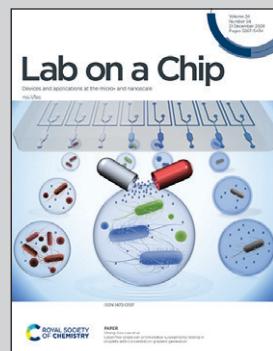


Showcasing research from Professor Takeharu Nagai's laboratory at the Department of Biomolecular Science and Engineering, SANKEN, The University of Osaka, Osaka, Japan. Artwork by Asako Chio.

Compact lens-free imager using a thin-film transistor for long-term quantitative monitoring of stem cell culture and cardiomyocyte production

This study addresses the need for practical devices to ensure quality control in long-term cell cultures and production processes. By incorporating a large thin-film transistor image sensor and rear side dark-field illumination, the authors developed a compact lens-free imager capable of monitoring six independent culture wells simultaneously in an incubator named INSPECTOR. INSPECTOR enables monitoring the confluence of pluripotent stem cell cultures and identifying the timing of cell state transitions during mesoderm induction, as well as tracing the beating frequency and conduction patterns of cardiomyocytes throughout long-term differentiation and maturation processes. This device facilitates quality control of cell products and has the potential to contribute to advancements in regenerative medicine and drug discovery.

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Compact lens-free imager using a thin-film transistor for long-term quantitative monitoring of stem cell culture and cardiomyocyte production†

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With advancements in human induced pluripotent stem cell (hiPSC) technology, there is an increasing demand for quality control techniques to manage the long-term process of target cell production effectively. While monitoring systems designed for use within incubators are promising for assessing culture quality, existing systems still face challenges in terms of compactness, throughput, and available metrics. To address these limitations, we have developed a compact and high-throughput lens-free imaging device named INSPECTOR. The device is as small as a standard culture plate, which allows for the installation of multiple units within an incubator. INSPECTOR utilises a large thin-film transistor image sensor, enabling simultaneous observation of six independent culture environments, each approximately 1 cm². With this device, we successfully monitored the confluence of hiPSC cultures and identified the onset timing of epithelial-to-mesenchymal transition during mesodermal induction. Additionally, we quantified the beating frequency and conduction of hiPSC-derived cardiomyocytes by using high-speed imaging modes. This enabled us to identify the onset of spontaneous beating during differentiation and assess chronotropic responses in drug evaluations. Moreover, by tracking beating frequency over 10 days of cardiomyocyte maturation, we identified week-scale and daily-scale fluctuations, the latter of which correlated with cellular metabolic activity. The metrics derived from this device would enhance the reproducibility and quality of target cell production.

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Introduction

The recent advances in human induced pluripotent stem cell (hiPSC) technology have enabled the production of various cell types constituting our human body, accelerating progress in regenerative medicine and drug discovery.^{1–3} The process of inducing target cells from hiPSCs is a prolonged procedure, lasting from days to weeks and sometimes even months. Furthermore, the optimised culture conditions and induction efficiency are known to vary from lab to lab, from one cell line to another, and sometimes depending on subtle differences in sample handling.^{4,5} Therefore, optimisation of the induction conditions is often required for each experimental

site, and strict control of the entire process is essential for achieving high reproducibility. Consequently, lab automation has attracted attention as a technology for efficiently handling a large number of cell culture samples with high reproducibility.^{6–8} On the other hand, it is crucial not only to handle cell culture efficiently and reproducibly but also to evaluate their quality effectively. If the induction process is evaluated on the end products alone, the entire process, which may have taken several weeks, might be rendered useless. The ability to monitor and evaluate quality during the induction process enables early assessment of culture conditions, rapid detection of abnormalities, and rigorous reproducibility, including at intermediate points.^{9–12} Thus, systems capable of long-term monitoring and quality evaluation of cell culture are of significant value.

Imaging devices installed inside incubators serve as the primary technology for long-term cell culture monitoring.¹³ At the same time, several requirements must be met to effectively monitor hiPSCs culture and their differentiation induction. The first requirement is compactness since incubators offer limited space. In particular, miniaturised microscopes and lens-free imaging systems meet this demand. Miniaturised microscopes are designed to maintain as high resolution as conventional microscopes in a smaller

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system, making them useful for analysing detailed structures.^{14–17} Lens-free imaging systems offer a wide field of view, further miniaturisation, and cost-effectiveness since they do not require magnification and objective lenses.^{18–24}

High throughput is also essential, given the need for parallel culture under multiple conditions, such as optimisation of differentiation induction. High-throughput devices capable of observing multiple regions using scanning mechanisms have been reported and commercialised.^{25,26} For instance, products such as the provi-CM20 and CM30 by Evident are representative and are already in use.²⁷ Additionally, technologies have been developed that allow simultaneous observation of multiple wells in a culture plate by aligning compact imaging units (illumination, lenses, and sensors) to match the positions of the wells.^{28–30} However, most of these devices are large and expensive, posing a challenge to miniaturisation. While these devices can be effectively utilised in labs equipped with large incubators and ample funding, there is still a significant demand for compact and high-throughput devices.

Finally, providing effective metrics for quality control is a crucial requirement. Previously, many in-incubator monitoring systems have attempted to achieve high resolution and implement fluorescence observation.^{15,16,19,20,28,29} For example, Son *et al.* developed an upright-type miniaturised microscope array mountable on a 12-well plate, achieving a single-cell resolution fluorescence imaging system that is relatively compact and high-throughput while the field of view (FOV) is small.²⁸ Such technology is expected to be utilised for detailed dynamic analysis at the cellular level. However, from the perspective of quality control in cell production processes for regenerative medicine and drug discovery, these devices may not be suitable. For instance, confluence is the most common parameter in cell culture monitoring, where a wide FOV that can capture many hiPSC colonies is more important than resolution. In addition, label-free quantifications are more crucial than fluorescence observation. Therefore, it is challenging to find a monitoring system that simultaneously meets all these requirements above.

Here, we propose an in-incubator cell imaging device that is compact, high-throughput, and capable of providing quantitative metrics for quality control of hiPSC culture and the induction processes of target cells. This device utilises lens-free imaging and notably uses a thin-film transistor (TFT) image sensor. TFT technology, primarily developed to meet the market demands of liquid crystal displays,^{31,32} is also widely used in biological research.^{33–35} Applications include electrophysiological studies,³⁶ X-ray imaging,^{37,38} bacterial colony classification,³⁹ and biochemical detection such as glucose sensors,^{40,41} immunosensors,⁴² and DNA sensors.⁴³ TFTs have the advantage of enabling the cost-effective production of large image sensors.^{32,39} In this study, we developed an image sensor with a large FOV by combining TFT and photodiode arrays. By placing cell culture chambers on the sensor, six independent culture environments can be simultaneously monitored with an FOV

of approximately 1 cm². Additionally, by implementing lens-free imaging with dark-field illumination from the rear and scattered light detection, we reduced the illumination system's size, achieving a device volume comparable to standard cell culture plates such as 6-well plates. We named this device INSPECTOR (in-incubator specialised compact lens-free TFT cell monitor).

A potential disadvantage of the TFT image sensor is its large pixel pitch, which reduces spatial resolution. At the same time, during the hiPSCs culture and target cell inductions, colony-level morphological changes and coordinated dynamic behaviours at the population level occur.⁴⁴ Therefore, even if it cannot analyse morphology at the single-cell level, achieving a wide FOV can potentially allow for the acquisition of effective metrics for quality control of the culture. To demonstrate the effectiveness of INSPECTOR, we quantified the confluence of hiPSC culture and analysed morphological changes at the colony level caused by epithelial-to-mesenchymal transition (EMT) during mesoderm induction. In addition, by implementing high-speed imaging modes, we quantified the beating dynamics of hiPSC-derived cardiomyocytes and tracked their fluctuations throughout their long-term production processes. These results indicate that INSPECTOR is a practical, high-throughput, and compact device for long-term monitoring and quality evaluation of hiPSC culture and its differentiation induction processes, highlighting its potential contribution to regenerative medicine and drug discovery.

Materials and methods

Fabrication of thin-film transistor image sensor panel

INSPECTOR's image sensor consists of a low-temperature polysilicon thin-film transistor (LTPS-TFT) circuit and an amorphous silicon (a-Si) pin photodiode formed on a glass substrate (Fig. S1†). Both the LTPS-TFT and the a-Si pin photodiode were fabricated using standard photolithography processes for liquid crystal display production. The LTPS-TFT channel layer (poly-Si) was deposited using chemical vapour deposition (CVD) and subsequently crystallised through an excimer laser annealing process. The a-Si pin photodiode was stacked on top of the TFT using CVD. The upper electrode of the a-Si pin photodiode is a transparent indium tin oxide electrode, allowing the photodiode to absorb light from above. The top layer is covered with a transparent acrylic resin for surface protection. The surface of the acrylic resin has been plasma-treated to improve the adherence of the cultured cells. All the above manufacturing processes took place at Kyocera Corporation.

Construction of the base module of INSPECTOR

The base module of INSPECTOR consists of an LED light source, a control board, a power board, a Li-ion battery, and a connection interface for the sensor module (Fig. S2†). The LED light source utilises a planar light source commonly used in LCD displays. The control board is equipped with an



integrated circuit comprising a field programmable gate array (FPGA), a microcontroller unit (MCU), and memory, and it can be connected to the sensor module *via* pogo pins. The control board performs various functions, such as receiving control signals from a PC, driving the sensor, reading light signals from the sensor, converting the light signals from analogue to digital, and transmitting the light signals to the PC. Communication with the PC is conducted *via* Bluetooth. The power board is connected to the control board through an FFC connector, manages the lithium-ion battery's voltage control, and supplies power to the control board and the light source. The outer frame of the base module is fabricated by machining polycarbonate. All the above manufacturing processes took place at Kyocera Corporation.

HiPSC culture

The hiPSC line 585A1 (ref. 45) was maintained according to the methods described in a previous report.⁴⁶ Specifically, 6-well culture plates were coated with 3 $\mu\text{g ml}^{-1}$ iMatrix-511 (982021, MATRIXOME) in D-PBS for 1 hour at 37 °C, and then hiPSCs were cultured in StemFit medium (AK02N, Ajinomoto). The medium was changed daily, and cells were passaged using TrypLE Select (A1285901, Thermo Fisher Scientific) every 4 days. Cell counting was performed using Countess (Thermo Fisher Scientific). At each passage, 2×10^5 cells were seeded. For the first 24 hours post-passage, 10 μM Y-27632 (259-00613, FUJIFILM Wako) was added to the StemFit medium. When culturing on INSPECTOR, the sensor was coated with 5 $\mu\text{g ml}^{-1}$ iMatrix-511 in D-PBS for 1 hour at 37 °C, and the desired number of cells was passaged. After the passage, cells were incubated in StemFit containing 10 μM Y-27632 for 24 hours, followed by daily medium changes without Y-27632.

Fluorescent staining and observation of hiPSCs

Two days post-seeding on the sensor module, hiPSC images were captured using INSPECTOR. Subsequently, cytoplasmic staining or nuclear staining was performed by adding 2 μM CMFDA (223-02151, FUJIFILM Wako) or 1 μM Kakshine⁴⁷ to the StemFit medium, respectively. Cells were incubated at 37 °C for 30 minutes, washed with PBS, and fixed with 4% PFA at room temperature for 15 minutes. The cytoplasm-stained cells were examined under a conventional upright microscope. For nuclear-stained cells, after removing the culture chamber, cells were mounted with Fluomount-G (0100-01, Southern Biotech), covered with a glass slide, and then high-resolution fluorescence imaging was conducted using AMATERAS.⁴⁸

Mesoderm induction

We used a base medium consisting of RPMI-1640 with HEPES (189-02145, FUJIFILM Wako) supplemented with B27 supplement without insulin (A1895601, Thermo Fisher Scientific) and penicillin-streptomycin (168-23191, FUJIFILM Wako), designated as RPMI/B27-ins. On the second day after

seeding hiPSCs on INSPECTOR, the culture medium was switched to RPMI/B27-ins with 6 μM CHIR99021 (038-23101, FUJIFILM Wako) to initiate mesoderm differentiation. Control samples were switched to RPMI/B27-ins without CHIR. The medium was not changed during the subsequent 48 hour observation on INSPECTOR.

Cardiomyocyte differentiation induction

Cardiomyocyte differentiation was induced based on temporal modulation of Wnt signalling in monolayer culture.^{49,50} RPMI/B27-ins containing 50 $\mu\text{g ml}^{-1}$ ascorbate is denoted as RPMI/B27-ins/asc. On day 0, the third day after hiPSC seeding, the medium was switched to RPMI/B27-ins/asc supplemented with 6 μM CHIR99021 to initiate differentiation. On day 1, the medium was changed to RPMI/B27-ins/asc. On day 3, for ventricular-like cardiomyocyte induction, the medium was switched to RPMI/B27-ins/asc containing 5 μM IWP-2 (034-24301, FUJIFILM Wako), 5 μM XAV-939 (247-00951, FUJIFILM Wako), and 1 μM BMS-453 (19076, Cayman Chemical).^{50,51} For atrial-like cardiomyocyte induction, the medium was switched to RPMI/B27-ins/asc containing 5 μM IWP-2, 5 μM XAV-939, and 1 μM retinoic acid (186-01114, FUJIFILM Wako).⁵⁰⁻⁵² From day 5 onwards, both ventricular-like and atrial-like cardiomyocytes were cultured under the same conditions. On day 5, the medium was changed to RPMI/B27-ins/asc. Subsequently, the medium was changed every other day to RPMI-1640 with HEPES containing B27 supplement (17504044, Thermo Fisher Scientific) and penicillin-streptomycin, which is denoted as RPMI/B27. On day 11 or 12, cells were detached using TrypLE Select and reseeded onto dishes or sensors coated with Matrigel (354277, Corning). After seeding, cells were incubated in RPMI/B27 containing 10% KnockOut Serum Replacement (10828010, Thermo Fisher Scientific) for 24 hours, followed by medium changes every other day with new RPMI/B27. For glucose-free medium, no glucose RPMI-1640 (185-02865, FUJIFILM Wako) containing B27 supplement, 5 mM sodium DL-lactate (L4263, Merck), and penicillin-streptomycin was used.

Drug treatment on cardiomyocytes

On day 23 of ventricular-like cardiomyocyte induction, cells were passaged using TrypLE Select and seeded at 2×10^5 cells per well on the sensor. After seeding, cells were cultured in RPMI/B27 containing 10% KnockOut Serum Replacement for 24 hours, followed by medium changes every other day with RPMI/B27. Drug response assays were conducted on day 30 of the induction. The medium was changed just before the experiment. Drug concentrations were adjusted by replacing half of the medium with a medium containing adequate concentrations of isoproterenol (I0260, Tokyo Chemical Industry) and propranolol (168-28071, Fujifilm). Measurements were taken 15 minutes after the medium change. All conditions were sequentially measured using the same sample, starting from the lowest to the highest concentration as follows: control, 10 nM Iso., 30 nM Iso., 100 nM Iso., 300 nM Iso., 300 nM Iso.



and 300 nM Pro., 300 nM Iso. and 1 μ M Pro., and 300 nM Iso. and 3 μ M Pro.

Image processing and quantification

To evaluate INSPECTOR's detection accuracy, we aligned the fluorescence image with the corresponding INSPECTOR image, manually identifying the same locations in both. Given the fluorescence image's 1.1 μ m pixel pitch, it was segmented into a 59 pixel grid, matching each grid area with an INSPECTOR pixel. Colony areas in the fluorescence image were delineated using a Gaussian filter for smoothing followed by baseline subtraction with the rolling ball algorithm, binarising the image by setting an intensity threshold, applying the watershed algorithm to segment, and extracting spatially connected areas as individual colonies. For confluence quantification, the rolling ball algorithm was used to obtain and subtract a baseline from the image, and the same threshold was manually set across all conditions. Regions of interest (ROI) were set within the well, excluding the chamber edges. The proportion of pixels with signals exceeding the threshold was calculated. For quantification of EMT during mesoderm induction, the Sobel filter was used to compute the first derivative and visualise the spatial gradient. ROIs were set within the well, excluding the chamber edges. The average value of the first derivative image in each ROI was divided by the mean brightness of the original image and then normalised with the value of the first time frame. To quantify cardiomyocyte beating dynamics, low pass filtering was applied to the temporal intensity fluctuations in each pixel to remove high-frequency noise and identify the baseline. The change ratio from the baseline was calculated pixel by pixel to normalise the intensity fluctuations. A fast Fourier transform analysis was performed on the pixel-level intensity fluctuations, and the dominant frequency in the movie was analysed by averaging the results from all pixels. Peak detection was performed on the pixel-level intensity fluctuation to visualise the contraction conduction based on the differences in peak timing. All the processes are carried out using Fiji⁵³ and Python 3.7.

Results

Design of INSPECTOR

INSPECTOR is a lens-free imaging device designed for the long-term monitoring of hiPSC culture and the production processes of target cells within an incubator. It is designed to employ a novel illumination strategy, which illuminates the cells from the rear and detects the scattered light using a TFT image sensor, optimising light capture (Fig. 1A). The device comprises a sensor module for cell culture and light detection, as well as a base module for light irradiation, data storage, and transfer (Fig. 1B, S1 and S2†).

The sensor module includes an image sensor deposited on a glass substrate and a connection to the base module. The image sensor pixels are composed of low-temperature poly-silicon thin-film transistors (LTPS-TFT) and thin-film amorphous Si photodiodes (Fig. S1†). 580 \times 328 pixels are

arrayed with a pitch of 65 μ m, yielding a sensor area of approximately 37.7 \times 21.3 mm (Fig. 1B and C). A 10 μ m gap between the pixels allows light irradiation from the rear (Fig. 1A and C).

Due to advancements in TFT manufacturing technology, the sensor module can be mass-produced cost-effectively and is designed as a disposable consumable.^{32,39} Consequently, cells can be cultured directly on the sensor, resulting in proximity to the sensor corresponding to the thickness of the surface acrylic resin layer of 4 μ m (Fig. 1A). The size of the sensor module is approximately 76 \times 26 mm, equivalent to a standard slide glass. As a result, common cell culture chambers can be attached. When a typical 8-well type chamber (#30118, SPL LIFE SCIENCES) is placed, the sensor can cover an area equivalent to 6 wells (Fig. 1D). Therefore, the six independent culture environments of approximately 1cm² can be simultaneously monitored. The remaining two wells are covered only by glass, allowing observation of control samples with a conventional inverted microscope (Fig. 1D).

The base module features an LED irradiation surface, an internal battery, data storage, and a connection to the sensor module (Fig. 1B and E, S2†). The proximity of cells to the sensor in the sensor module reduces image blurring, which might otherwise be caused by distance. This proximity eliminates the need for coherent light illumination or computational image reconstruction, often required in lens-free imaging.²⁴ Therefore, to minimize the device and simplify the imaging processes, we employed a novel illumination strategy. Specifically, we employed dark-field illumination from the rear and scattering detection, thereby miniaturising the base module's illumination system. To suppress background signals during cell observation and enhance the detection of scattered light, the LEDs are designed to irradiate obliquely through a louvre film (Fig. 1A). Data obtained from the sensor module is saved in the base module. Light irradiation, data acquisition, and data transfer are remotely controlled by an external PC connected *via* Bluetooth low energy (BLE). If a direct BLE connection with the PC outside the incubator proves challenging, a BLE antenna wired to the PC can be installed inside the incubator (Fig. S3†). These features enable real-time monitoring of the culture.

Fig. 1F illustrates the appearance of INSPECTOR installed inside an incubator. The XY plane is approximately 48 cm² (9.5 \times 5.0 cm), with a maximum height of about 4.5 cm to the top of the lid, occupying a volume of less than 220 cm³. Compared to a standard 6-well plate (e.g., 12.8 \times 8.5 \times 2.2 cm), the XY plane of INSPECTOR is less than half the size, yet the height is approximately double, achieving a compact design with a comparable volume. Therefore, despite limited space inside an incubator, multiple devices can be easily installed, facilitating effective high-throughput monitoring.

To assess the spatial resolution of INSPECTOR, we analysed a USAF 1951 target pattern (R3L1S4P, Thorlabs) on transparent glass comprising three lines of specific widths that reflect light at intervals matching their widths (Fig. 1G–J). We evaluated whether these lines could be distinctly recognised (Fig. 1I and J). By removing the sensor module's cell culture



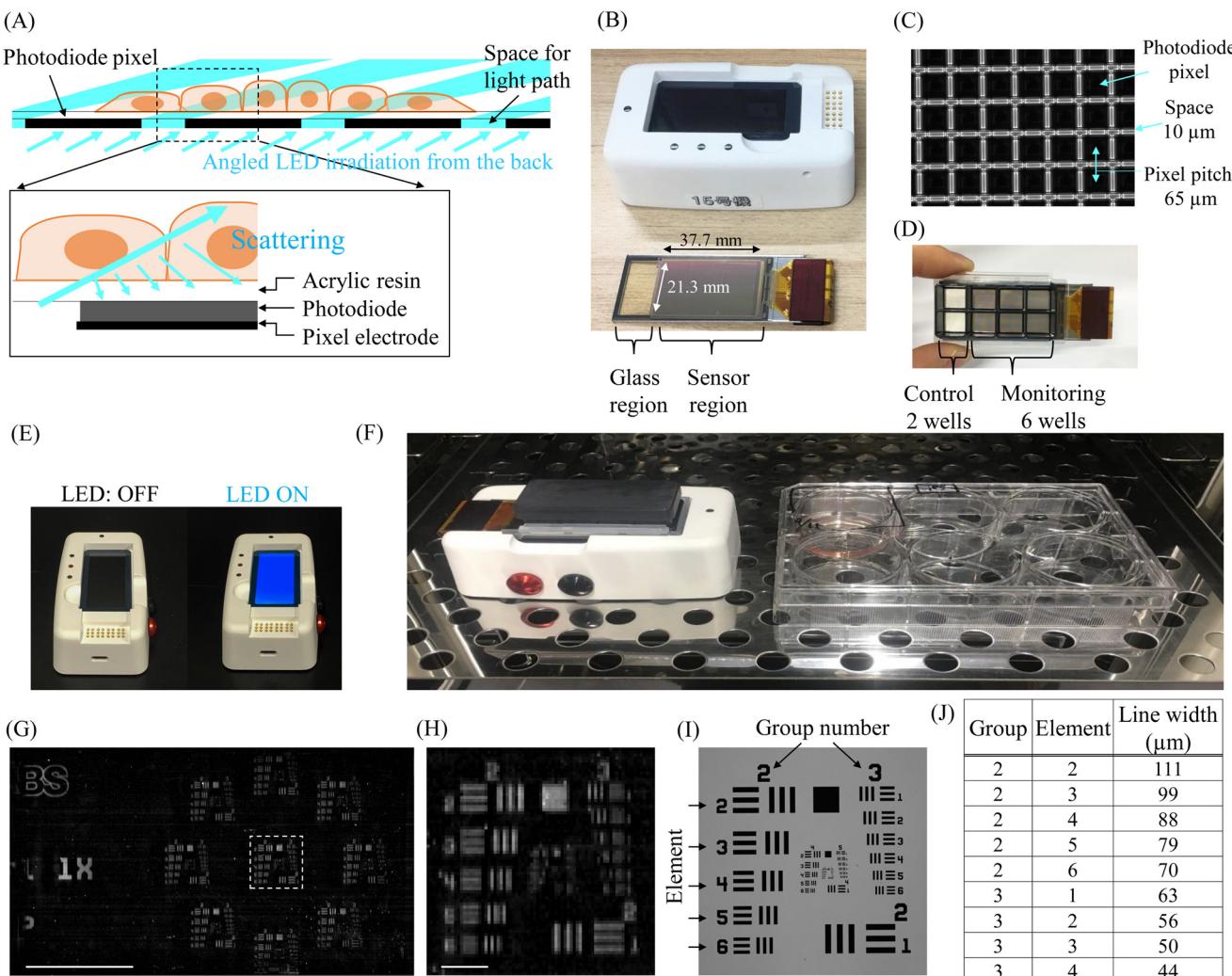


Fig. 1 Design and appearance of INSPECTOR: (A) schematic illustration of the imaging mechanism. Scattered light from cells is detected on the surface of the sensor pixels. (B) Appearance of the base module (top) and the sensor module (bottom). (C) Magnified image of the image sensor pixels. (D) Image of a sensor module attaching an 8-well type cell culture chamber. (E) LED illumination from the base module. (F) The appearance of INSPECTOR putting inside an incubator. A 6-well plate was also placed to compare the sizes. (G) An entire FOV of the INSPECTOR image observing the target pattern USAF 1951. (H) An enlarged image of the white dot square region in G. (I) A transmitted bright field high-resolution image of the same region as H. (J) A table of line width in each group and element. Scale bars: 1 cm in G, 1 mm in H.

chamber and substituting a glass slide with the pattern, Fig. 1G illustrates the full FOV where INSPECTOR detected light from the pattern. An enlarged view (Fig. 1H) reveals that vertical lines of 56 μm width (group 3, element 2) and horizontal lines of 79 μm width (group 2, element 5) were distinguishable, supporting that INSPECTOR spatial resolution approaches its pixel pitch, aligning with the optical detection principle of the device.

In the following sections, we will present examples of applications of INSPECTOR for long-term monitoring of hiPSC culture and its differentiation induction.

Long-term confluence monitoring of hiPSC culture

Confluence is a widely used quantitative indicator for monitoring cell cultures. Particularly in hiPSC cultures, quantifying confluence is valuable for determining the appropriate passage timing for maintaining undifferentiated cells^{45,46} and adjusting

the culture density for optimal differentiation induction.^{49–52,54,55} Therefore, we attempted to quantify confluence during hiPSC culture using INSPECTOR.

INSPECTOR detects scattered light, producing an image with a dark background and bright regions where scattering objects such as cells and chamber wells exist. Fig. 2A shows a full FOV image of undifferentiated hiPSCs cultured on and observed by INSPECTOR. Six culture areas can be easily identified from the scattered light at the edges of the wells. Since hiPSCs proliferate, forming dense colonies, they are expected to scatter light strongly and can be sufficiently detected by the resolution of INSPECTOR (Fig. S4†).

Indeed, distributed bright signals were detected within the wells (Fig. 2A). Upon enlarging specific regions within wells, signals presumably originating from hiPSC colonies were identified with a sufficient signal-to-noise ratio (Fig. 2B). To confirm that these signals originate from hiPSC colonies,



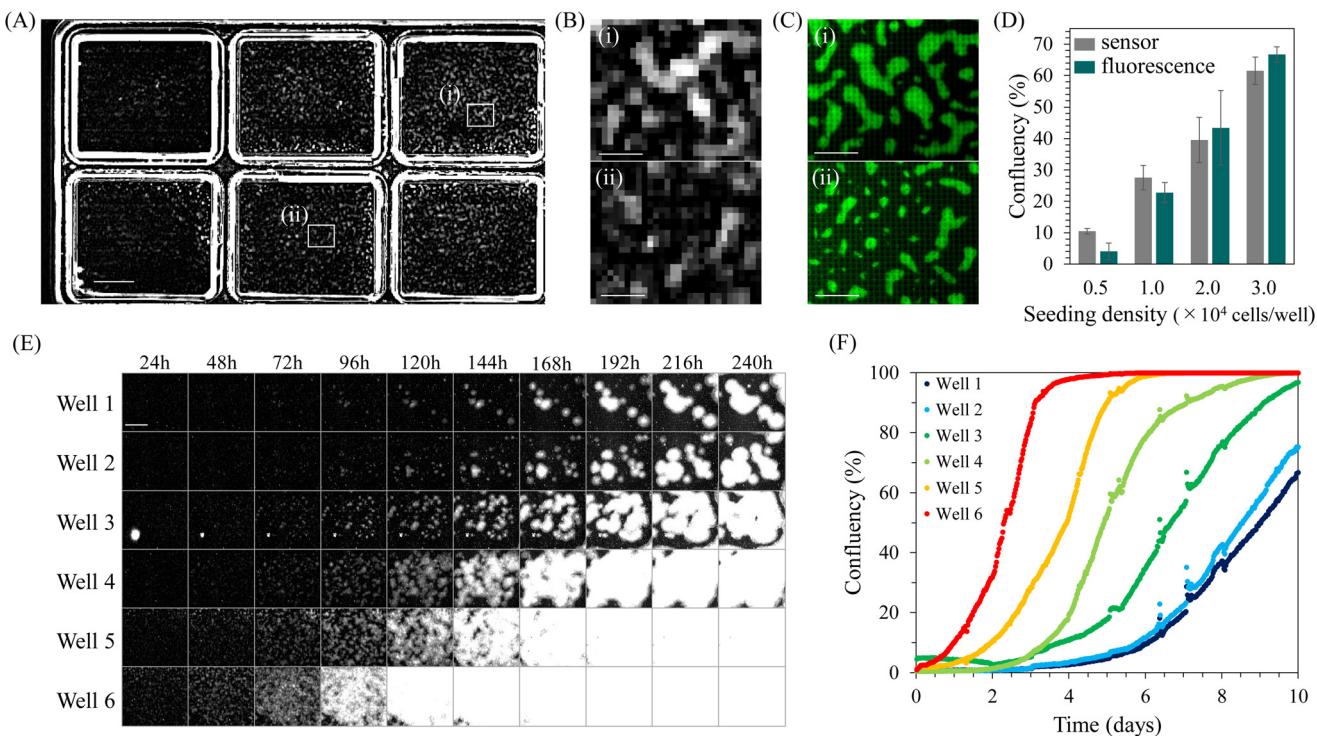


Fig. 2 Quantification of confluence in hiPSC culture: (A and B) a full field-of-view image (A) and zoom in images (B) of hiPSC culture obtained by INSPECTOR. The white squares (i) and (ii) in A are enlarged in B. (C) Cytoplasm-stained fluorescence images of the same areas as (i) and (ii) in A and B. (D) Comparison in confluence quantified based on the INSPECTOR images and fluorescence images from the same samples. Error bars represent standard deviation ($n = 4$). (E) Example images of hiPSC culture with different seeding cell densities taken during long-term monitoring. (F) Plots in confluence over time. Image backgrounds in A, B and E were subtracted by image processing. Scale bars: 3 mm in A and E, 500 μ m in B and C.

cytoplasm of the same sample was fluorescently stained and observed under a conventional upright microscope (Fig. 2C). Comparing the two, it was found that regions of hiPSC colonies confirmed by fluorescence are also detected by INSPECTOR despite the coarser images due to the pixel pitch (Fig. 2B and C).

To evaluate the cell detection accuracy of INSPECTOR, we performed a detailed comparison with a nuclear-stained fluorescence image offering subcellular resolution (Fig. S5†). Initially, we quantified the fluorescence intensity for areas corresponding to each INSPECTOR pixel and found a positive correlation with INSPECTOR's pixel intensity (Fig. S5G†). We established thresholds for signal intensities to assess cell presence, using fluorescence-based judgment as a benchmark. INSPECTOR achieved approximately 90% detection accuracy (Fig. S5G†). An investigation into detection errors revealed that false positives often arose from background noise (cyan arrowheads in Fig. S5B and E†) or from INSPECTOR's overestimation of large colony contours (cyan arrowheads in Fig. S5C and F†). False negatives primarily occurred when INSPECTOR missed small colony regions (white arrowheads in Fig. S5C and F†). Further analysis demonstrated INSPECTOR's signal intensity with colony size, detecting all colonies over 21 000 μ m², though efficiency diminished with smaller colonies (Fig. S5H and I†). These findings confirm INSPECTOR's high detection accuracy for hiPSCs, which typically form large colonies, despite some inaccuracies.

Next, we quantified confluence from the images obtained by INSPECTOR and compared the results with those quantified based on fluorescence images. As a result, INSPECTOR showed confluence values comparable to those quantified from fluorescence images (Fig. 2D). This result demonstrates the sufficient capability of INSPECTOR to quantify the confluence of hiPSC culture.

Given the above results, we next conducted long-term monitoring of confluence during the undifferentiated culture of hiPSCs. Here, hiPSCs were cultured at six different seeding densities, and images were captured every 30 minutes for 10 days (Fig. 2E and F). Fig. 2E displays the extracted regions of each well, showing the colonies grow larger over time. Plotting quantified confluencies successfully visualised the difference in timing of increase depending on the seeding density (Fig. 2F). These results indicate that INSPECTOR can be effectively utilised for long-term quantitative monitoring of confluence during hiPSC culture.

Monitoring of EMT during mesoderm induction

Next, we attempted to quantify changes in the cellular state during differentiation induction using INSPECTOR. As an example, we specifically observed mesoderm induction from hiPSCs. The mesoderm is a germ layer that gives rise to various cells and organs, such as the heart, blood cells, kidney, muscle, osteocytes, and adipocytes. It can be induced

from pluripotent stem cells *in vitro*.^{54–56} During the induction, EMT occurs as an initial cell state change. Due to EMT, cells tightly bound in colonies acquire migratory properties, spreading outside the colonies, causing the colony morphology to collapse and cells to distribute in a flattened manner.⁵⁵

Here, we attempted to quantify this EMT using INSPECTOR. HiPSCs were induced into mesoderm through Wnt signal stimulation using 6 μ M CHIR99021 (CHIR), and the process was monitored every 30 minutes over a period of 48 hours. In a control sample without CHIR, the morphology of colonies remained largely unchanged over 48 hours, accompanied by an increase in average intensity (Fig. 3A). In contrast, in the CHIR-treated samples, colonies collapsed in their shape during the latter stage, and the original morphology was lost after 48 hours (Fig. 3B). We used a first derivative filter to calculate the local spatial gradient and quantified this morphological change (Fig. 3C and S6†). Strong gradients around the contour of the maintained colonies were detected in the control sample (Fig. S6A†), while a smaller gradient was quantified in the CHIR-treated group (Fig. S6B†), indicating that morphological differences were successfully quantified. Fig. 3C and D showed that the local gradient in the control increased monotonically over time, suggesting continued cell proliferation and the formation of denser colonies. In contrast, the spatial gradient in the CHIR-treated samples decreased from the midway point around 30 hours after mesoderm induction (Fig. 3D).

asterisk). This indicates that the acquisition of migratory properties and changes in colony shape due to EMT begin at this time.

These results demonstrate that INSPECTOR can accurately identify the initiation timing of EMT, a typical cellular state change occurring during mesoderm induction from hiPSCs. By using this timing as an indicator, researchers can effectively conduct intermediate evaluation and assess the reproducibility of the differentiation induction process.

Quantification of beating dynamics of hiPSC-derived cardiomyocytes

Next, to illustrate how we can quantify functional indicators of differentiation products, we attempted to quantify the beating dynamics of cardiomyocytes. Cardiomyocytes are a type of cell that can be effectively induced from hiPSCs *in vitro* and thus are expected to play significant roles in drug discovery and regenerative medicine.^{57–59} HiPSC-derived cardiomyocytes exhibit spontaneous beating at the cellular level.⁶⁰ Their beating dynamics are closely related to electrophysiology and are useful for evaluating cellular states and drug responsiveness.⁶¹

First, to accurately capture the rapid beating dynamics of cardiomyocytes, we implemented high-speed imaging modes. The maximum imaging speed is determined by the image readout speed, which can be improved by reducing the FOV. Here, we implemented two FOV restriction modes: single-

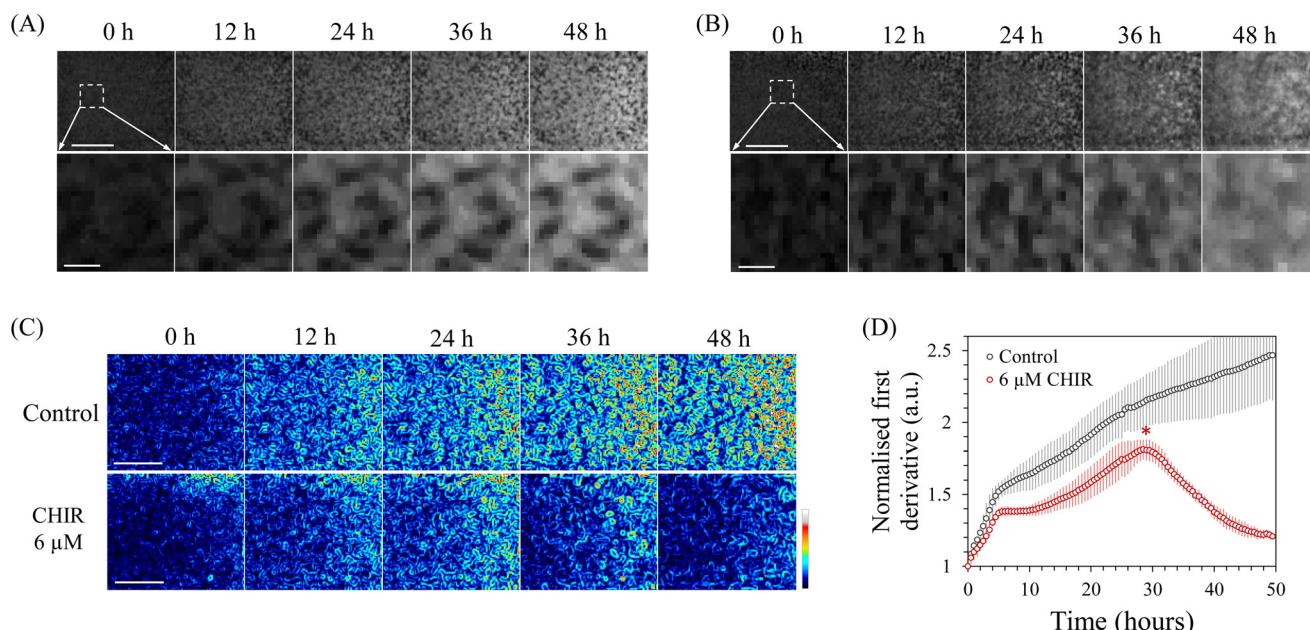


Fig. 3 Quantification of morphological change of colonies during mesoderm induction: (A and B) cropped INSPECTOR image of a control sample (A) and 6 μ M CHIR treated sample (B). The top images show almost all of a region within a chamber, and the bottom images zoom in on the dot square area in the top image. (C) Visualisation of intensity gradient by normalised first derivative images of (A) top and (B) top. Pixel intensity is visualised as colour. (D) Quantification of change in mean intensity of the normalised first derivative images. The asterisk represents the timing of morphological change corresponding to the initiation of EMT. Error bars represent standard deviation ($n = 4$). Scale bars: 3 mm in A top, B top, and C, 500 μ m in A bottom and B bottom.



Table 1 The single-area modes

	Area (pixel)	Area (mm)	Speed
Mode 1	32 × 32	2.08 × 2.08	10.8 fps
Mode 2	48 × 48	3.12 × 3.12	10.1 fps
Mode 3	64 × 64	4.16 × 4.16	8.2 fps

area and six-area modes. The single-area mode limited the FOV to a small area, 2.08 × 2.08, 3.12 × 3.12, or 4.16 × 4.16 mm square, which enabled observations at 10.8, 10.1, or 8.2 fps respectively (Table 1). In the six-area mode, six different areas corresponding to the six wells on the sensor can be captured at high speeds simultaneously, allowing for the comparison of multiple culture conditions. Each area was 0.78 × 0.78, 1.04 × 1.04, or 1.56 × 1.56 mm square in size, achieving 25.4, 12.9, or 8.8 fps respectively (Table 2).

Using these high-speed imaging modes, we observed cardiomyocytes induced from hiPSCs directly on the sensor. At days 10–11 of cardiomyocyte induction, spontaneous contraction-relaxation cycles were observed in most of the cells within the control well, as observed under a conventional inverted microscope, indicating highly efficient cardiomyocyte induction (Movie S1†). High-speed imaging with the single-area mode of INSPECTOR revealed periodic increases in image intensity (Fig. 4A). This increase likely reflects enhanced scattered light intensity when the cardiomyocytes contract and become denser. Analysing this intensity change with the fast Fourier transform (FFT) enabled quantification of the beating frequency (Fig. 4B). As a result, multiple peaks appeared at frequencies that were integer multiples of the lowest frequency observed. This occurs because the original intensity change displayed sharp spikes, unlike a sine curve (Fig. S7†). The peak showing the lowest frequency corresponded to the beating frequency (Fig. 4B asterisk). In addition, the Single-area mode with a FOV exceeding 4 mm² allowed for spatiotemporal analysis, which enabled the visualisation of the beating conduction patterns (Fig. 4C and D). Interestingly, around days 10–11 of the differentiation induction, reentry patterns characterised by circulating beating propagation were observed in approximately half of the samples (Fig. 4E and F, Movie S2†). The reentry pattern is thought to arise from fragmented propagation waves in heterogeneous environments.⁶² The observed reentry was likely due to temporary spatial heterogeneity occurring during the cardiomyocyte differentiation process. These results demonstrate that INSPECTOR can effectively quantify the beating frequency and conduction as functional indicators of cardiomyocytes derived from hiPSCs.

Table 2 The six-area modes

	Area (pixel)	Area (mm)	Speed
Mode 1	12 × 12	0.78 × 0.78	25.4 fps
Mode 2	16 × 16	1.04 × 1.04	12.9 fps
Mode 3	24 × 24	1.56 × 1.56	8.8 fps

Next, we assessed the drug response in cardiomyocytes using INSPECTOR by quantifying their beating frequency. We reseeded hiPSC-derived cardiomyocytes as monolayer cultures with spatially uniform distribution (Movie S3†). These cells were then used on day 30 of cardiomyocyte induction. No reentry patterns were observed, and the propagation speed was significantly faster than that on days 10–11, showing almost simultaneous contractions (Fig. 4G and H). Here, we analysed the chronotropic response to β-adrenergic receptor perturbation using isoproterenol as the agonist and propranolol as the antagonist (Fig. 4I). As a result, when the isoproterenol was applied, a concentration-dependent increase in beating frequency was confirmed. Moreover, maintaining a high concentration of isoproterenol and subsequently adding the propranolol led to a concentration-dependent decrease in beating frequency. These results are consistent with the reported chronotropic response to isoproterenol and propranolol.⁵⁸ Therefore, it can be concluded that INSPECTOR can evaluate drug response based on the beating frequency of hiPSC-derived cardiomyocytes.

Detection of beating onset during cardiomyocyte differentiation from hiPSCs

In the context of cardiomyocyte production for regenerative medicine and drug discovery, efficient and rapid induction reduces production costs and is thus valuable. When optimising the induction conditions, the timing of beating initiation indicates the speed of cardiomyocyte induction. Therefore, we attempted to detect the onset of spontaneous beating during cardiomyocyte induction using INSPECTOR. Here, we observed two differentiation conditions that induced cardiomyocytes with a bias toward atrial- and ventricular-like phenotype. We monitored these conditions from day 5 to 12 of differentiation using the six-area mode with high-speed imaging at 8.8 fps, capturing images every two hours.

In the atrial and ventricular induction conditions, initiation of spontaneous beating was successfully detected on days 9 and 7, respectively (Fig. 5A). Generally, hiPSC-derived cardiomyocytes initiate beating between days 7 and 12, which is consistent with our results.^{49–52} The observed beating frequencies in the ventricular condition fluctuated between 0.6 and 1 Hz (Fig. 5A green plots). In contrast, the atrial condition initially exhibited a slightly higher frequency of approximately 2.3 Hz, which rapidly decreased to below 1.0 Hz. Similar fluctuations repeated after the medium change (Fig. 5A magenta plots). These results demonstrate that INSPECTOR can detect the onset of spontaneous beating and the differences in beating frequency fluctuations. Thus, these timing and patterns can be utilised to assess the speed, reproducibility, and quality of cardiomyocyte induction.

Long-term monitoring of beating frequency during cardiomyocyte maturation

Next, we conducted a long-term time-lapse observation of the maturation process of cardiomyocytes. A prolonged culture is



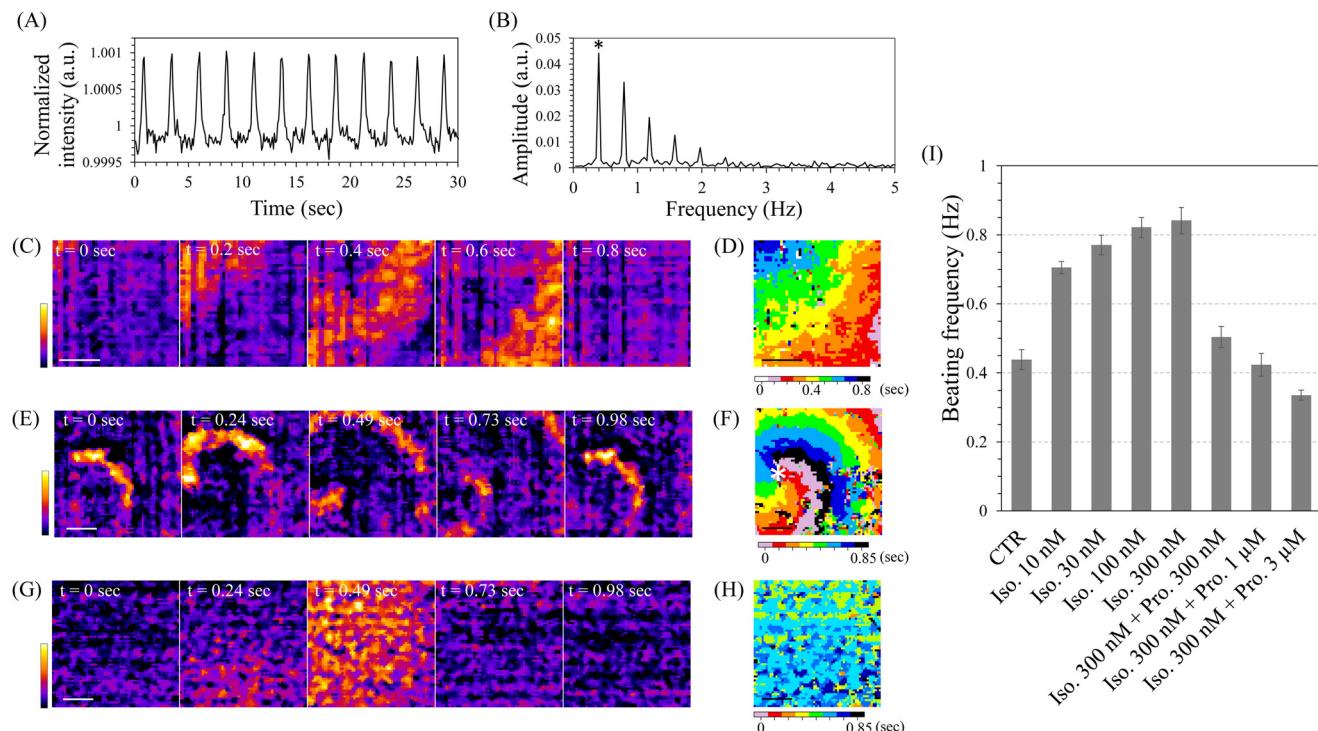


Fig. 4 Quantification of beating dynamics of hiPSCs-derived cardiomyocytes: (A) quantifying intensity fluctuation in a movie of hiPSC-derived cardiomyocytes on day 11 of differentiation. (B) A result of fast Fourier transfer analysis of A. The asterisk represents a frequency corresponding to the beating rate. (C, E and G) Visualisation of contraction conduction. Intensity changes as a unidirectional propagation (C) and a reentry propagation (E) on day 11 and as an almost simultaneous contraction on day 30 (G) were shown in colour. (D, F and H) The peak timings of intensity increases in C, E, and G were represented as colours to visualise spatial contraction patterns. The asterisk in F represents the centre of the rotation (phase singularity). (I) Beating frequency upon drug treatment. Iso.: isoproterenol, Pro.: propranolol. Error bars represent standard deviation ($n = 5$). Scale bars: 1 mm.

essential for producing mature cardiomyocytes.⁶³ Here, cardiomyocytes induced toward atrial- and ventricular-like phenotypes were reseeded on the sensor on day 11 of the induction and observed for approximately two weeks starting from day 12. Images were captured every 2 hours at 8.8 fps using the six-area mode (Fig. 5B). As a result, stable beatings were consistently detected from day 14 under the ventricular condition and from day 15 under the atrial condition. The cell state recovery and the time until reaching the confluence monolayer upon proliferation (Fig. S8†) may contribute to this delayed detection of beating after passage. Subsequently, cardiomyocytes exhibited two major patterns in the fluctuation of beating frequency: long-term and short-term changes.

As for the long-term change, in the ventricular condition, the beating frequency gradually decreased to stabilise around 0.5 Hz (Fig. 5B green plots). In contrast, under the atrial condition, the frequency gradually increased and reached 1.4 Hz by day 26 but had not yet stabilised (Fig. 5B magenta plots). HiPSC-derived atrial cardiomyocytes typically exhibit a higher beating frequency than ventricular cardiomyocytes, which is consistent with our results.⁵² However, whether the frequency has stabilised can only be determined through long-term monitoring, as demonstrated by our observation.

The short-term change was characterised by rapid frequency increases observed every other day, followed by a gradual return to baseline within approximately one day (Fig. 5B-i and ii). This frequency increase coincided with the timing of medium changes, as indicated by arrowheads in Fig. 5B. We speculate that this change could be attributed to fluctuations in pH resulting from glucose metabolism. It is known that the beating frequency of cardiomyocytes is pH-dependent; it increases under alkaline conditions and decreases under acidic conditions.^{64,65} Additionally, lactate accumulates in the medium due to glucose metabolism of cardiomyocytes, leading to a decrease in medium pH.^{66–68} The culture medium contained 5 mM glucose (high-Gluc medium). Therefore, the observed phenomenon is consistent with the assumption that the medium, initially neutral after a medium change, gradually becomes acidic due to lactate accumulation from glucose metabolism.

To test this hypothesis, we used a medium without glucose (Fig. 5C). Cardiomyocytes are known to metabolise lactate to produce ATP *via* the TCA cycle and survive for extended periods without glucose.⁶⁶ Thus, we switched to a glucose-free, 5 mM lactate (no-Gluc) medium from day 17 in the ventricular condition and from day 18 in the atrial condition. Interestingly, in the no-Gluc medium condition, the frequency rapidly decreased upon each medium change



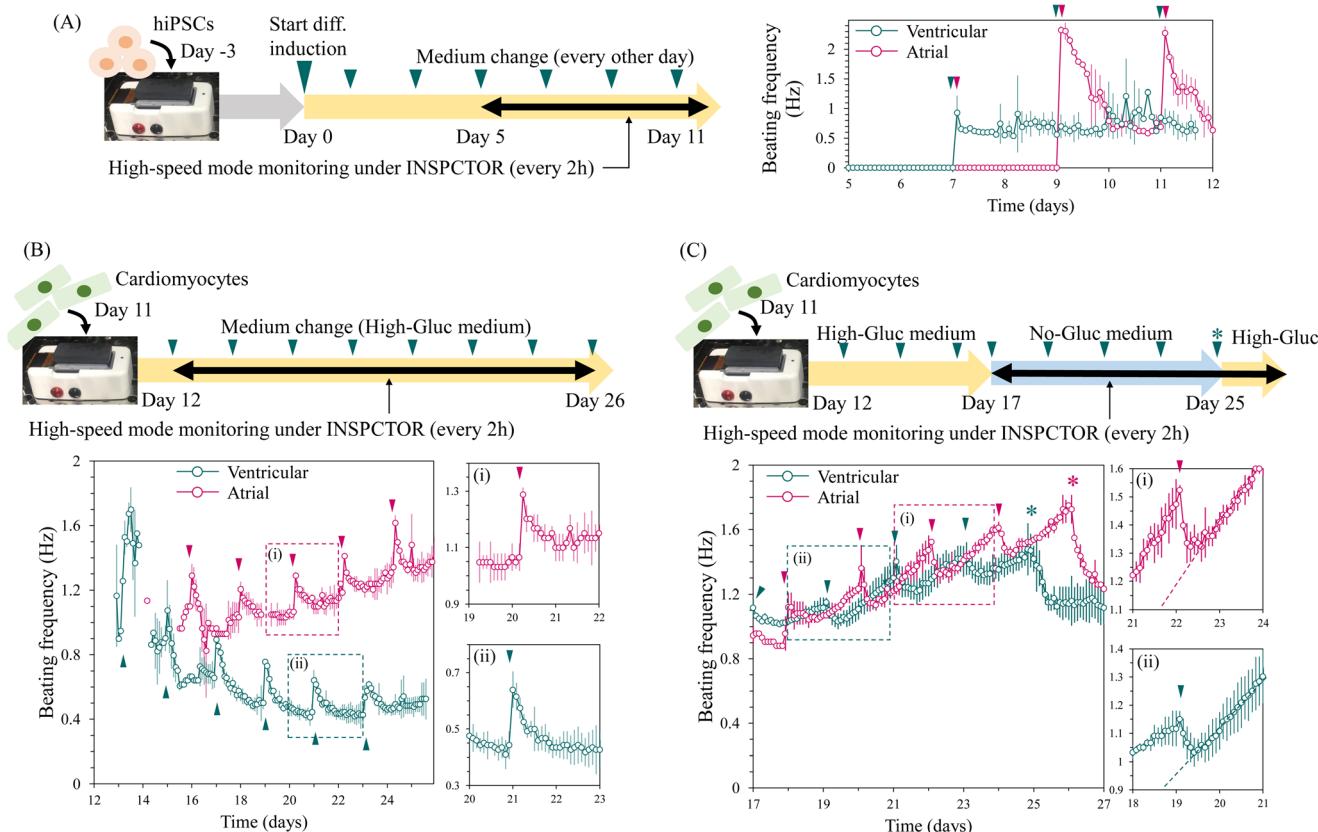


Fig. 5 Long-term monitoring of beating dynamics of hiPSCs-derived cardiomyocytes: (A–C) beating frequencies of cardiomyocytes during the atrial (magenta) and the ventricular (green) differentiation induction were quantified. The results of an early stage (A), a later stage in the high-Gluc medium (B), and a later stage in the no-Gluc medium (C) were represented. Regions of (i) and (ii) in the left plots of B and C were magnified in the right plots. Arrowheads represent the timing of the medium change. Asterisks in C represent the timing to return the medium to the high-Gluc medium. Error bars represent standard deviation ($n = 3$).

(Fig. 5C). After reaching the baseline, the frequency increased almost linearly (Fig. 5C-i and ii). These changes may reflect a decrease in lactate concentration upon metabolic consumption and a shift towards alkaline pH over the two days before the medium change. These observations do not contradict our hypothesis that metabolic activity leads to changes in pH and beating frequency. In addition, under both atrial and ventricular conditions in the no-Gluc medium, the frequency continued to increase without stabilising but began to decrease after the medium was changed to the high-Gluc medium (Fig. 5C asterisk). This suggests that metabolic differences that occurred in the no-Gluc medium induced the long-term and non-stabilizing increase in beating frequency.

These results demonstrate that INSPECTOR effectively tracks beating frequency during the maturation process of cardiomyocytes. The long-term changes allow us to determine if the beating frequency has stabilised, which indicates the appropriate timing of downstream analysis, such as drug evaluation. Additionally, the daily responses to medium changes suggest the capability to quantify metabolic-dependent fluctuations. Thus, further research is anticipated

to provide more concrete interpretations and practical indicators of cellular metabolic state in the future.

Discussion

We have developed a lens-free imaging device named INSPECTOR. Despite being as compact as a standard 6-well plate, it can simultaneously observe six independent culture environments. This throughput is attributable to the strength of TFT technology, which enables the cost-effective production of large image sensors.^{32,39} Additionally, INSPECTOR does not require an objective lens and utilises LEDs for illumination, making it an inexpensive device. Multiple devices can be installed at a low cost, enabling easy expansion of monitoring samples. Particularly in long-term differentiation induction processes, it is common for multiple batches to be cultured in parallel, as the next batch often begins before the completion of a previous batch. Therefore, rather than increasing the throughput by enlarging the device, utilising multiple compact devices with adequate throughput is more practical for operational efficiency. Based on these factors, INSPECTOR can be considered an effective and affordable device for monitoring



differentiation induction within the limited space of an incubator.

An ideal incubator imaging system for quality control in long-term cell culture combines compact size, high throughput, and a larger FOV. Current systems fall into three categories: single-FOV, multi-FOV, and scanning mechanism systems. Single-FOV systems, while compact and capable of a wide FOV, often fall short of throughput capabilities.^{14–24} On the other hand, systems with scanning mechanisms excel in throughput and FOV but typically struggle with bulkiness.^{25–27} This study introduces a multi-FOV system designed to address these limitations, offering a balanced solution that maintains compactness while enhancing throughput and maximising FOV.

Several multi-FOV in-incubator imaging systems have been developed to improve observation throughput by incorporating dedicated imaging units for each culture well. For instance, miniaturised modular-array microscopy (MAM) utilises 12 small imaging units to serve a 12-well plate,²⁸ while EmSight employs imaging units designed for Fourier ptychographic microscopy to simultaneously observe six samples in a 6-well plate.²⁹ Each system, however, faces distinct challenges. MAM, targeted primarily at single-cell analysis, offers a limited FOV of 0.75×0.45 mm and requires individual focusing for each well, struggling with focal drift during medium exchanges due to temperature fluctuations. Conversely, EmSight, despite providing a wide FOV and high spatial resolution through variable illumination angles and digital refocusing, involves complex hardware with six sets of objective lenses, tube lenses, and CMOS cameras, complicating miniaturisation and simplification efforts.

Unlike traditional array-type multi-FOV systems that require a dedicated imaging unit for each well, INSPECTOR employs a simpler, more efficient design, utilising just one image sensor and a single planar light source. This system features innovative backlight illumination through pixel gaps, which avoids the shadows cast by chamber walls typical in top-down illumination setups. This enables uniform lighting across all wells without the need for individual light sources. The lens-free design of INSPECTOR also contributes to its compact size, as it eliminates the bulk of typically added lenses. Additionally, the use of a large TFT image sensor allows for multi-FOV imaging with a single sensor, simplifying the overall design. A significant advantage of INSPECTOR is its ability to culture cells directly on the sensor, removing the necessity for focusing mechanisms and thus preventing focal drift during prolonged observations. These advancements render INSPECTOR not only simpler and smaller but also more practical compared to other multi-FOV systems.

As highlighted previously, the INSPECTOR system achieves the simultaneous observation of six wells with a large FOV while retaining the simplicity and compactness typical of a single-FOV system. This design effectively fulfils the three key requirements for optimal in-incubator long-term cell culture

monitoring: compact size, high throughput, and a larger FOV. Hence, INSPECTOR exemplifies a high-level integration of these critical attributes, making it an exemplary tool for advanced quality control in long-term cell culture.

INSPECTOR is designed for the disposable use of the sensor module. Unlike applications involving long-term differentiation induction with low subculture frequency, handling cells with high subculture frequency could significantly increase the costs associated with disposable components. The sensor module can withstand autoclave sterilisation and can be reused if properly cleaned. Furthermore, without intensive cleaning, reusing the same well by reseeding cells after detachment with trypsinisation, as a common practice to economise on culture plates, is feasible. However, in either case, remaining dead cells or debris can scatter light and become a noise source, so caution is required for reuse. Theoretically, it is also possible to reuse the sensor module by applying a thin, easily peelable film to the surface. However, reflections due to differences in refractive index between the film and the image sensor surface would need to be managed, possibly using optical adhesives. Moreover, a thick film layer can blur images, necessitating either an extremely thin film or computational methods to correct blurring. Overall, reusing the sensor module seems feasible, depending on the approach and purpose.

One of the main challenges with our INSPECTOR is the low image resolution caused by the $65\text{ }\mu\text{m}$ pitch of the TFT image sensor. Due to the large pixel size, some cells may not be directly illuminated. In fact, a positive correlation was observed between colony size and INSPECTOR signal intensity; however, detection efficiency significantly decreased for small colonies that were the same size or smaller than the pixel size (Fig. S5†). On the other hand, light scattered by the illuminated cells can indirectly irradiate surrounding cells, potentially enhancing detection efficiency. Indeed, the signal intensity in large colonies increased, enabling reliable colony detection, although it often extended beyond the actual borders of the colonies (Fig. S5†).

The primary concern is how many valuable metrics can be obtained given this resolution. Moreover, unlike in studies of molecular mechanisms with fluorescence imaging, non-label quantification is required for quality control in differentiation induction. Despite the above concerns, we successfully demonstrated the acquisition of multiple useful metrics in addition to confluence. For example, INSPECTOR has demonstrated its effectiveness during mesoderm induction. Mesoderm induction is a shared initial process of producing various mesodermal-derived cell types. This induction period strongly influences subsequent fate determination, making the optimisation of this process crucial.^{10,12,49–52,54–56} By quantifying changes in colony morphology, INSPECTOR successfully identified the onset timing of EMT during mesoderm induction. This can serve as a versatile metric in various mesodermal-derived cell differentiation induction processes. In the future, elucidating the relationship between EMT quantification and



target cell induction efficiency could provide additional metrics for evaluating induction quality beyond just reproducibility. Additionally, by implementing high-speed imaging modes in INSPECTOR, we effectively monitored the onset timing of spontaneous beating, conduction patterns, and beating frequency over extended periods during cardiomyocyte production. The onset timing of spontaneous beating serves as an indicator of cardiomyocyte induction speed. The difference in conduction patterns is known to cause variations in maturation speed,⁶⁹ potentially making them valuable indicators for assessing maturation. Furthermore, reentry, where conduction circulates in space, is considered a mechanism for various arrhythmias,^{70,71} suggesting potential applications for evaluating drug efficacy against arrhythmias. Additionally, we demonstrated that beating frequency fluctuation reflects metabolic activity. Changes in metabolic state are known to be essential for the maturation of cardiomyocytes.⁶³ Although a detailed analysis was not conducted in this study, quantifying the dynamics of beating frequency fluctuation could indicate cellular metabolic state and be used to determine maturity in the future.

Our TFT image sensor has a 65 μm pitch (390 ppi), but recent developments are advancing toward higher resolution TFTs in response to the demand for head-mounted displays. State-of-the-art high-resolution TFTs include Samsung's 2250 ppi (ref. 72) and BOE's 2117 ppi.⁷³ Although these are designed for colour displays, they would correspond to a pixel pitch of approximately 4 μm if used for monochrome applications like INSPECTOR. The pixel size and photodiode spacing are determined by the precision of photolithography processes, and the aforementioned ultra-high-resolution TFTs achieve a pixel spacing of approximately 1 μm . The development of high-resolution imaging devices using such high-resolution TFT image sensors is anticipated in the future.

Looking ahead, INSPECTOR is anticipated to be versatile in monitoring diverse cell populations. Although not ideal for analysing small or intricate structures, its design excels in observing larger objects due to its wide FOV. Consequently, its potential applications include monitoring diverse types such as organoids,² wound healing assays,⁷⁴ bacterial colonies,³⁹ and so on.

Conclusion

In this study, we developed INSPECTOR, a compact and high-throughput imaging device, for quality control in long-term stem cell culture and differentiation. By quantifying images and movies captured by INSPECTOR without the need for staining, we successfully obtained effective metrics, even from low-resolution images derived from its TFT image sensors. We demonstrated these capabilities by monitoring hiPSC culture, mesodermal differentiation, and cardiomyocyte production. In the future, we expect that incorporating cutting-edge image and movie analysis, such as deep learning, will enable the provision of effective

metrics for various other types of differentiation induction processes.^{9,10,39,75} We believe this research will significantly contribute to advancements in regenerative medicine and drug discovery.

Data availability

A set of imaging data for this article is available at SSBD: repository at <https://doi.org/10.24631/ssbd.repos.2024.06.364>.

Author contributions

Conceptualisation: T. Nagai, T. Natsume, and T. Kakizuka. Device design: T. Natsume. Sample preparation: T. Kakizuka. Data acquisition: T. Kakizuka. Data analysis and visualisation: T. Kakizuka. Manuscript writing – original draft: T. Kakizuka. Manuscript writing – review & editing: T. Kakizuka, T. Nagai, and T. Natsume.

Conflicts of interest

There are no conflicts to declare.

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