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## A new non-scanning confocal microscopy module for functional voltage-sensitive dye and Ca<sup>2+</sup> imaging of neuronal circuit activity

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## A new non-scanning confocal microscopy module for functional voltage-sensitive dye and $\text{Ca}^{2+}$ imaging of neuronal circuit activity

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**Tominaga T, Tominaga Y.** A new non-scanning confocal microscopy module for functional voltage-sensitive dye and  $\text{Ca}^{2+}$  imaging of neuronal circuit activity. *J Neurophysiol* 110: 553–561, 2013. First published April 24, 2013; doi:10.1152/jn.00856.2012.—Recent advances in fluorescent confocal microscopy and voltage-sensitive and  $\text{Ca}^{2+}$  dyes have vastly improved our ability to image neuronal circuits. However, existing confocal systems are not fast enough or too noisy for many live-cell functional imaging studies. Here, we describe and demonstrate the function of a novel, non-scanning confocal microscopy module. The optics, which are designed to fit the standard camera port of the Olympus BX51WI epifluorescent microscope, achieve a high signal-to-noise ratio (SNR) at high temporal resolution, making this configuration ideal for functional imaging of neuronal activities such as the voltage-sensitive dye (VSD) imaging. The optics employ fixed 100- $\times$  100-pinhole arrays at the back focal plane (optical conjugation plane), above the tube lens of a usual upright microscope. The excitation light travels through these pinholes, and the fluorescence signal, emitted from subject, passes through corresponding pinholes before exciting the photodiodes of the imager: a 100- $\times$  100-pixel metal-oxide semiconductor (MOS)-type pixel imager with each pixel corresponding to a single 100- $\times$  100- $\mu\text{m}$  photodiode. This design eliminated the need for a scanning device; therefore, acquisition rate of the imager (maximum rate of 10 kHz) is the only factor limiting acquisition speed. We tested the application of the system for VSD and  $\text{Ca}^{2+}$  imaging of evoked neuronal responses on electrical stimuli in rat hippocampal slices. The results indicate that, at least for these applications, the new microscope maintains a high SNR at image acquisition rates of  $\leq 0.3$  ms per frame.

confocal microscopy; hippocampal slice; voltage-sensitive dye; optical imaging;  $\text{Ca}^{2+}$  imaging

CONFOCAL MICROSCOPY (Amos and White 2003; Pawley 2006) is of increasing importance in functional imaging of neuronal circuit (Bullen et al. 1997; Ikegaya et al. 2004; Ohki et al. 2005). However, existing confocal systems are not fast enough for many high-speed imaging applications because the temporal resolution is limited (Saggau 2006). The aim of our study was to develop confocal optics that can be used for functional imaging that requires fast image acquisition rates such as voltage-sensitive dye (VSD) imaging (Cohen et al. 1978; Grinvald and Hildesheim 2004; Peterka et al. 2011; Peterlin et al. 2000).

VSD imaging of rapid changes of membrane potential, such as action potentials, requires a frame rate of  $\geq 1$  kHz (Grinvald et al. 1982; Ichikawa et al. 1993; Tominaga et al. 2000, 2009; Vranesic et al. 1994). Because VSD signals are often as small

as  $10^{-4}$  of baseline fluorescence, large numbers of photons must be sampled during short time intervals to achieve a sufficient signal-to-noise ratio (SNR). This is because photon shot noise expressed as fraction of baseline fluorescence decreases with 1 per square root of the number of photons (Knöpfel et al. 2006). This is difficult to achieve with a conventional confocal microscope because their frame rate is limited and they are not sampling large numbers of photons.

Most confocal microscopes, including laser-scanning, Nipkow disk, and swept-field confocal microscopy, rely on some type of scanning mechanism to acquire an image, thus limiting image acquisition speed (Kino 1995). The optical system that we have developed eliminates the need for a scanning mechanism, using, instead, fixed arrays of pinholes that correspond to a photodiode array on the imager. Our complementary metal-oxide semiconductor (CMOS) imaging system (Tominaga et al. 2005; same as MiCAM Ultima BrainVision) is composed of only a 100- $\times$  100-pixel photodiode array but has a very high frame rate (10 kHz) at high SNR (70 dB). Each photodiode has a maximum surface area of 10,000  $\mu\text{m}^2$ . This imager enabled us to use a non-scanning pinhole array with each pinhole corresponding to a unique pixel of the imager.

The confocal microscopy module was designed to fit the standard camera port of an epifluorescent microscope (BX51WI; Olympus). We tested the microscope by imaging a hippocampal slice preparation of rat by bulk VSD and  $\text{Ca}^{2+}$  indicator staining. The results demonstrate that the module can achieve fast acquisition in a confocal configuration without introducing noise associated with traditional scanning confocal microscopes.

### MATERIALS AND METHODS

**Optical configuration.** A schematic diagram of the non-scanning confocal microscope is shown in Fig. 1. The confocal microscope consists of an infinity correction epifluorescent microscope (BX51WI) and our confocal microscopy module. Light from a xenon arc lamp (75-W Xe-lamp; U-LH75EAP0; Olympus) was introduced via a large core fiber optic cable [core diameter = 3 mm, numerical aperture (NA) = 0.48] and passed through an excitation filter of a mirror unit (compatible with Olympus microscope mirror unit, U-MF/XL). An electrical shutter (Copal 0 Shutter) was placed in the fiber optics condenser unit. The excitation light was reflected through a secondary objective lens onto a pinhole array plate (Fig. 1B) positioned in the intermediate image plane. The excitation light passing through each pinhole (P and P') was focused onto corresponding focus points (O and O') on the sample. Fluorescent light, emitted from each focal point on the sample, was then projected onto the same pinhole array by the objective lens ( $\times 20$  NA 1.0; Olympus) and tube lens ( $f = 180$  mm) of the epifluorescent microscope. So that the focal plane of the sample and the pinhole array plate were in optical phase conjugation, the fluorescence passing through each pinhole was collected by a

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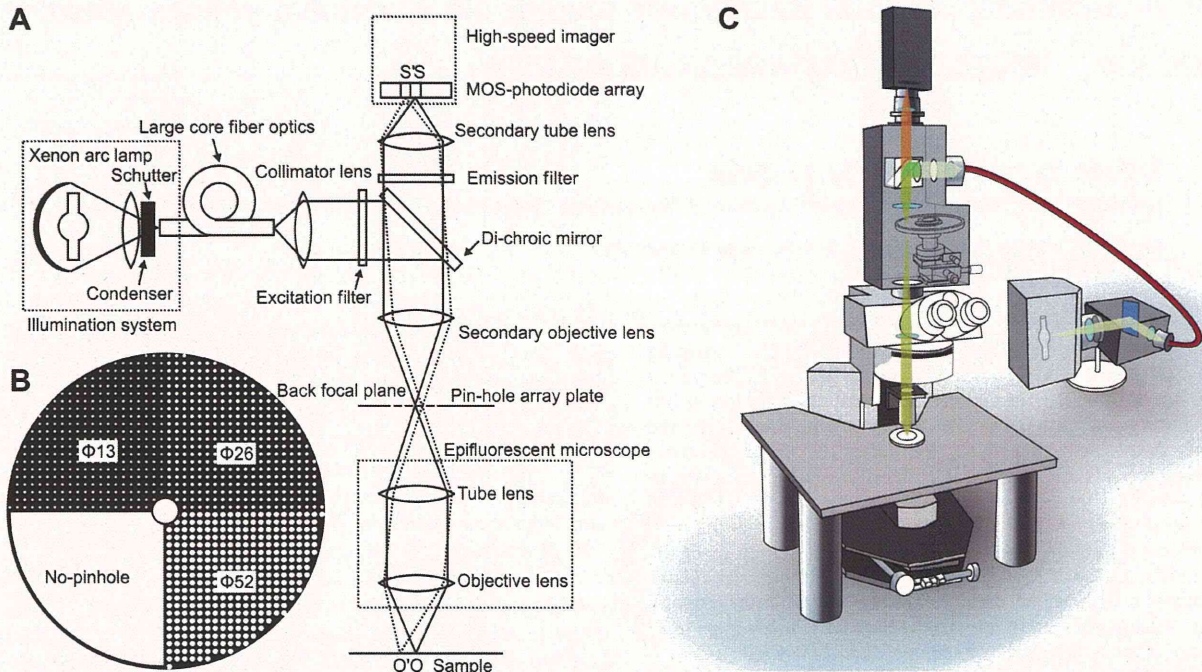


Fig. 1. Optical configuration of the nonscanning confocal microscope. *A*: schematic of the new confocal microscope. The confocal epifluorescent optics were designed to fit a conventional upright microscope with an objective lens and a tube lens (surrounded by dashed line). A pinhole array was positioned at the back focal plane, on which epifluorescent optics were placed. These optics consist of a secondary objective lens and secondary tube lens, between which a mirror unit (excitation filter, dichroic filter, and emission filter) was inserted. The fluorescence was then projected onto a complementary metal-oxide semiconductor (CMOS) high-speed imager. Lights from a sample in objective plane (O and O') correspond to lights sampled by photodiodes (S and S'). *B*: schematic drawing of the pinhole array plate consisting of 4 sectors with different pinhole patterns [pinhole diameters ( $\Phi$ ) of 13, 26, or 52  $\mu\text{m}$  and a no-pinhole sector]. *C*: illustration of the new confocal microscopy module attached to a conventional upright microscope (BX51WI; Olympus). A microscope was placed on a movable table combined with fixed table. The confocal module was placed on the upper optical part of the microscope. The excitation light was introduced by a large core fiber optic cable attached to a xenon light source with an electrical shutter.

secondary objective lens ( $f = 90 \text{ mm}$ ) and then projected onto each corresponding photodiode (S and S') of the CMOS-photodiode array after traveling through a secondary tube lens ( $f = 90 \text{ mm}$ ; XLFuor2X; Olympus) and a mirror unit containing a dichroic mirror and emission filter. The secondary objective lens and the secondary tube lens were at onefold magnification. The optics was adjusted for the  $\times 20$  objective lens ( $\times 20/0.95$  or  $\times 20/1.0$ ; Olympus), and the resulting field of view was approximately  $500 \times 500 \mu\text{m}$ . An overview of the microscope is shown in Fig. 1C.

**Pinhole array.** Our pinhole array (Fig. 1B) was made, from the type of glass plates used in the commercial disk confocal unit (DSU; Olympus), by a chromium deposition technique. The disk was divided into four sectors. Three of the sectors contained pinholes with 13-, 26-, or 50- $\mu\text{m}$  diameters. The fourth sector did not have any chromium deposition. The glass plate (disk) was fixed on a rotary shaft and mounted on an  $x$ - $y$  translation table (Sigma Koki, Tokyo, Japan), and the rotation angle was adjusted ( $\theta$  adjustment) to align each pinhole with its corresponding pixel of the imager.

The diameters of the pinholes were chosen to match the size of the first minimum of the Airy disk as follows:

$$\Phi = 1.22 \times \lambda / \text{NA}, \quad (1)$$

where  $\lambda$  is the wavelength of the excitation light (0.53  $\mu\text{m}$ : excitation light for di-4-ANEPPS) and NA equals the ratio of NA (1.0) and magnification ( $\times 20$ ) of the objective lens. According to Eq. 1, the calculated  $\Phi$  was 12.9  $\mu\text{m}$ .

To compensate for the loss of light transmission through the 13- $\mu\text{m}$  pinholes, we also used larger pinholes. This was accomplished by dividing the disk into four sectors: one without pinholes and three covered with a grid pattern of pinholes (13, 26, or 52  $\mu\text{m}$  in diameter) arranged in a square grid with an interval of 100- $\mu\text{m}$  distance. The

center part,  $1 \times 1 \text{ cm}$  that contains  $10^4$  pinholes in a sector, was used as a pinhole array.

**Slice preparation and staining with VSD.** All animal experiments were performed according to protocols approved by the Animal Care and Use Committee of Tokushima Bunri University. Hippocampal slices were prepared from 4- to 5-wk-old male rats decapitated under deep isoflurane anesthesia after perfusion with ice-cold artificial cerebrospinal fluid (aCSF; 124 mM NaCl, 2.5 mM KCl, 2 mM  $\text{CaCl}_2$ , 2 mM  $\text{MgSO}_4$ , 1.25 mM  $\text{NaH}_2\text{PO}_4$ , 26 mM  $\text{NaHCO}_3$ , and 10 mM glucose; pH 7.4) bubbled with 95/5%  $\text{O}_2/\text{CO}_2$  gas. The brains were quickly removed and cooled in aCSF. After cooling for 5 min, the hippocampus and the surrounding cortex was dissected and sliced into 400- $\mu\text{m}$  transverse sections using a vibratome (Leica VT1000). Following incubation in gassed aCSF for 3–5 min, each slice was transferred onto a fine-mesh membrane filter (Omnipore Membrane Filter, JHWP01300; Millipore) held in place by a thin Plexiglas ring (inner diameter, 11 mm; outer diameter, 15 mm; thickness, 1–2 mm). These slices were transferred to a moist chamber continuously supplied with a humidified  $\text{O}_2$ - $\text{CO}_2$  gas mixture. The temperature was held at 32°C for 1 h and then maintained at room temperature thereafter.

**Bulk staining and imaging with the VSD.** After the 1-h incubation, each slice was stained for 25 min with 100  $\mu\text{l}$  of the VSD solution, containing 0.2 mM di-4-ANEPPS (Molecular Probes) in 2.5% ethanol, 0.13% Cremophor EL (Sigma), 1.17% distilled water, 48.1% fetal bovine serum (Sigma), and 48.1% aCSF. After washing, sections were incubated at room temperature for  $\geq 1 \text{ h}$  before they were imaged.

**Single-cell staining and imaging with the VSD.** Individual pyramidal cells in the CA1 area of the sections were visualized using a conventional CMOS camera (SKDCE-2EX; Sigma Koki) with oblique illumination. Patch-clamp recordings, in whole cell mode, were made



using a patch-clamp amplifier with a capacitive headstage (Axoclamp 700B; Axon Instruments, Foster City, CA). Borosilicate glass pipettes (Sutter Instruments, Novato, CA) were pulled using a P-97 Flaming/Brown pipette puller (Sutter Instruments). The cesium-based pipette internal solution, used for whole cell voltage-clamp experiments, consisted of (in mM): 120 CsMeSO<sub>3</sub>, 10 HEPES, 4 MgCl<sub>2</sub>, 4 NaATP, 0.4 NaGTP, 10 Na-phosphocreatine, 5 QX-314, 10 EGTA; pH was adjusted to 7.2 (2–4 MΩ). Whole cell recordings were low-pass filtered at 3 kHz and digitized at 10 kHz (ITC-18; InstruTECH). Data were fed into a computer for online and offline analysis using laboratory-developed software on IGOR Pro (WaveMetrics). In voltage-clamp mode, a test membrane potential step (–10 mV) was always applied before electrical stimulation, and traces with series resistance (*R<sub>s</sub>*) <20 MΩ were accepted. After establishing optimal whole cell clamp conditions, the internal solution of the patch pipette was replaced with internal solution containing VSD (1 mg di-2-ANEPPS/1 ml internal solution) for 20–30 min. The internal solution was then replaced with internal patch pipette solution. For the patch pipette perfusion, we used a pipette holder with perfusion port and suction port. Thin-tipped polypropylene tubing was inserted into the perfusion port and connected to a syringe. The tip of the perfusion tube was positioned 400–600 μm from the tip of the patch pipette. The internal pipette solution was replaced by applying pressure through the syringe after making the whole cell condition.

For VSD imaging, we used a mirror unit consisting of a 530 ± 15-nm excitation filter, a 590-nm dichroic mirror, and 610 ± 20-nm emission filter.

**Staining and imaging with Ca<sup>2+</sup> indicator.** Fifty micrograms of Oregon Green 488 BAPTA-1, AM (special package; Molecular Probes) was dissolved into 10 μl of 0.5% Cremophor EL-DMSO mixture and added to 2-ml aCSF to yield final concentrations of

0.0025% (20 mM) AM-ester dye, 0.002% Cremophor EL, and 0.5% DMSO. Slices were bath-stained for 40 min at 37°C with an aliquot (100 μl per section) of the Ca<sup>2+</sup> indicator solution. Sections were washed and incubated at room temperature for ≥1 h before imaging.

For imaging of the Ca<sup>2+</sup> indicator, we used a mirror unit for FITC (FITC-3540B, BrightLine series; Semrock) consisting of a 482 ± 17-nm excitation filter, a 513-nm dichroic mirror, and a 536 ± 20-nm emission filter.

**RESULTS**

**Optical sectioning.** To test the optical sectioning of the new confocal module, we measured light intensity reflected on a total reflection mirror placed under the objective lens and plotted the light intensity at the center of the imager as a function of focus height along the z-axis (Fig. 2A). When in the no-pinhole configuration (Fig. 1B, No-pinhole), the microscope worked as a conventional bright-field microscope. The intensity of the reflected light linearly decreased around the focal plane as the focus was raised along the z-axis, as determined under transmitted light illumination. In contrast, the pinhole configurations (diameters: 13 μm on sector 1, 26 μm on 2, and 53 μm on 3) resulted in peak light intensity around the focus plane.

Next, we tested the microscope using a calibration kit of fluorescent bead preparation [Rainbow Fluorescent Particle Slide 6 Sizes (0.56, 0.96, 3.0, 5.5, 10.0, and 15.5 μm), FPS-M57-6; SpheroTech]. We used particle size 5.5 for the calibration. Figure 2B shows a composed (slit) image consisting of a line

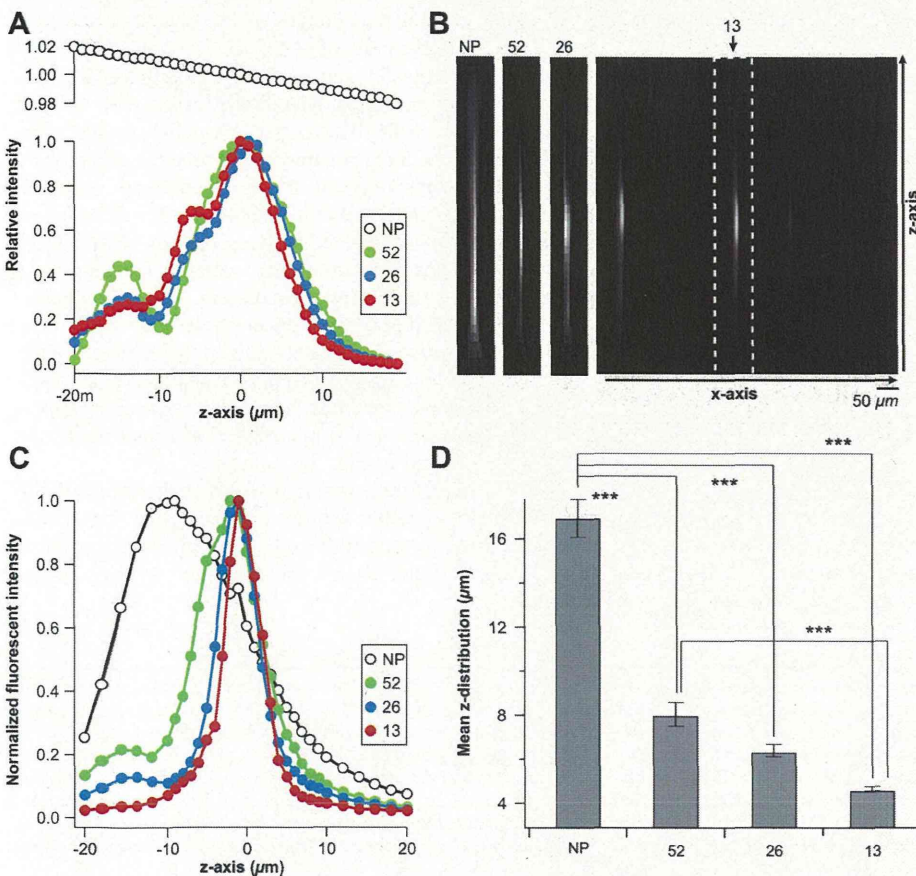


Fig. 2. Basic optical characteristics of the new confocal microscopy module. *A*: light intensity detected at the imager after reflection from a total reflection mirror placed under an objective lens in the nonpinhole configuration (NP; open circles) and with arrays of pinholes of 13-μm (13; red), 26-μm (26; blue), and 52-μm (52; green) diameter. On the z-axis, 0 indicates the focal point determined with transmitted light. *B*: fluorescent images of the standard fluorescent bead preparation, swept by a single horizontal line of the imager, collected using the no-pinhole, 52, 26, and 13 configurations. *C*: the profile of fluorescence intensity of beads shown in *B* for the no-pinhole and 52-, 26-, and 13-μm pinhole configurations. *D*: pooled data of the half-length of the profile shown in *C*. \*\*\**P* < 0.001, Tukey post hoc test.



of pixels collected from different focal planes (1- $\mu\text{m}$  step, 40 steps). Figure 2B, 13, shows a slit image taken when the 13- $\mu\text{m}$  pinhole array was used. Here, 2 or 3 lines are observed along the  $z$ -axis, each reflecting an image of a bead. The same image, indicated by dashed line, was taken with the 26- $\mu\text{m}$  (Fig. 2B, 26) and 52- $\mu\text{m}$  (Fig. 2B, 52) pinhole arrays. Finally, an image of the same bead, taken using no-pinhole configuration (*sector 0*), is shown in Fig. 2B, NP. The changes in light intensities along a line and within the same pixel are plotted in Fig. 2C. The mean  $z$ -distribution was measured as the half-width of the fluorescence intensity profiles along the  $z$ -axis, recorded for 20 different beads. These data are shown in Fig. 2D. The half-width was significantly decreased when pinholes were used, and the size of pinhole inversely affected the  $z$ -resolution of the beads. The results indicate that the configuration used in this study effectively improved depth resolution even when the largest pinholes were used.

**Images and noise consideration.** Fluorescent images of VSD-stained hippocampal slices were obtained with the confocal module using all four pinhole configurations (no-pinhole and diameters equal to 52, 26, and 13  $\mu\text{m}$ ). The intensity of fluorescence at each pixel decreased as the size of the pinhole decreased; that must be proportional to the fourth power of the pinhole radius. To acquire images within the dynamic range of the imager, the images were acquired at four different accumulation times (0.1, 1, 10, and 100 ms per frame). The consecutive images were obtained at different focal depths in 5- $\mu\text{m}$  steps.

The images obtained with the 13- $\mu\text{m}$  pinhole array had the finest resolution, and the images tended to blur as the diameter of the pinholes increased. The images obtained with the 13- $\mu\text{m}$  pinhole appeared different at every 5- $\mu\text{m}$  step along the  $z$ -axis, whereas images taken without pinholes looked similar across the 30- $\mu\text{m}$  change in the  $z$ -axis. The time courses of the optical signal obtained with these conditions are presented on the right-hand side of the images of Fig. 3A. Image intensities decreased and baseline-normalized noise increased as pinholes became finer (Fig. 3A).

To identify the source of the noise in the optical signals, the standard deviation of the noise [i.e., root-mean-square (RMS) noise] was measured in pixels from images obtained using different acquisition conditions [e.g., acquisition rates ranging from 0.1 to 200 ms and presence or absence of a neutral density (ND) filter]. To sample a range of fluorescence intensities, a vertical line of pixels through the center of the images was used for this calculation and was plotted on the same graph as shown in Fig. 3B. The RMS noise increased as the fluorescence increased for each pinhole configuration. A double-logarithmic plot of this relationship shows that normalized RMS noise by  $F_0$  (RMS noise/ $F_0$ ) decreased as a first order function of  $F_0$  below an  $F_0$  equivalent to 1% saturation of imager (approximately  $1 \times 10^6$  electrons) and decreased proportionally to the square root of  $F_0$

when  $F_0$  was  $>1\%$ . That is, when  $F_0$  was  $<1\%$  of the maxima of imager photodetector, the major cause of the noise was equal to the constant amount of noise ( $\sim 0.02\%$ , corresponding to 230 electrons) for the imaging device (Dr. Michinori Ichikawa, BrainVision, personal communication). In contrast, when  $F_0$  was  $>1\%$ , the major source of noise was the shot noise, which is proportional to square root of light intensity.

**VSD signals obtained from a hippocampal slice.** Figure 4, A and B, compares the VSD signals in the CA1 area of hippocampal slices on electrical stimulation of the Schaffer collateral pathway. The *top* shows the data acquired in the no-pinhole configuration, and the *bottom* shows the data acquired using the 52- $\mu\text{m}$  pinhole array. Both signals were acquired at a frame rate of 0.3 ms per frame. The intensity of the initial fluorescence with the two conditions was matched by applying a ND filter (6% transmission) to the no-pinhole condition. The optical signal, normalized by initial fluorescence, was almost identical in both the nonpinhole and 52- $\mu\text{m}$  pinhole condition. The optical signal of representative pixels, along the axis parallel to pyramidal cells, is shown in Fig. 4C for the non-pinhole (A) and 52- $\mu\text{m}$  pinhole (B) configurations. As indicated by the superimposed traces, the signals obtained from both configurations were nearly the same.

Figure 4D shows a confocal fluorescent image of a pyramidal cell, perfused with VSD (di-2-ANEPPS), from a whole cell clamp preparation collected at a frame rate of 10 ms per frame using the 52- $\mu\text{m}$  pinhole array. The sensitivity of the optical signal was about 10%/100-mV change (Fig. 4E), similar to the value previously reported for di-4-ANEPPS (Loew et al. 1992). Figure 4 shows representative images of the optical signals detected for membrane potentials of  $-20$ ,  $20$ ,  $-50$ , and  $50$  mV.

**$\text{Ca}^{2+}$  imaging.** Figure 5A shows the optical signal acquired from CA1 area of a hippocampal slice being loaded with Oregon Green 488 BAPTA-1, AM. The data presented here were acquired at a frame rate of 5 ms per frame using the 52- $\mu\text{m}$  pinhole array and show the response of the CA1 area on electrical stimulation of the Schaffer collateral pathway. The data show several cells in the pyramidal cell layers that exhibited large changes in fluorescence on stimulation, whereas the responses were less conspicuous in the stratum radiatum. It is also interesting to note that the initial fluorescence levels were high in s. alveus, and the responses remained obvious in this region (Fig. 5, A and B). Figure 5C shows the peak optical signals for each pixel. The large, steep response shown in the upper right corner represents the  $\text{Ca}^{2+}$  response in a single cell, corresponding to the optical signal represented in Fig. 5B, trace 4.

The 26- $\mu\text{m}$  pinhole array was also effective at detecting the response of these cells when acquired at 10 ms per frame, but the data were similar to those presented here. In the nonpinhole condition, it was difficult to see the specimen as the images were blurred.

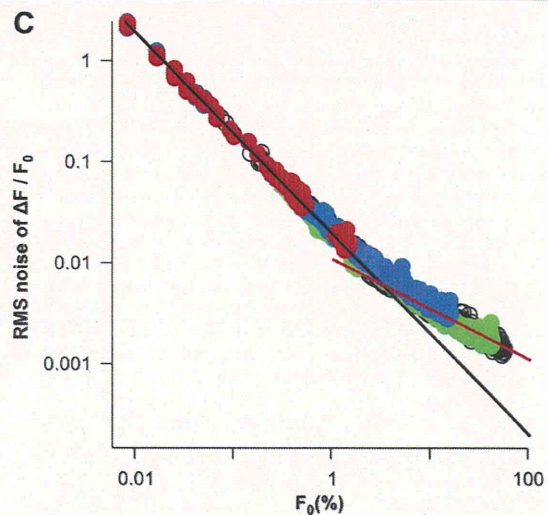
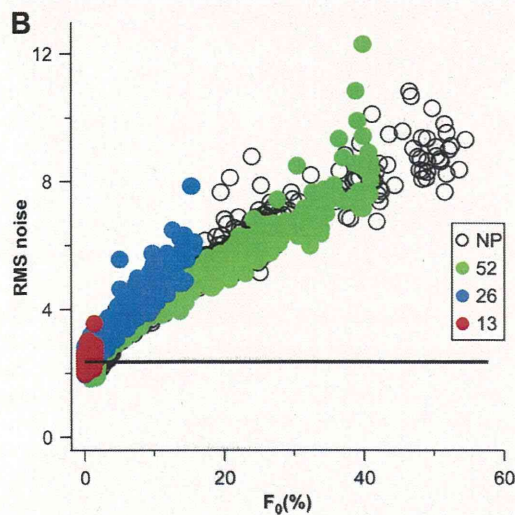
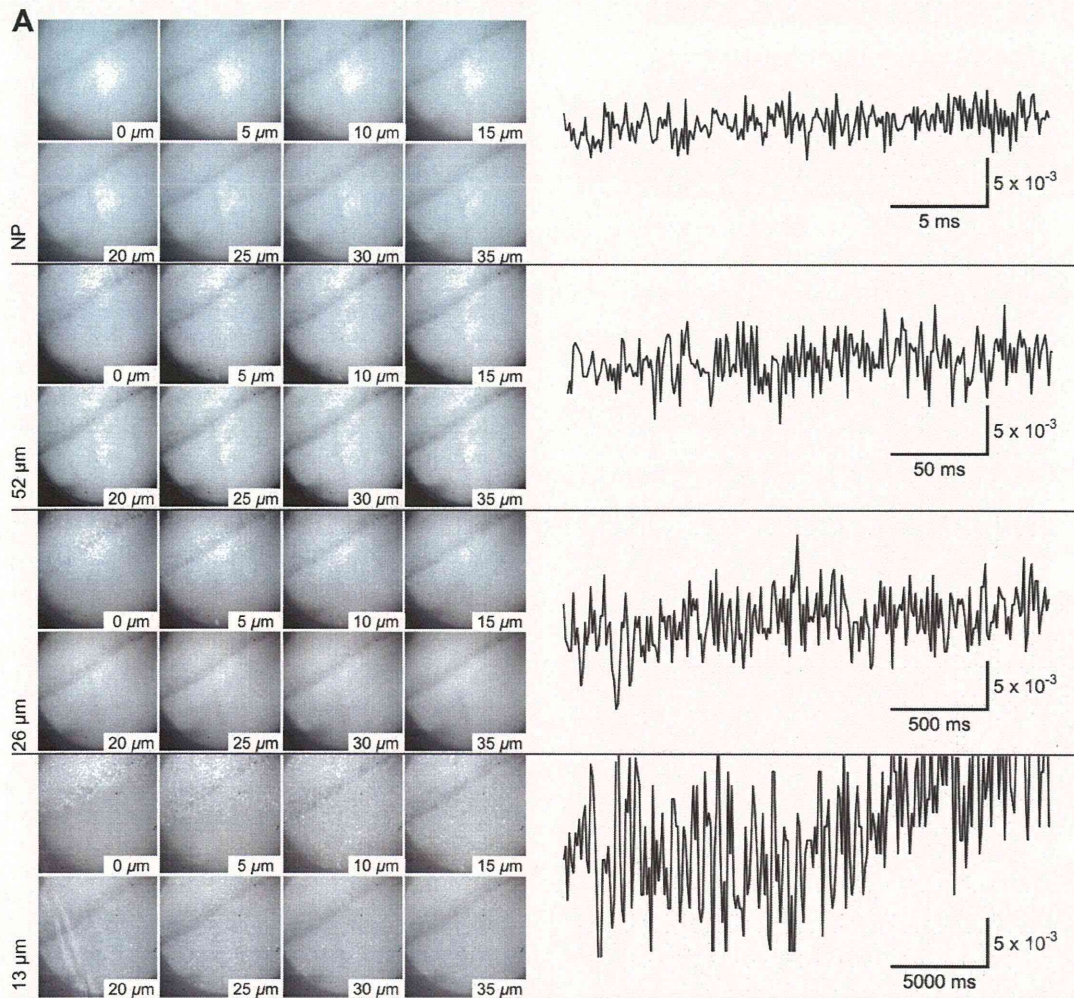
Fig. 3. Fluorescent images of rat hippocampal slices stained with voltage-sensitive dye (VSD) and acquired by the confocal microscopy module using different pinhole configurations. A: images of VSD-stained hippocampal slice collected using the different pinhole configurations at different positions in the  $z$ -axis. The *right* traces show the optical signal from a single pixel of the image. B: root-mean-square (RMS) noise in the time course of the fluorescence plotted against initial fluorescent intensity ( $F_0$ ). Fluorescent intensity was expressed relative to the saturation level of imager (approximately 1 M electron per pixel). RMS noise was calculated as a sum of square roots of measurements at time  $t$  ( $F_t$ ) deviation from the  $F_0$  (i.e., the standard deviation of the optical signal). The measurements plotted in the graph were obtained with different pinhole conditions (black, no-pinhole; blue, 13  $\mu\text{m}$ ; green, 26  $\mu\text{m}$ ; red, 52  $\mu\text{m}$ ) at acquisition rates ranging from 0.1 to 200 ms for different pixels in the view. C: normalized RMS noise ( $\Delta F/F_0$ ) plotted against  $F_0$  in double-logarithmic scale for the same measurements as B. The black line shows the constant RMS value of 0.02% obtained from the darkest RMS noise divided by  $F_0$  so that the slope of the line (a constant/ $F_0$ ) on the graph equals to  $-1.0$ . The red line shows the slope of photon shot noise proportional to square root of  $F_0$ .



DISCUSSION

*Advantages and limitation of the system.* Using a fixed-pinhole array, our confocal module achieves confocal signals (Amos and White 2003; Pawley 2006) at all of the 10,000 pixels on an imager simultaneously. Because the pinholes are

stationary, there is no scanning noise (Saggau 2006) as seen in traditional confocal microscopy such as confocal laser-scanning microscopes (CLSM) and Nipkow disk confocal microscopes (Kino 1995). In addition, because the image acquisition rate is not restricted by a scanning mechanism, our system can





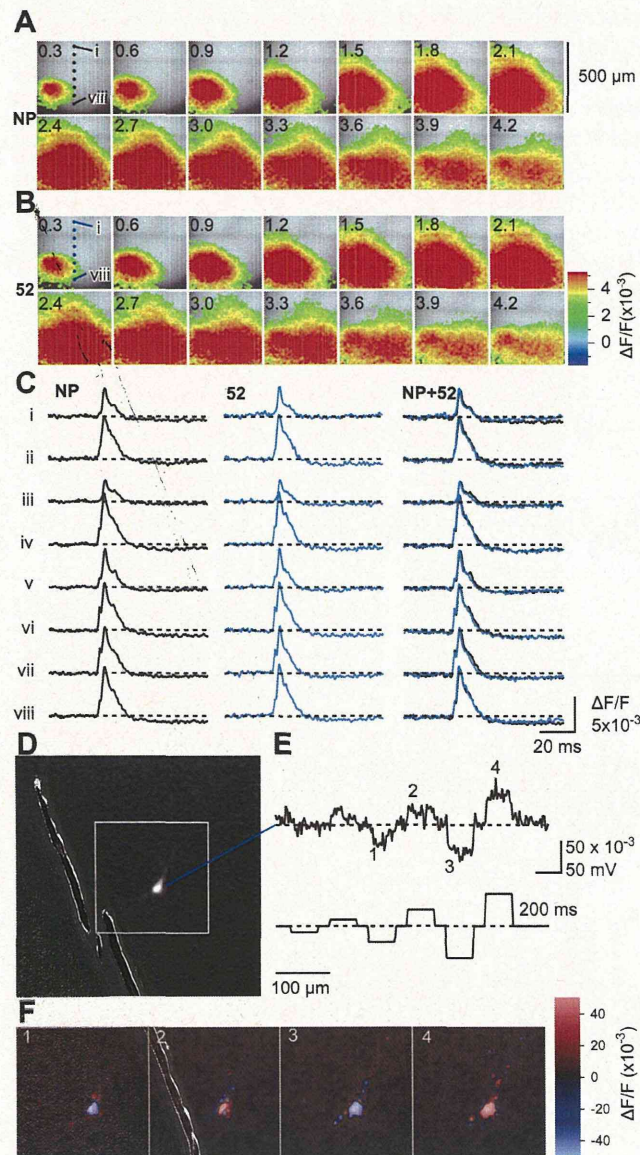


Fig. 4. Representative optical signals recorded from VSD-stained hippocampal slices on an electrical stimulation of the Schaffer collateral pathway using the conventional (no-pinhole) and 52- $\mu\text{m}$  pinhole configurations. Consecutive images are of fluorescence change ( $\Delta F/F$ ), in pseudocolor, recorded after electrical stimulation to Schaffer collateral pathway using the no-pinhole (A) and 52- $\mu\text{m}$  diameter (B) pinhole configurations at the frame rate of 0.3 ms per frame. The number of each image is the time after an electrical stimulation given to the Schaffer collateral (0 ms). The initial fluorescence was matched by using a neutral density filter (6%) when no pinhole was used. C: optical traces of the pixels indicated in A and B (i–viii) in no-pinhole condition and 52- $\mu\text{m}$  pinhole condition. Superimposed traces are shown in NP+52. D: fluorescent image of VSD-stained pyramidal cell in the CA1 area. The images were taken at the frame rate of 10 ms per frame when 52- $\mu\text{m}$  pinhole configuration was used. E: optical traces under voltage-clamp condition at the center of the cell are shown on the right-hand side of the images. F: images of the fluorescent change in pseudocolor code at the acquisition times indicated in the E traces. One corresponds to  $-20\text{-mV}$ , 2 to  $20\text{-mV}$ , 3 to  $-50\text{-mV}$ , and 4 to  $50\text{-mV}$  membrane potential.

achieve higher acquisition rates, which are limited only by the imaging system. In its implementation, the system has a spatial resolution of only  $100 \times 100$  pixels. The use of more dense pinholes and a camera with a larger number of pixels can

decrease pixilation. As in normal wide-field epifluorescence microscopes, it is not very straight-forward to achieve very high excitation light densities. Moreover, only a small fraction of the light that is projected from the light source onto the pinhole array passes the pinholes (1.3, 4.6, and 18.4% of the illuminated area for 13-, 26-, and 52- $\mu\text{m}$  pinholes, respectively). Use of a more intense light source will increase effective excitation intensities and, hence, SNR.

*Comparison with other high-speed imaging systems.* For the CLSM, the product of frame rate and number of pixels is limited by the speed of the scanning process such that frame rates  $>50$  Hz demand for a strict reduction of the number of measurement points. For off-the-shelf disk-scanning confocal system, the temporal resolution is often limited by the imager, and there are similar tradeoffs between the spatial resolution and frame rates. Confocal spot detection can avoid this trade-off and has been used to visualize  $\text{Ca}^{2+}$  microdomain dynamics (Escobar et al. 1994; DiFranco et al. 2012; DiGregorio et al. 1999). This method has the clear advantage of measuring the optical signal from a single point at high speed with good SNR (Bradley et al. 2009; Fink et al. 2012). Moreover, recent optoacoustic device-driven random-access fluorescence microscopy (Bullen and Saggau 1999; Otsu et al. 2008; Saggau et al. 1998), which enables the measurement of several points, has the advantage that it can measure several points with a performance similar to the single-point measurements. These measurements can then be converted into a multiphoton depth image (Duemani Reddy et al. 2008). The nonlinear measurements, using multiphoton microscopy, were extended to record second harmonic generation signals (Dombeck et al. 2004, 2005; Nuriya et al. 2006).

The light-sheet microscopy, with special focus on high-speed image acquisition such as the selective plane illumination microscopy (SPIM; Huisken 2012; Huisken and Stainier 2009) and/or the objective-coupled planar illumination (OCPI) microscope (Turaga and Holy 2012) can provide high-speed image acquisition while still allowing the thinner optical dissection and volumetric imaging. With these techniques, the image acquisition speed is determined by the imager acquisition rate rather than the scanning devices, similar to the present system. Moreover, because SPIM and OCPI focus excitation light using a thin sheet, often through a cylindrical lens, they offer advantages in terms of excitation light intensity compared with our system.

Light-sheet microscopy has been used mostly for the imaging of clear objects, such as embryos, because other specimens produce shadows that interfere with visualization. Recent introduction of multidirectional SPIM (mSPIM; Huisken and Stainier 2007) is overcoming this problem. In addition, the spatial resolution has now been improved by using nonlinear excitation with stimulated emission depletion (STED; Friedrich et al. 2011).

Compared with these examples of high-speed imaging microscopy, the current system has a clear disadvantage in terms of excitation light intensity. However, the simple configuration of our module offers an inexpensive high-speed imaging option for users that require a wider field of view such as that required for functional neuronal circuit imaging. In this regard, the module shares many of the advantages of confocal spot detection and fast frame acquisition of SPIM. Adding to this advan-



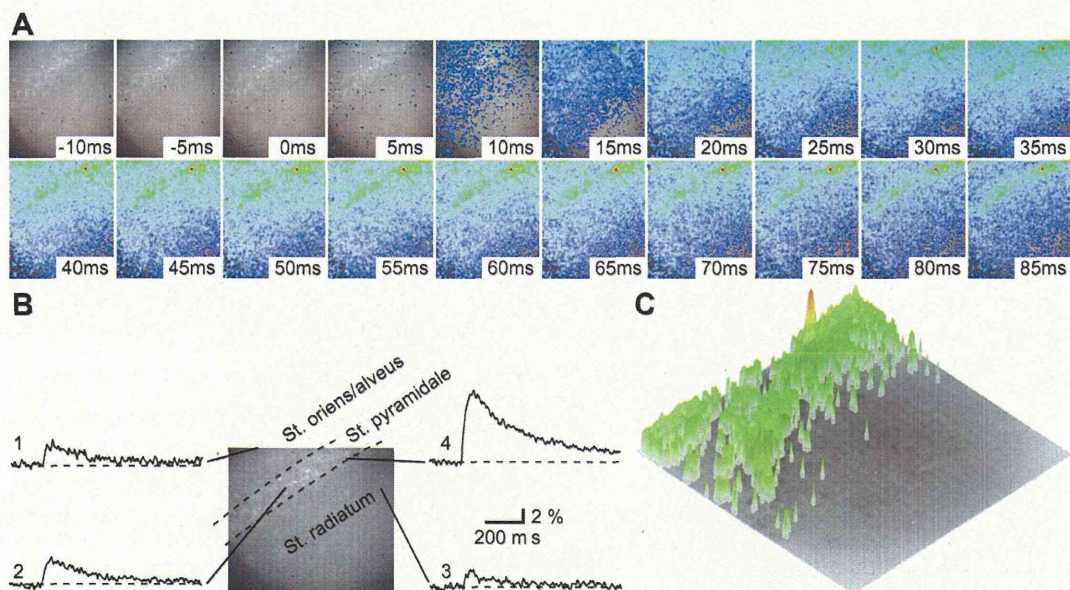


Fig. 5. Fluorescent  $\text{Ca}^{2+}$  imaging of indicator-stained hippocampal slices. *A*: consecutive images showing the spread of  $\text{Ca}^{2+}$  in response to an electrical stimulation (0 ms) of the Schaffer collateral. *B*: the  $\text{Ca}^{2+}$  signal at 4 representative pixels in the image. Shown are 1 for stratum oriens/alveus, 2 for s. pyramidale, 3 for s. radiatum, and 4 for a cell showing huge  $\text{Ca}^{2+}$  change. *C*: the 3-dimensional representation of the peak signal measured in each pixel. A highest spot is seen in a cell body of a neuron corresponding to 4 in *B*. The images were taken using the 26- $\mu\text{m}$  pinhole configuration at a frame rate of 5 ms per frame.

tage, oblique illumination with a separate optical axis could be useful, especially for low-magnification wide-field optical recording.

**Light sources.** For single-point measurements, CLSM can use lasers as light source. Because CLSM offers flexibility in choice of power and range of excitation wavelengths, these microscopes can achieve high SNR. However, stability of the laser and speckle noise caused by its coherent nature can also cause noise. In contrast, pinhole-array confocal microscopy requires a wide field of illumination. Here, lasers are less suitable due to their narrow beam diameter. Instead, excitation light is typically supplied by a xenon arc lamp, halogen light, or high-power LED with temperature compensation (Nishimura et al. 2006) because of their wider area of illumination. These are stable and inexpensive, but the light intensity is relatively limited, rendering them less suitable for point-scanning applications. Fortunately, rapid progress in high-power LED technology could overcome this limitation.

**Usefulness in VSD and  $\text{Ca}^{2+}$  imaging.** Among the various imaging techniques, techniques based on a fast VSD (Cohen et al. 1974; Salzberg et al. 1973) offer high spatial and temporal resolution. VSD rapidly converts changes in the membrane potential, such as action potentials, into optical signals using changes in either the absorption of transmitted light or emission of fluorescence (Cohen et al. 1978; Grinvald et al. 1988; Grinvald and Hildesheim 2004; Peterka et al. 2011). Recent progress in imaging devices together with the speed of the VSD response have allowed us to follow the submillisecond time scale of neuronal activities by VSD imaging. However, because the fractional optical signal is small, the difficulty in obtaining a good SNR is the most critical. Efforts have been aimed at improving the SNR by increasing the photon efficacy of wide-field excitation optics to increase the fluorescence intensity and achieve rapid image acquisition (Ratzlaff and Grinvald 1991; Tominaga et al. 2000).

The magnitude of the signal we obtained from hippocampal slices bulk-stained with VSD (di-4-ANEPPS) was approximately  $4 \times 10^{-3}$ . This is far smaller than the VSD signal expected (a relative change of few percent per 100 mV) from a single plasma membrane because of the absence of nonresponsive fluorescence background (Loew et al. 1992). In addition, VSD signals obtained from the CA1 area showed maximum response on Schaffer collateral stimulation in the layer of s. radiatum. However, as we have shown earlier, the signal was almost uniform if the membrane potential was elevated by high-potassium stimulation when the action potential was inhibited by TTX (Tominaga et al. 2009). These problems are related to the fact that the baseline VSD signal from bulk-stained samples is often high relative to the change. In other words, there are many membrane components that do not respond to stimulation but are stained with VSD. The use of our confocal microscope module did not resolve this issue, as shown in Fig. 4. However, recent progress in the development of a voltage-sensitive fluorescent protein (Akemann et al. 2010, 2012; Baker et al. 2008; Knöpfel 2008, 2012; Kralj et al. 2011, 2012) may offer a solution, as the combination of this technique with our confocal module should produce clearer signals.

As is shown in Figs. 4 and 5, our new confocal microscope is capable of VSD and  $\text{Ca}^{2+}$  imaging at very fast acquisition rates while maintaining a high SNR and should have applications for use with new imaging techniques where the high frame rate and low noise are needed.

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## DISCLOSURES

There are no conflicts of interest, financial or otherwise, to declare by the author(s).

## AUTHOR CONTRIBUTIONS

T.T. and Y.T. conception and design of research; T.T. designed the optics; Y.T. developed the software; T.T. and Y.T. performed experiments; T.T. and Y.T. analyzed data; T.T. and Y.T. interpreted results of experiments; T.T. and Y.T. prepared figures; T.T. and Y.T. drafted manuscript; T.T. and Y.T. edited and revised manuscript; T.T. and Y.T. approved final version of manuscript.

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