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Light-Sheet Imaging to Elucidate Cardiovascular Injury and Repair

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Abstract

Purpose of review—Real-time 3-dimensional (3-D) imaging of cardiovascular injury and regeneration remains challenging. We introduced a multi-scale imaging strategy that uses light-sheet illumination to enable applications of cardiovascular injury and repair in models ranging from zebrafish to rodent hearts.

Recent findings—Light-sheet imaging enables rapid data acquisition with high spatiotemporal resolution and with minimal photo-bleaching or photo-toxicity. We demonstrated the capacity of this novel light-sheet approach for scanning a region of interest with specific fluorescence contrast, thereby providing axial and temporal resolution at the cellular level without stitching image columns or pivoting illumination beams during one-time imaging. This cutting-edge imaging technique allows for elucidating the differentiation of stem cells in cardiac regeneration, providing an entry point to discover novel micro-circulation phenomenon with clinical significance for injury and repair.

Summary—These findings demonstrate the multi-scale applications of this novel light-sheet imaging strategy to advance research in cardiovascular development and regeneration.

Keywords

light-sheet imaging; cardiovascular injury; regeneration; doxorubicin

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Compliance with Ethical Standards

Conflict of Interest

Yichen Ding, Juhyun Lee, Jeffrey J. Hsu, Chih-Chiang Chang, Kyung In Baek, Sara Ranjbarvazirj, Reza Ardehali, René R. Sevag Packard, and Tzung K. Hsiai declare no conflict of interest.

Human and Animal Rights and Informed Consent

This article does not contain any studies with human subjects performed by any of the authors. All animal rights have been approved by AAALAC and USDA.

Introduction

Real-time 3-dimensional (3-D) imaging of cardiovascular injury and repair remains a significant challenge. Conventional microscopes are confined to low light penetration and a small working distance for optical sectioning and parallelization that limit long-term biological interrogation (1–3). In addition, mechanical slicing likely distorts the intrinsic tissue integrity, and under-sampling occurs following 3-D reconstruction (4). Despite the advent of 3-D live imaging techniques such as positron emission tomography (PET) (5, 6), computed tomography (CT) (7, 8), and magnetic resonance image (MRI) (9, 10), these techniques are limited by spatial resolution (11–15). In addition, they are unable to capture fluorescently-labelled molecules that are used frequently in biomedical research. For these reasons, we hereby introduce a multi-scale and 3-D approach to imaging organisms ranging from live zebrafish embryos ($\sim 0.4 \times 0.5 \times 0.6 \text{ mm}^3$) to mouse hearts ($\sim 2 \times 2 \times 4 \text{ mm}^3$). This technique, called light-sheet fluorescence microscopy (LSFM) (16–20), has demonstrated high spatiotemporal resolution and specific labeling without needing to stitch image columns (21–23) or pivot the illumination beams and has minimal photo-bleaching and rapid acquisition time (24).

While fluorescence microscopy technologies in general have become increasingly powerful in terms of resolution, speed, and penetration, they are most often performed in thin and transparent samples (25, 26). The size and opacity of entire embryos, which are often a few millimeters in size, render it challenging to achieve single-cell resolution of $\sim 10 \mu\text{m}$ (several hundred microns deep inside the intact embryos). Current imaging techniques such as confocal microscopy are limited by their intrinsic depth penetration and z-axis resolution. Images acquired with the conventional microscopy techniques incur 1) significant background noise due to out-of-focus illumination, and 2) low axial resolution due to a large depth of field. The dynamic range and axial penetration depth are also reduced due to $\sim 95\%$ of fluorescent blockage (4). Unlike the confocal and wide-field microscopy, LSFM has the capacity to localize the 4-D (3-D spatial + 1-D time or spectra) cellular phenomena with multi-fluorescence channels (27–31). Due to the intrinsic properties of the current optical microscopy design, depth-of-focus and resolution offset each other. Furthermore, optical sectioning for dynamic sample remains a challenging issue. Initially developed to image *Caenorhabditis elegans* (32, 33), zebrafish embryos (34, 35), and *Drosophila* (36, 37), LSFM has now been adapted to image rodent organs such as the mouse hippocampus (38–40) and mouse cochlea (41) with the introduction of optical clearing techniques (42). Thus, LSFM system has allowed for uncovering both mechanical and structural phenotypes at the cellular level (16–18, 20). The Comparative advantages and disadvantages among different optical imaging modalities are listed in Table 1. In this review, we will introduce the basic optical principle of LSFM. Next, we will provide three areas of applications: 1) analysis of vascular injury and regeneration in zebrafish embryos, 2) quantification of doxorubicin chemotherapy-induced cardiac injury and repair, and 3) 3-D localization of cardiac progenitor lineage in neonatal mouse hearts.

1. Operating principle of light-sheet system and sample preparation

The custom-made light-sheet microscope (Figure 1a–c) utilized a continuous-wave laser (Laserglow Technologies, Canada) as the illumination source. The detection module was composed of the scientific CMOS (sCMOS, ORCA-Flash4.0, Hamamatsu, Japan) and a set of filters (Semrock, New York, USA). This detection module was perpendicularly installed to the illumination plane. The sample holder was oriented by a five-axis mounting stage for scanning the biological specimen. Both illumination and detection modules were controlled by a computer with a dedicated solid-state drive and the Redundant Array of Independent Disks level 0 (RAID 0) storage for fast data streaming. The lateral resolution of the LSFM is mainly governed by the numerical aperture (NA) of the objective lens, as denoted by $d_{lateral}=0.61\lambda/NA$, where λ indicates the wavelength of excitation light. The axial resolution is determined by the waist of Gaussian beam and detection NA. The waist ω_0 and Rayleigh range Z are used to define the light-sheet (Figure 1d): $\omega_0=\lambda f/\pi\omega$, $Z=\lambda f^2/\pi\omega^2$, where f is the focal length of the excitation objective, and ω denotes the half of the width of illumination beam before focusing. Generally, both waist ω_0 and Rayleigh range Z drop as the ω increases.

For the imaging data presented here, the detection objective was imaged through the liquid-air interface which introduced a spherical aberration-based point spread function (PSF) extension (43). Each image was acquired within a 10–50 ms exposure time. The stepping size of mechanical scanning was 1–5 μm , less than one half of the light sheet thickness in accordance with the Nyquist-Shannon sampling theorem. Thus, the resolving power of the LSFM in cross-section varied from 1 μm to 10 μm , while the waist ω_0 ranged from 2 to 9 μm . All of the raw data were processed to remove stationary noise. The simplified tissue clearing method was used to process the mouse hearts as described in (44).

2. Light-sheet imaging of vascular injury and regeneration in zebrafish embryos

We studied zebrafish blood cell migration in response to vascular injury and regeneration by using the transgenic *Tg(fli1:GFP; gata1:DsRed)* and *Tg(fli1:GFP; cmc12:mCherry)* lines, in which we were able to track blood cells in the vasculature in response to tail amputation at 3 days post fertilization (dpf). Using the transgenic *Tg(fli1:GFP;gata1:DsRed)* line in which the *fli1* promoter drives expression of GFP in all vasculature throughout embryogenesis and *gata1* promoter drives the expression of DsRed in blood cells (Figure 2a), we applied dual-channel detection to simultaneously track 1) the vascular loop connection between the dorsal aorta (DA) and the dorsal longitudinal anastomotic vessel (DLAV) at 3 dpf, and 2) the blood cells circulating to the injured site (Figure 2b). At 100 fps, the measured angle and displacement allowed computation of the velocity of the targeted cells as $2.9 \pm 0.1 \mu\text{m}/\text{ms}$ (Figure 2c). This method may allow studies of non-linear shear rates in a low Reynolds number system ($\text{Re}: 100 \sim 1000$). Genetic manipulation of the ADAM10 inhibitor (which inhibits proteolytic cleavage of the Notch extracellular domain) also enabled us to elucidate Notch-mediated vascular regeneration to connect the loop between DLAV with DA (Figure 2d vs. 2b). Thus, LSFM enables the discovery of novel micro-circulation phenomenon with clinical significance for injury and repair.

3. Light-sheet imaging to analyze doxorubicin chemotherapy-induced cardiac injury and repair

In adult zebrafish, the regenerating myocardium electrically couples with uninjured myocardium (45) and represents an evolutionarily-conserved model of cardiomyopathy (46). However, the small size and the two-chambered zebrafish heart limits precise morphologic assessment. In the setting of chemical clearing to achieve tissue transparency and laser light transmission, advances of light-sheet imaging coupled with automated image segmentation based on histogram analysis led to rapid and robust 3-D cardiac reconstruction, thereby unraveling the architecture of doxorubicin chemotherapy-induced cardiac injury and regeneration in adult zebrafish (47). By combining light-sheet imaging with automated segmentation, we observed dynamic changes in cardiac volumes in response to doxorubicin-induced cardiac toxicity. Precise 3-D reconstruction further enabled quantitation of cardiac volumes at days 3, 30, and 60 following chemotherapy treatment (Figure 3a). Compared to control fish, doxorubicin treatment led to an acute decrease in myocardial and endocardial volumes at day 3 ($P < 0.01$), demonstrating global cardiac injury (Figure 3b). This was followed by ventricular remodeling at day 30 ($P < 0.01$) and complete regeneration and restoration of normal architecture at day 60. Our results demonstrate the suitability of light-sheet imaging combined with automated segmentation as a high-throughput method to monitor 3-D cardiac ultrastructural changes in adult zebrafish, with translational implications to drug discovery and modifiers of chemotherapy-induced cardiomyopathy.

4. Light-sheet to localize cardiac progenitor lineage in neonatal mouse hearts

Accumulating evidence supports the ability of the mammalian heart to generate new cardiomyocytes during development and in response to injury (48, 49). However, 3-D distribution of cardiac progenitor cells to cardiac tissue formation remain poorly understood for both cardiac morphogenesis and regeneration (48, 49). We applied two reporter systems to demonstrate LSFM-imaged cardiac progenitor lineage. A rainbow multi-color reporter system (50) was used to retrospectively identify the source of new cardiomyocytes in a mouse heart at postnatal day 1 (P1) (Figure 4). Cre-mediated recombination of paired lox P sites in the $\alpha MHC^{Cre}; R2\delta^{VT2/GK}$ mouse model resulted in expression of all four fluorescent proteins (Cerulean, GFP, mOrange and mCherry, Figure 4b–c). We applied a sub-voxel light-sheet imaging method to precisely localize and subsequently quantify the numbers of cells and size of the individual clones (Figure 4d). Another fetal mouse model (embryonic day 16.5, E16.5) was $Mesp1^{Cre/+}$ crossed with $Rosa26^{tdT/+}$ reporter system to study the contribution of tdT+ cell (red hot) in an intact fetal mouse heart (Figure 4e–h). 3-D LSFM imaging showed the majority of the cardiac cells that were labeled with tdT arose from a mesoderm posterior1 homolog ($Mesp1$) origin and allowed tracing of their 3-D distribution, proliferation, and tissue formation. Thus, light-sheet imaging allows for clonal analysis of cardiomyocyte generation during development and injury.

Conclusions and Discussion

We have established a holistic strategy for multi-scale applications of LSFM to elucidate cardiovascular injury and repair in both zebrafish and mouse models. This method provides an efficient and robust platform for detailed analysis of cardiovascular phenomena ranging

from tissue clearing *ex vivo* to 4-D imaging *in vivo*. This framework builds on the high axial and temporal resolution provided by LSM, which allows for long-term, 3-D and 4-D visualization of *in vivo* cellular events and organogenesis with minimal photo-bleaching or photo-toxicity.

Light-sheet imaging allows one to image the entire vasculature and cardiac architecture without the need for mechanical slicing or stitching of image columns. By this means, LSM bypasses the need to move the tissue volume or light-sheet along the propagation of the illumination, thereby allowing us to carry out multi-scale imaging in a wide range of specimens, from embryos to adults, with the light-sheet thickness ranging from 1 μm to 18 μm (18, 20). Our methodology expands the field of view from hundreds of micrometers to tens of millimeters to cover the entire mouse heart with sufficient spatiotemporal resolution to track cardiovascular development in rodents. Our imaging strategy simplifies the operation during image acquisition, reduces the complexity of pre- and post-processing of cardiac images, and enhances the robustness of the imaging system to provide dual-channel *in vivo* imaging at over 100 fps. This strategy brings advanced imaging to studies of tissue injury and regeneration with multi-scale applications to fundamental studies of cardiovascular development and translational work in congenital cardiac anomalies and disease.

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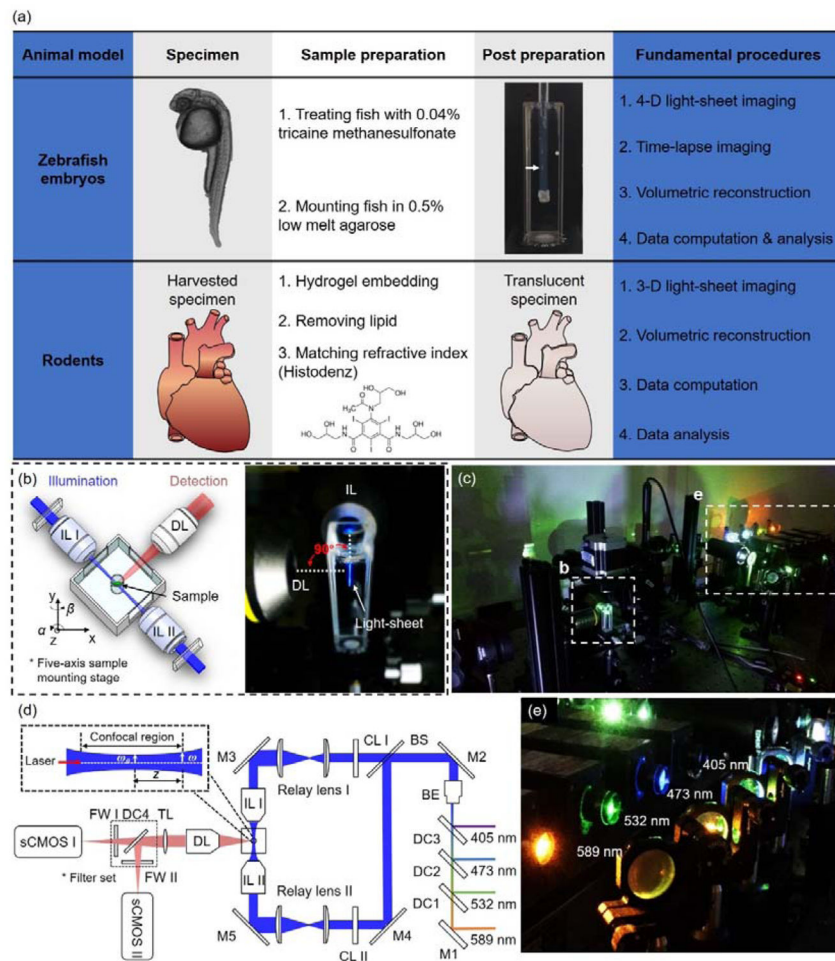


Figure 1.

Fundamental concept of the light-sheet imaging strategy. (a) The critical procedures of the multi-scale imaging are indicated for both embryonic zebrafish and mouse studies. (b) The specimen is mounted at the intersection of the illumination lens (IL) with the detection lens (DL). The laser light-sheet is excited from the IL in a 2-D plane which is orthogonal to the detection axis. The LSFM system provides a long working distance with air objective lenses in comparison to water-dipping lenses in conventional light-sheet systems. (c–d) A photo and a schematic illustrate the layout of the light-sheeting imaging system. A cylindrical lens (CL) converts the laser beam to a sheet of laser light that can transversely illuminate a thin layer of the sample. The illuminated 2-D thin layer (fluorescent detection in red) is captured by the high-frame rate sCMOS camera. The filter wheels (FW I and II) in front of sCMOS cameras are used for multi-color acquisitions. (e) A photo depicts an array of laser beams aligned for multi-channel fluorescent detection. M: mirror; BS: beam splitter; BE: beam expander; TL: tube lens; DC: dichroic mirror; FW: filter wheel.

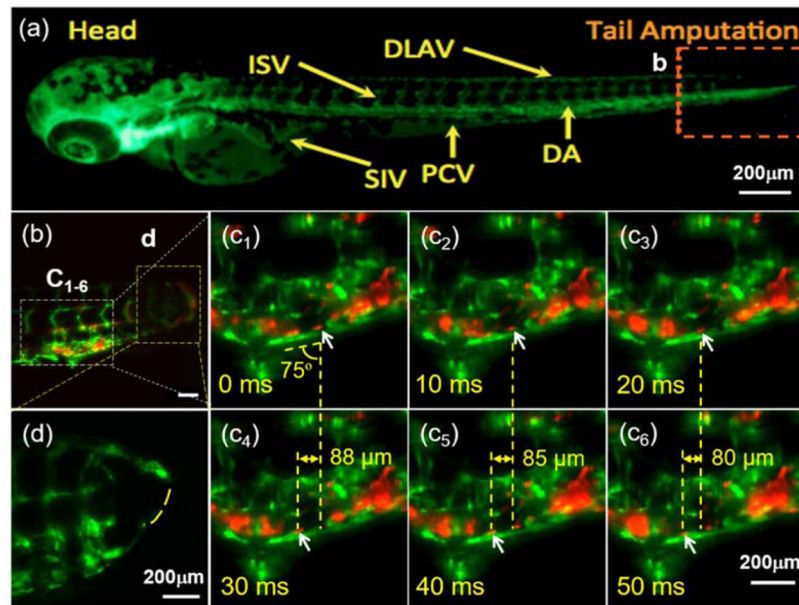


Figure 2.

Light-sheet microscopic illumination of vascular regeneration and circulating blood cells in response to tail amputation. (a) An inverted fluorescence image of a transgenic *Tg(fli1:GFP:gata1:DsRed)* zebrafish embryo showing the vasculature (green) at 3 dpf. ISV: intersegmental vessel; DLAV: dorsal longitudinal anastomotic vessel; SIV: subintestinal vessel; PCV: posterior cardinal vein; DA: dorsal aorta. Box b indicates the site of tail amputation. (b) LSFM captures blood cells (red) proximal to the site of amputation and regeneration. Dashed yellow boxes indicate locations of higher power images in the subsequent panels (c_{1-6}). Arrows indicate the position of an individual RBC (red) in relation to the vascular endothelial layer (green) acquired by LSFM at 100 fps. The average angle between the vein (green PCV) and vertical axis of the frame is 75° , and the relative displacement along the horizontal axis of the frame in each 30 ms period is $88\ \mu\text{m}$ (c_4 - c_1), $85\ \mu\text{m}$ (c_5 - c_2) and $80\ \mu\text{m}$ (c_6 - c_3), respectively. These measures correspond to a net velocity of $2.9 \pm 0.1\ \mu\text{m}/\text{ms}$ for that blood cell of interest. (d) The dashed line indicates the incomplete vascular regeneration between DLAV and DA in a separate zebrafish embryo treated with an inhibitor of ADAM10 (GI254023X, Sigma) which blocks Notch signaling. Scale bars: $200\ \mu\text{m}$.

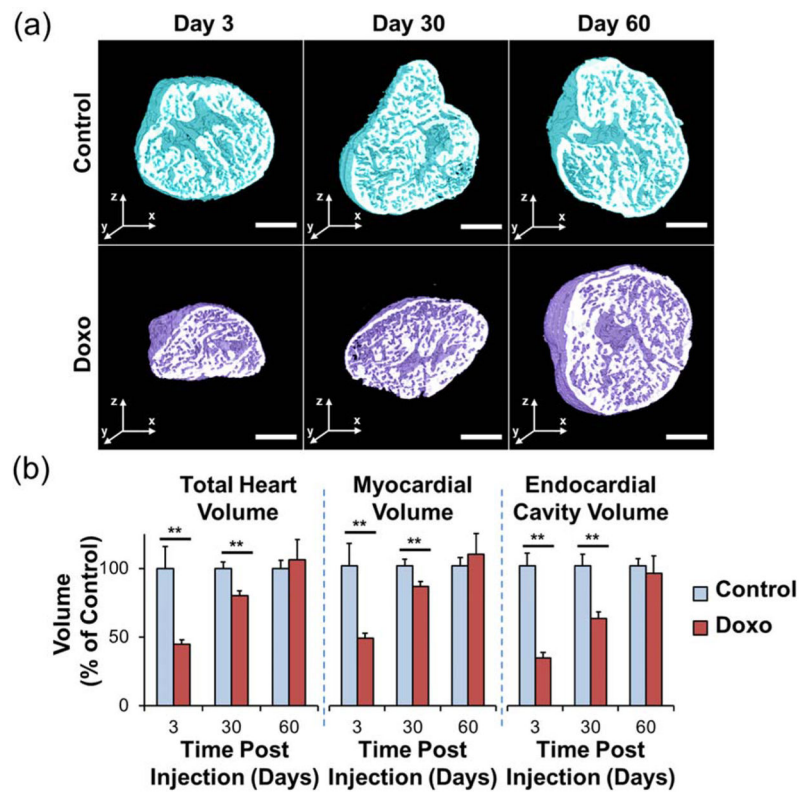


Figure 3. Cardiac architecture following doxorubicin treatment. Following intraperitoneal treatment with doxorubicin or control vehicle, adult zebrafish hearts were harvested at days 3, 30, and 60. (a) Throughout the duration of the study, control hearts exhibited a preserved architecture. In contrast, doxorubicin-treated hearts demonstrated a profound cardiac remodeling leading to acute decrease in size at day 3, followed by gradual increase at day 30, and normalization at day 60. (b) Cardiac architecture characterization by quantitative analysis of the total heart, myocardial, and endocardial volumes compared to control values demonstrating the cardiac repair process following response to chemotherapy-induced injury. Legend. ** $P < 0.01$. Doxo: doxorubicin. Scale bar: 200 μm . (Reproduced with permission from: Packard et al. Sci Rep. 2017;7) (47).

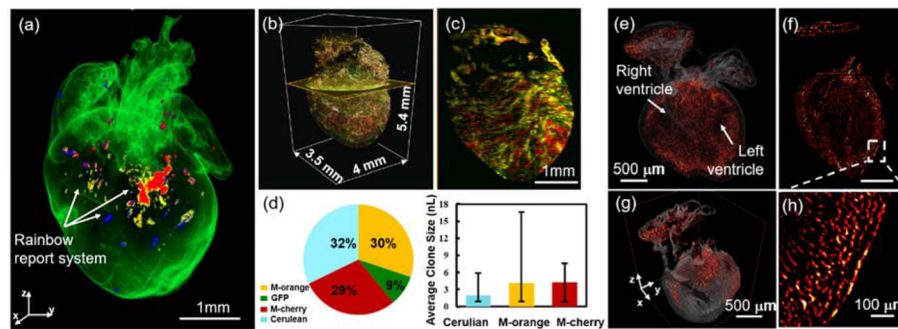


Figure 4.

3-D tracking of the cardiac progenitor lineage in the fetal and neonatal mouse hearts. (a) Expression of Cre in the neonatal mouse heart induces random recombination between mutated paired lox P sites leading to expression of Cerulean, GFP, mOrange and mCherry respectively. (b–c) Sub-voxel imaging of the rainbow heart was captured in four channels. Multi-view reconstruction was performed to enhance the spatial resolution for tracking and localizing the differentiation of cardiomyocytes in the intact neonatal heart. (d) Quantification of the number of labeled cells is illustrated in the pie chart, and the sizes of individual clones in a heart by the bar graphs. (e–f) Spatial distribution of tdT+ cells (red hot) in a fetal mouse heart which was *Mesp1*^{Cre/+} crossed with *Rosa26*^{tdT/+} reporter. (g) A 3-D orthogonal slice and (h) 2-D inset reveal the contribution of tdT+ cells in the heart. Scale bars: (a & c) 1 mm; (e–g) 500 μm; (h) 100 μm.

Table 1

Comparative advantages and disadvantages among different imaging modalities

| Method | Resolution (Lateral-Axial, nm) | Penetration (μm) | Acquisition time per image | Advantage | Disadvantage |
|-------------------|--------------------------------|-------------------------------|----------------------------|---|---|
| LSFM | 200–500 | >1000 | ms-s | <ul style="list-style-type: none"> • Low phototoxicity • Isotropic resolution | <ul style="list-style-type: none"> • Sample preparation • Big data |
| WFM | 250 | --- | ms-min | <ul style="list-style-type: none"> • Low cost • Easily built | <ul style="list-style-type: none"> • Low contrast • Out of focus blur |
| CFM | 200–400 | 150 | s-min | <ul style="list-style-type: none"> • Optical sectioning | <ul style="list-style-type: none"> • Scanning |
| SD-CFM | 200–400 | 150 | ms-min | <ul style="list-style-type: none"> • Fast speed | <ul style="list-style-type: none"> • Fixed pinhole |
| MPM | 300–500 | 1000 | s-min | <ul style="list-style-type: none"> • Penetration depth • Inherent confocality | <ul style="list-style-type: none"> • High power pulsed laser |
| STED | 80–400 | 50 | min | <ul style="list-style-type: none"> • Super-resolution at confocal speed | <ul style="list-style-type: none"> • Intense depletion laser • Photobleaching |
| PALM/STORM | 50–100 | 0.1 | min-h | <ul style="list-style-type: none"> • Extremely high resolution | <ul style="list-style-type: none"> • Slow speed • Pointillism artifacts |
| FRET | 4–10 | --- | ms-s | <ul style="list-style-type: none"> • Molecular interactions | <ul style="list-style-type: none"> • Sensitive to <i>pH</i> |

Legend: LSFM: light-sheet fluorescent microscopy; WFM: wide-field microscopy; CFM: confocal microscopy; SD-CFM: spinning disk confocal microscopy; MPM: multi-photon microscopy; STED: stimulated emission depletion; PALM: photoactivated localization microscopy; STORM: stochastic optical reconstruction microscopy; FRET: Förster resonance energy transfer. Adapted from [2].