration of influenza virus was mixed with the fluorescence substrate MUNANA in 1 mM reaction buffer (1 M diethanolamine, 4 mM CaCl_2). The reaction mixture was introduced into a flow cell constructed from the FRAD and CYTOP-coated glass. After sealing with FC40 and Fomblin, the FRAD was incubated at 47 $^\circ\text{C}$ on a hotplate. Fluorescence images were obtained using a smartphone on the MobIP or a conventional fluorescence microscope (Olympus IX83, Olympus Co.).

3. Results and discussion

Femtoliter reactor array device (FRAD)

The femtoliter reactor array device (FRAD) was prepared as previously reported.⁴⁴ In short, a fluorinated polymer was spin-coated at a thickness of 3 μm on a coverslip and fabricated to display an array of microholes ($\phi = 4.5 \mu\text{m}$, volume = 48 fL) through photolithography treatment (see the Materials and methods section for details). The micron holes are used as reactors, entrapping water-in-oil droplets under fluorinated oil (FC40). The total number of reactors was 6.0×10^6 on a $20 \times 20 \text{ mm}$ square surface. A flow cell was assembled from a fabricated device, a top glass layer with inlet and outlet holes, and a spacer sheet.

Mobile imaging platform (MobIP)

We built a MobIP based on an Xperia smartphone (SO03-J, SONY, Japan). We 3D-printed an optomechanical platform structure on which the following components were mounted: a mechanical stage, high power light emitting diodes (LEDs) (365 nm at 3 W), a long-pass thin-film filter ($\lambda = 410 \text{ nm}$), and an aspheric lens ($f = 4 \text{ mm}$, N.A. = 0.64). The mechanical stage had vertical and lateral actuator systems for adjusting the focal position of a flow cell and multipoint imaging. The LED was placed at the side of the FRAD. Incident light was introduced from the side wall of the coverslip of the FRAD,

through a black rubber slit clipping a fabricated coverslip to separate the parallel component of the incident light.^{34,46} The reflection indices of CYTOP and water are 1.34 and 1.33, respectively, so that the incident light propagated inside the coverslip *via* waveguide-based total internal reflection when the incident angle was over 42° . Illumination by an evanescent field at the water/glass interface largely reduced the background noise and enabled detection of the fluorescence signal from single enzymes or single virus particles with an ordinal long pass filter. An aspheric lens was placed between the long pass filter and the smartphone camera for focusing the image as previously reported.^{31,32} We tested several types of aspheric lenses and found that C340TMD-A (Thorlabs) was the most suitable for imaging in the present setup. In addition, a microactuator system was implemented for lateral positioning of the FRAD in relation to the camera. The actuator system and LED were controlled with a microcomputer unit connected to the smartphone *via* Bluetooth Low Energy (BLE). When needed, multiple point imaging was conducted using the lateral actuator system (Fig. S2†).

Digital enzyme assay of bovine alkaline phosphatase (bALP) on the MobIP

To test the performance of the MobIP, we conducted a digital enzyme assay of bALP, which has a high dephosphorylation activity (about 1400 reactions per s per molecule) and was thus suitable as a model enzyme for an initial proof-of-concept experiment. The bALP solution was mixed with a fluorogenic substrate [4-methylumbelliferone (4MUP)] and introduced into the FRAD, followed by oil sealing (Fig. 2a). After incubation for 20 min at room temperature, the FRAD was inserted in the MobIP. The concentration of bALP was diluted to 1 pM, which corresponds to 0.029 enzyme molecules per reactor, ensuring digital assay conditions where single enzyme molecules are stochastically entrapped in reactors.

Fig. 2b (left) shows a fluorescence image of the digital enzyme assay for bALP with the MobIP. Discrete fluorescent bright spots were observed. Because the spherical aberration caused image distortion in the marginal area, the circular field within 690 μm from the center (the dashed circle in Fig. 2b), where individual fluorescent reactors were identifiable, was selectively analyzed in each image. To confirm that the bright spots in the MobIP image corresponded to positive reactors that give a fluorescence signal over mean + 3 SD of the background signal, the same FRAD used in Fig. 2b was subjected to imaging with conventional fluorescence microscopy with a $\times 20$ objective lens. The whole image from the FRAD was scanned, and the corresponding image was identified (the right image in Fig. 2b). In this assessment, conventional fluorescence imaging found 139 positive reactors in total. Meanwhile, the MobIP detected 134 reactors, all consistent with the positive reactors from the microscopy image. Five reactors were not detected with the MobIP due to their dark signals and image distortion because of positioning near the marginal area.



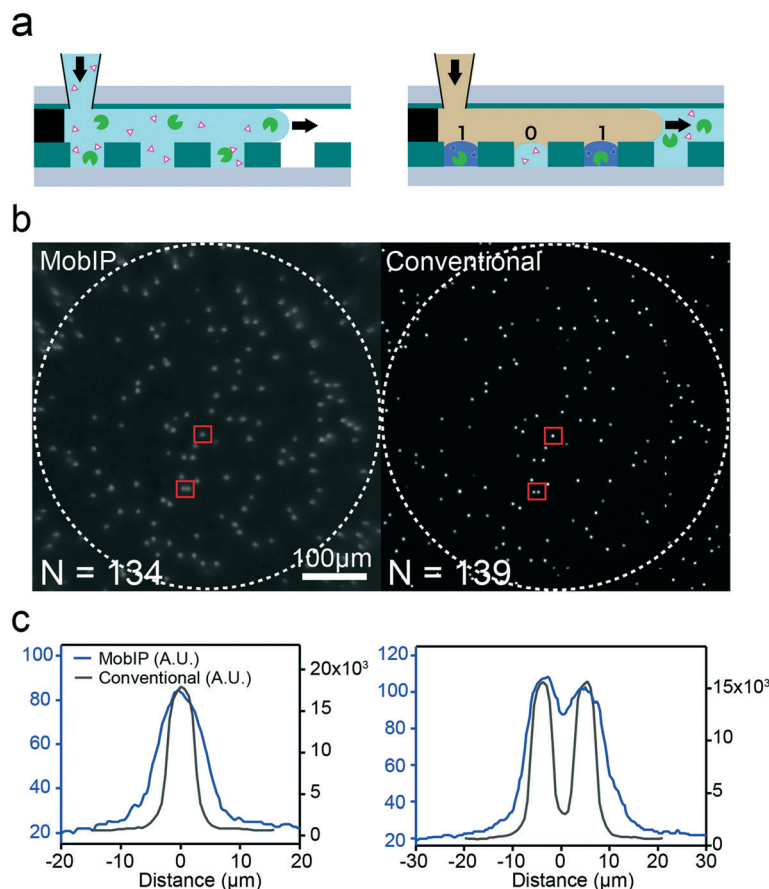


Fig. 2 Digital bioassay of bovine ALP (bALP) with a mobile imaging platform (MobIP). (a) Schematic image of a digital enzyme assay for bALP. (b) A typical fluorescence image obtained with the MobIP (left) and the corresponding image taken using conventional fluorescence microscopy (right). Pixel sizes of the MobIP and conventional fluorescence microscopy are 338 and 650 nm, respectively. White dashed circles indicate the area subjected to analysis. Positive reactors were identified as fluorescent spots giving a signal over mean + 3 SD of the background signal: $N = 134$ for the MobIP and $N = 139$ for the conventional microscope. (c) Intensity profiles of single (left) or double (right) positive reactors from the red squares in Fig. 2b.

The number of positive reactors detected in the above analysis was consistent with the expected values from the concentration of bALP. The MobIP found 134 positive signals among 5286 reactors in the analysis area, giving 0.025 as the average positive count per reactor (λ). This is consistent with the λ value (0.029) expected from the bALP concentration, and supports the detection and quantification performance of the MobIP.

The spatial resolution of images obtained with the MobIP was characterized from intensity profiles. Images of single or double positive reactors were arbitrarily selected from the image obtained with the MobIP (as marked with red squares in Fig. 2b), and their intensity profiles are shown in Fig. 2c. The corresponding images from conventional microscopy were also analyzed. Fitting these profiles with a Gaussian function determined that the full width at half maximum (FWHM) in the MobIP images was $8.72 \pm 0.74 \mu\text{m}$, while that of the microscopy image was $4.60 \pm 0.02 \mu\text{m}$. The spatial resolution for the MobIP was lower than that of the conventional microscopy. One possible reason for that is the spherical and other aberrations from the aspherical lens. Another possible

reason is that the images were processed for contrast enhancing or smoothing using built-in image processing software on the smartphone. Even with this relatively lower spatial resolution, the MobIP resolved fluorescence images of the two adjacent positive reactors separated by $9 \mu\text{m}$ (Fig. 2c). It should be noted that the spatial resolution is dependent on the position in the image field: it worsened near the marginal area. For further improvement, optimization of optical systems as well as development of software to completely control image acquisition, as well as processing, is required.

Digital influenza virus counting (DIViC) with the MobIP

We tested digital influenza virus counting (DIViC), based on previous work⁴⁴ (Fig. 3a) in which influenza virus particles were detected from the catalytic activity of neuraminidase (NA)—a surface glycoprotein of the influenza virus. Because DIViC does not require antibodies for detection but depends on the catalytic activity of the influenza virus by itself, DIViC requires only simple mixing of specimen solution with a fluorogenic substrate and introduction into the FRAD. Thus, DIViC is swifter and simpler than conventional antibody-



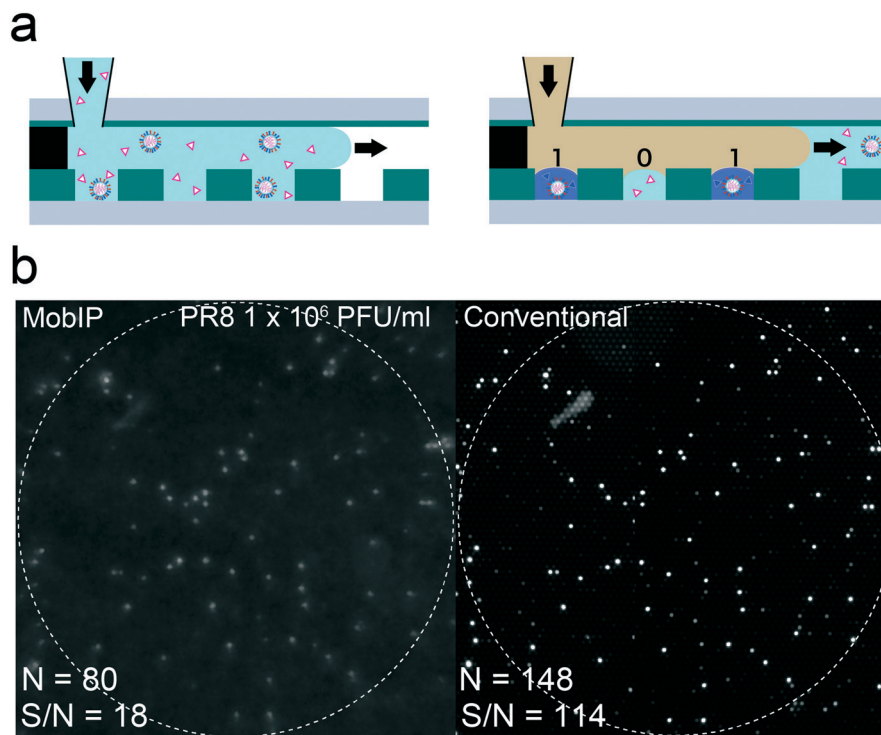


Fig. 3 Digital influenza virus counting (DIViC) imaged with the mobile imaging platform (MobIP). (a) Schematic image of DIViC. (b) Fluorescence image obtained with the MobIP (left) and a corresponding image taken with a conventional microscope (right). White dashed circles indicate the area subjected to analysis. Positive reactors were identified as fluorescent spots giving a signal over mean + 3 SD of the background signal: $N = 80$ for the MobIP and $N = 148$ for the conventional microscope. The sample was influenza type A virus (A/PR/8/1934(H1N1)) at 10^6 PFU ml^{-1} .

based assays. 2'-(4-Methylumbelliferyl)- α -D-*N*-acetylneuraminic acid (MUNANA) was used as the fluorogenic substrate, which produces a fluorescent dye [4-methylumbelliferone (4MU)] upon catalysis. We diluted the A/PR/8/1934(H1N1) samples to 1.0×10^6 PFU ml^{-1} , and MUNANA was added to give a final concentration of 1 mM.

We previously determined the catalytic activity of NA to be around 240 turnovers per s per virus particle,⁴⁴ six times lower than the turnover rate of bALP (1400 turnovers per s per molecule). Considering that bALP takes 20 min to give a sufficiently high signal on the MobIP, we predicted that the MobIP would take 2 h to detect the signal for DIViC. When incubated at room temperature for 20 min, obvious signals were not detected. In order to accelerate the reaction, the FRAD was incubated at 47 °C before imaging. This was based on the expectation of 4 times (or more) acceleration, given that the factor of activity for heating increments of +10 °C (Q_{10} factor) is generally 2 for enzymes.

The left panel of Fig. 3b shows the fluorescence image obtained with the MobIP after 10 min incubation (left). For comparison, the whole image of the same FRAD was scanned with a conventional microscope (right) after imaging with the MobIP. The corresponding image was identified in the same way as the digital enzyme assay for bALP (the right panel of Fig. 3b). All the fluorescence spots found with the MobIP were observed in the image with a conventional microscope. However, the total number of positive reactors found in the

MobIP was only 60% of those found in the image from the conventional microscope. This is notably lower than the digital enzyme assay of bALP with the MobIP. The close comparison of the images taken with the MobIP and conventional microscope revealed that fluorescent spots with lower signals were blurred with a higher background noise in the MobIP image. This was principally due to the relatively lower signal-to-noise ratio (S/N) in the MobIP images (18) than that for the conventional microscope (114). Fig. S3† shows the same images in Fig. 3b that is normalized with a median signal value among all pixel values of individual images, with an attempt to show images with a different contrast for comparison between images obtained with the MobIP and conventional microscope.

Different from bALP, the NA activity of the influenza virus shows high particle-to-particle variance: the variance coefficient of the NA activity among virus particles is reported to be 37%, while that for bALP is only 9%.⁴⁴ Therefore, the fraction of low activity virus particles was undetected in imaging with the MobIP.

Next, we tested DIViC on the MobIP at several virus titers, ranging from 1.0×10^6 to 3.0×10^4 PFU ml^{-1} (Fig. 4a–c). The total numbers of reactors subjected to analysis were approximately 30 000, 45 000, and 100 000 at 1.0×10^6 , 1.0×10^5 , and 3.0×10^4 PFU ml^{-1} , respectively. The numbers of detected virus particles were plotted against virus titer as a percentage against the total number of analyzed reactors (Fig. 4d). The



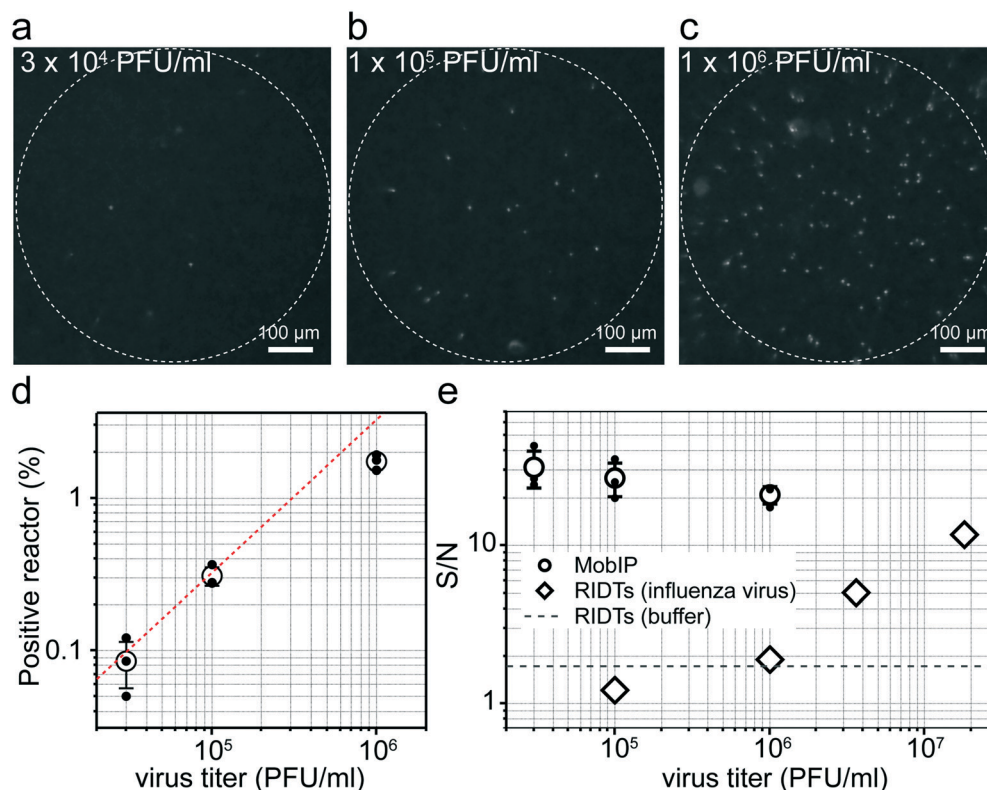


Fig. 4 Image of digital influenza virus counting (DIViC) with the mobile imaging platform (MobIP) at different virus titers. (a–c) Representative fluorescence images of DIViC obtained with the MobIP at different virus titers [A/PR/8/1934(H1N1)] of 3.0×10^4 , 1.0×10^5 , and 1.0×10^6 PFU ml^{−1}. (d) Positive reactor against virus titer. Linear fitting gave a count-to-PFU ratio of 375. (e) The ratio of fluorescence intensity of positive reactors to background noise against virus titer. For comparison, the S/N ratio of a commercial rapid influenza diagnostic test was determined (Fig. S4†). The open circle and diamond show the S/N at various virus titers for the MobIP and RIDTs, respectively. The dotted lines show the S/N of reaction buffer not containing influenza virus for RIDTs as reference.

plot showed fine linearity against the virus titer, giving a count-to-PFU ratio (CTPR) of 375. We also conducted DIViC with a conventional microscope for comparison, which gave a CTPR of 683. The detection efficiency of DIViC on the MobIP, compared with that of microscope-based DIViC, was 55%, essentially consistent with the above-mentioned value (60%). The CTPR value determined with the microscope was evidently higher than that previously reported for DIViC (189).⁴⁴ However, this value is within the range of values reported in the literature (8.0 to 2000) [for details see Table S1† in ref. 44]. The instability of the CTPR value can be attributed to the sensitivity of the CTPR to preparation and storage procedures.

The ratio of the fluorescence signal from positive reactors to background noise (signal-to-noise ratio: S/N) was essentially constant with the analyte concentration of the virus titer, as observed in other digital bioassays. Meanwhile, slight decay was observed at higher titers (Fig. 4e). This is due to the background noise increase that is attributable to stray light from fluorescence emissions from positive reactors that increase at higher virus titers.

Then, we compared the sensitivity of DIViC on the MobIP with rapid influenza diagnostic tests (RIDTs) (Fig. S4†), with the idea of using DIViC on the MobIP for diagnostic tests.

While RIDTs showed clear signals at 3.6×10^6 PFU ml^{−1}, the signal color band was not observed at 1.0×10^6 PFU ml^{−1} or lower, where S/N was around the S/N of the reaction buffer (Fig. 4e). Here, the signal intensity and background noise were analyzed from photomicrographs of RIDT devices (see Fig. S4†). The lowest virus titer, which gives distinct fluorescent spots in DIViC on the MobIP, was 3.0×10^4 PFU ml^{−1}. This comparison reveals that DIViC with the MobIP was at least 100 times more sensitive than with the RIDTs. It should be noted that at or less than 1.0×10^4 PFU ml^{−1}, fluorescent spots were not found at all. Such disappearance of fluorescent spots was not observed in DIViC on a conventional microscope. Although the reason is unclear, it should be attributed to the image processing protocol of the built-in software in the smartphone.

Finally, we investigated the feasibility of DIViC on the MobIP for analysis of clinical samples. To explore the possibility of clinical tests with less burden for patients, we tested two oral gargle samples from patients who were identified as influenza-positive by using RIDTs of their nasal swab. The types of influenza virus in the gargle were identified by RT-PCR as type A (H3N2) and type B, as reported.⁴⁴ The fluorescence images of DIViC with the MobIP showed positive reactors (Fig. 5a and b). When compared with the assay with the



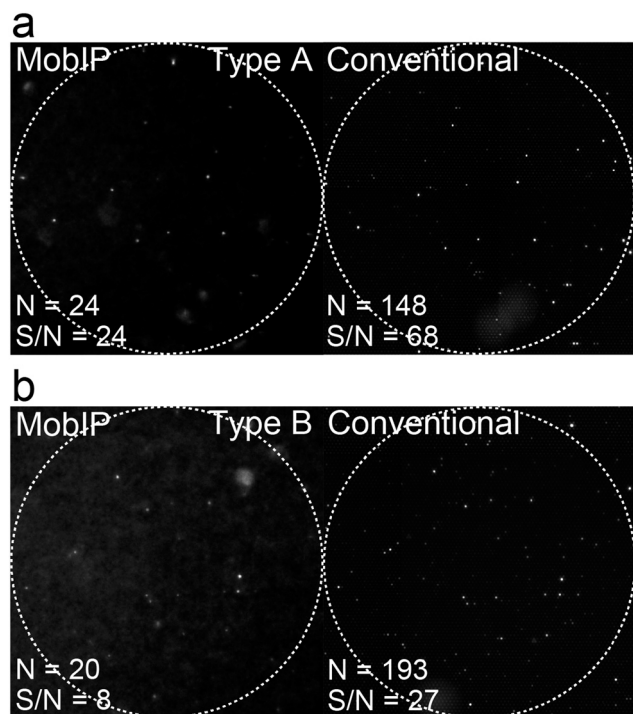


Fig. 5 Image of digital influenza virus counting (DIViC) with the mobile imaging platform (MobIP) by the use of clinical samples. Clinical sample test by DIViC with the MobIP. Fluorescence image obtained with the MobIP (left) and a corresponding image taken with a conventional microscope (right) for gargle containing influenza type A (a) and type B (b) identified by RT-PCR. White dashed circles indicate the area subjected to analysis. *N* means the number of positive reactors. S/N means the signal to noise ratio.

pure sample, the fluorescence signal was lower, suggesting that the neuraminidase activity of influenza virus in the clinical samples is lower than that of the laboratory strain. In addition, the background signal was also relatively higher, probably due to impurities or contaminants derived from the patient's mouth fluid. Same as the assays with pure solution, all positive reactors found with the MobIP were confirmed with a conventional microscope. The detection efficiency of positive reactors of clinical samples A and B with the MobIP compared with conventional microscopy was 16% and 10%, respectively. The lower detection efficiency could be attributed to the lower activity of the clinical samples and the higher background signal. Thus, it was demonstrated that the MobIP enables DIViC of clinical samples although the lower S/N of fluorescence images has to be improved for the development of diagnostic assays.

4. Conclusions

We developed a simple smartphone-based mobile imaging platform (MobIP) for the detection of digital bioassays. To achieve low noise fluorescence imaging with a simple optical setup, we employed evanescent field illumination by introducing incident light from the side of a femtoliter reactor array device (FRAD). Background noise was largely reduced,

allowing detection of fluorescence from femtoliter reactors encapsulating a single molecule of bovine alkaline phosphatase (bALP) or a single particle of influenza virus. Although the images obtained with the MobIP were distorted due to spherical aberration and the spatial resolution of the MobIP is lower than that of conventional microscopy, the MobIP enabled a quantitative digital bioassay of bALP and digital influenza virus counting (DIViC). The detection efficiency of DIViC with the MobIP was 60%, in relation to DIViC using conventional fluorescence microscopy. This difference is due to the relatively lower S/N of imaging with the MobIP than that of conventional microscopy and the intrinsically higher particle-to-particle variance of neuraminidase activity. Even with this detection sensitivity, DIViC with the MobIP showed 100 times greater sensitivity than a commercial rapid influenza diagnostic test.

These results suggest that DIViC with a MobIP is suitable for a new generation of point-of-care testing, enabling extremely sensitive detection of the influenza virus, and therefore diagnostic tests at the early phase of infection. The demonstration of the assay with clinical samples supports this result.

However, there are several technical challenges for further development of diagnostic tests. Firstly, the optical system has to be optimized to improve the spatial aberration and the spatial resolution. This would enable image analysis of more densely arrayed reactors with smaller volume, leading to a wider dynamic range as well as a swifter assay. The higher background signal found in the assays with clinical samples also remains to be addressed for a reliable diagnostic test with clinical samples.

Author contributions

H. N. conceived and coordinated this project; Y. M. and H. U. designed the optics and experimental platform; Y. M. developed the custom smartphone app and performed the digital assay and data analysis; K. V. T. gave technical support and conceptual advice; and H. N. and Y. M. wrote the manuscript.

Conflicts of interest

The authors declare no conflicts of interest.

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References

- 1 D. Witters, B. Sun, S. Begolo, J. Rodriguez-Manzano, W. Robles and R. F. Ismagilov, *Lab Chip*, 2014, **14**, 3225–3232.
- 2 Y. Zhang and H. Noji, *Anal. Chem.*, 2017, **89**, 92–101.
- 3 L. Cohen and D. R. Walt, *Annu. Rev. Anal. Chem.*, 2017, **10**, 345–363.
- 4 Y. Rondelez, G. Tresset, T. Nakashima, Y. Kato-Yamada, H. Fujita, S. Takeuchi and H. Noji, *Nature*, 2005, **433**, 773–777.
- 5 Y. Rondelez, G. Tresset, K. V. Tabata, H. Arata, H. Fujita, S. Takeuchi and H. Noji, *Nat. Biotechnol.*, 2005, **23**, 361–365.
- 6 D. R. Walt, *Chem. Soc. Rev.*, 2010, **39**, 38–50.
- 7 S. Sakakihara, S. Araki, R. Iino and H. Noji, *Lab Chip*, 2010, **10**, 3355–3362.
- 8 R. B. Liebherr, M. Renner and H. H. Gorris, *J. Am. Chem. Soc.*, 2014, **136**, 5949–5955.
- 9 D. Wu, E. Katilius, E. Olivas, M. Dumont Milutinovic and D. R. Walt, *Anal. Chem.*, 2016, **88**, 8385–8389.
- 10 K. Zhang, D. K. Kang, M. M. Ali, L. Liu, L. Labanieh, M. Lu, H. Riazifar, T. N. Nguyen, J. A. Zell, M. A. Digman, E. Gratton, J. Li and W. Zhao, *Lab Chip*, 2015, **15**, 4217–4226.
- 11 Z. Guan, Y. Zou, M. Zhang, J. Lv, H. Shen, P. Yang, H. Zhang, Z. Zhu and C. J. Yang, *Biomicrofluidics*, 2014, **8**, 014110.
- 12 R. A. White, 3rd, P. C. Blainey, H. C. Fan and S. R. Quake, *BMC Genomics*, 2009, **10**, 116.
- 13 H. C. Fan and S. R. Quake, *Anal. Chem.*, 2007, **79**, 7576–7579.
- 14 B. J. Hindson, K. D. Ness, D. A. Masquelier, P. Belgrader, N. J. Heredia, A. J. Makarewicz, I. J. Bright, M. Y. Lucero, A. L. Hiddessen, T. C. Legler, T. K. Kitano, M. R. Hodel, J. F. Petersen, P. W. Wyatt, E. R. Steenblock, P. H. Shah, L. J. Bousse, C. B. Troup, J. C. Mellen, D. K. Wittmann, N. G. Erndt, T. H. Cauley, R. T. Koehler, A. P. So, S. Dube, K. A. Rose, L. Montesclaros, S. Wang, D. P. Stumbo, S. P. Hodges, S. Romine, F. P. Milanovich, H. E. White, J. F. Regan, G. A. Karlin-Neumann, C. M. Hindson, S. Saxonov and B. W. Colston, *Anal. Chem.*, 2011, **83**, 8604–8610.
- 15 S. H. Kim, S. Iwai, S. Araki, S. Sakakihara, R. Iino and H. Noji, *Lab Chip*, 2012, **12**, 4986–4991.
- 16 D. M. Rissin, C. W. Kan, T. G. Campbell, S. C. Howes, D. R. Fournier, L. Song, T. Piech, P. P. Patel, L. Chang, A. J. Rivnak, E. P. Ferrell, J. D. Randall, G. K. Provuncher, D. R. Walt and D. C. Duffy, *Nat. Biotechnol.*, 2010, **28**, 595–599.
- 17 I. E. Araci, M. Robles and S. R. Quake, *Lab Chip*, 2016, **16**, 1573–1578.
- 18 L. Mazutis, A. F. Araghi, O. J. Miller, J. C. Baret, L. Frenz, A. Janoshazi, V. Taly, B. J. Miller, J. B. Hutchison, D. Link, A. D. Griffiths and M. Ryckelynck, *Anal. Chem.*, 2009, **81**, 4813–4821.
- 19 A. Gansen, A. M. Herrick, I. K. Dimov, L. P. Lee and D. T. Chiu, *Lab Chip*, 2012, **12**, 2247–2254.
- 20 Z. Wu, C. H. Zhou, L. J. Pan, T. Zeng, L. Zhu, D. W. Pang and Z. L. Zhang, *Anal. Chem.*, 2016, **88**, 9166–9172.
- 21 R. Watanabe, N. Soga, D. Fujita, K. V. Tabata, L. Yamauchi, S. Hyeon Kim, D. Asanuma, M. Kamiya, Y. Urano, H. Suga and H. Noji, *Nat. Commun.*, 2014, **5**, 4519.
- 22 N. Soga, R. Watanabe and H. Noji, *Sci. Rep.*, 2015, **5**, 11025.
- 23 R. Watanabe, T. Sakuragi, H. Noji and S. Nagata, *Proc. Natl. Acad. Sci. U. S. A.*, 2018, **115**, 3066–3071.
- 24 K. Akama, K. Shirai and S. Suzuki, *Anal. Chem.*, 2016, **88**, 7123–7129.
- 25 S. M. Scherr, D. S. Freedman, K. N. Agans, A. Rosca, E. Carter, M. Kuroda, H. E. Fawcett, C. E. Mire, T. W. Geisbert, M. S. Unlu and J. H. Connor, *Lab Chip*, 2017, **17**, 917–925.
- 26 D. Sevenler, G. G. Daaboul, F. Ekiz Kanik, N. L. Unlu and M. S. Unlu, *ACS Nano*, 2018, **12**, 5880–5887.
- 27 H. Takehara, K. Miyazawa, T. Noda, K. Sasagawa, T. Tokuda, S. H. Kim, R. Iino, H. Noji and J. Ohta, *Jpn. J. Appl. Phys.*, 2014, **53**, 04EL02.
- 28 H. Takehara, M. Nagasaki, K. Sasagawa, H. Takehara, T. Noda, T. Tokuda and J. Ohta, *Jpn. J. Appl. Phys.*, 2016, **55**, 3S2.
- 29 M. Zarei, *TrAC, Trends Anal. Chem.*, 2017, **91**, 26–41.
- 30 A. Roda, E. Michelini, M. Zangheri, M. Di Fusco, D. Calabria and P. Simoni, *TrAC, Trends Anal. Chem.*, 2016, **79**, 317–325.
- 31 S. K. Vashist, P. B. Lippa, L. Y. Yeo, A. Ozcan and J. H. T. Luong, *Trends Biotechnol.*, 2015, **33**, 692–705.
- 32 Q. Wei, H. Qi, W. Luo, D. Tseng, S. J. Ki, Z. Wan, Z. Gorocs, L. A. Bentolila, T. T. Wu, R. Sun and A. Ozcan, *ACS Nano*, 2013, **7**, 9147–9155.
- 33 M. Kuhnemund, Q. Wei, E. Darai, Y. Wang, I. Hernandez-Neuta, Z. Yang, D. Tseng, A. Ahlford, L. Mathot, T. Sjoblom, A. Ozcan and M. Nilsson, *Nat. Commun.*, 2017, **8**, 13913.
- 34 H. Zhu, S. Mavandadi, A. F. Coskun, O. Yaglidere and A. Ozcan, *Anal. Chem.*, 2011, **83**, 6641–6647.
- 35 S. J. Yeo, K. Choi, B. T. Cuc, N. N. Hong, D. T. Bao, N. M. Ngoc, M. Q. Le, K. Hang Nle, N. C. Thach, S. K. Mallik, H. S. Kim, C. K. Chong, H. S. Choi, H. W. Sung, K. Yu and H. Park, *Theranostics*, 2016, **6**, 231–242.
- 36 L. Shen, J. A. Hagen and I. Papautsky, *Lab Chip*, 2012, **12**, 4240–4243.
- 37 S. Feng, D. Tseng, D. Di Carlo, O. B. Garner and A. Ozcan, *Sci. Rep.*, 2016, **6**, 39203.
- 38 J. E. Kong, Q. Wei, D. Tseng, J. Zhang, E. Pan, M. Lewinski, O. B. Garner, A. Ozcan and D. Di Carlo, *ACS Nano*, 2017, **11**, 2934–2943.
- 39 P. B. Lillehoj, M. C. Huang, N. Truong and C. M. Ho, *Lab Chip*, 2013, **13**, 2950–2955.
- 40 H. Guner, E. Ozgur, G. Kokturk, M. Celik, E. Esen, A. E. Topal, S. Ayas, Y. Uludag, C. Elbuken and A. Dana, *Sens. Actuators, B*, 2017, **239**, 571–577.
- 41 Y. Liu, Q. Liu, S. Chen, F. Cheng, H. Wang and W. Peng, *Sci. Rep.*, 2015, **5**, 12864.
- 42 T. Laksanasopin, T. W. Guo, S. Nayak, A. A. Sridhara, S. Xie, O. O. Olowookere, P. Cadinu, F. Meng, N. H. Chee, J. Kim,



- C. D. Chin, E. Munyazesa, P. Mugwaneza, A. J. Rai, V. Mugisha, A. R. Castro, D. Steinmiller, V. Linder, J. E. Justman, S. Nsanzimana and S. K. Sia, *Sci. Transl. Med.*, 2015, **7**, 273re271.
- 43 V. Yelleswarapu, J. R. Buser, M. Haber, J. Baron, E. Inapuri and D. Issadore, *Proc. Natl. Acad. Sci. U. S. A.*, 2019, **116**, 10.
- 44 K. V. Tabata, Y. Minagawa, Y. Kawaguchi, M. Ono, Y. Moriizumi, S. Yamayoshi, Y. Fujioka, Y. Ohba, Y. Kawaoka and H. Noji, *Sci. Rep.*, 2019, **9**, 1067.
- 45 A. J. Einfeld, G. Neumann and Y. Kawaoka, *Nat. Protoc.*, 2014, **9**, 2663–2681.
- 46 S. Ramachandran, D. A. Cohen, A. P. Quist and R. Lal, *Sci. Rep.*, 2013, **3**, 2133.

