Light sheet theta microscopy for rapid high-resolution imaging of large biological samples

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Abstract

Background: Advances in tissue clearing and molecular labeling methods are enabling unprecedented optical access to large intact biological systems. These developments fuel the need for high-speed microscopy approaches to image large samples quantitatively and at high resolution. While light sheet microscopy (LSM), with its high planar imaging speed and low photo-bleaching, can be effective, scaling up to larger imaging volumes has been hindered by the use of orthogonal light sheet illumination.

Results: To address this fundamental limitation, we have developed light sheet theta microscopy (LSTM), which uniformly illuminates samples from the same side as the detection objective, thereby eliminating limits on lateral dimensions without sacrificing the imaging resolution, depth, and speed. We present a detailed characterization of LSTM, and demonstrate its complementary advantages over LSM for rapid high-resolution quantitative imaging of large intact samples with high uniform quality.

Conclusions: The reported LSTM approach is a significant step for the rapid high-resolution quantitative mapping of the structure and function of very large biological systems, such as a clarified thick coronal slab of human brain and uniformly expanded tissues, and also for rapid volumetric calcium imaging of highly motile animals, such as Hydra, undergoing non-isomorphic body shape changes.

Keywords: Light sheet microscopy, Whole brain imaging, Quantitative imaging, Hydra, Calcium imaging, Tissue clearing, Expansion microscopy
still entail highly redundant illumination of out-of-focus parts of the samples and also reduced axial resolution and imaging depth, thus limiting their utility for high-resolution imaging of large cleared samples such as whole mouse brains (Fig. 1a). On the other hand, light sheet microscopy (LSM)-based approaches, with their orthogonal single-plane illumination and simultaneous whole-plane detection, are proving to be highly effective due to minimal photo-bleaching and high imaging speeds (2–3 orders of magnitude more than confocal) [11–13]. However, these advantages of orthogonal illumination-detection geometry also require unhindered optical access from the

![Diagram of light sheet microscopy](image)

**Fig. 1** Light sheet theta microscopy (LSTM) for high-resolution quantitative imaging of large intact samples. a Light sheet microscopy (LSM) employs orthogonally illumination-detection optics, which limits the lateral dimensions of imaging volume. iSPIM, SCAPE/OPM and line scan confocal microscopy are partially effective in alleviating this limitation, however at the cost of reduction in usable working distance (magenta arrowheads) and image quality (e.g., SCAPE collects low-quality signal from non-native focal planes, and line scan confocal results in lower axial resolution and high photo-bleaching.). The proposed LSTM uses non-orthogonal (< 90°) illumination light sheets to effectively image very large samples, while maintaining high imaging speed and depth and uniform high resolution. b One or two light sheets intersect with the detection plane in a line illumination profile, which is synchronously scanned with the rolling shutter detection of an sCMOS camera to achieve optical sectioning. c Two scanning approaches: 1-axis scanning (1-AS) by perpendicular translation and simultaneous 2-axis scanning (2-AS, default LSTM) by translation along and perpendicular to the illumination axis such that the thinnest part is utilized for uniform planar illumination. d Comparison of point spread function (PSF) in 1-AS, default LSTM, and LSM configurations. Left: x-z maximum intensity projections of ~1 μm fluorescent microbeads imaged using the same detection (10x/0.6NA/8mmWD) and illumination (4x/0.28NA/28.5 mmWD) objectives. Axial full width at half maximum values (FWHM) across the field of view (blue LSTM in default 2-AS mode, green LSTM in 1-AS mode, red LSM). LSTM achieves uniform axial resolution (~4–6 μm FWHM) over the entire field of view, whereas both the 1-AS and LSM provide lower peripheral resolution (1-AS ~ 5–13 μm; LSM ~ 4–11 μm). Right: x-z projections (20 μm) of an image volume from a DAPI-stained human brain tissue. Additional file 5: Video 2 [29] provides 3D reconstructions. The graph compares the signal for a central and a peripheral region of interest. Scale bars: 100 μm
sides of the samples, thus limiting the lateral dimensions (along the illumination light sheet) of the imaging volumes (Fig. 1a). For example, we previously reported an optimized implementation of LSM, called CLARITY optimized light sheet microscopy (COLM) [5, 14], which allowed high-resolution imaging of entire intact mouse brains in a few hours imaging time, although with progressively reduced image quality towards the middle of the samples [5] due to the scattering of illumination light sheets. Similar attempts of LSM imaging of clarified rat brains resulted in much poorer image quality in large parts of the brain [15].

Alternative optical configurations of LSM have been explored to address these limitations, including the rotation of the illumination and the detection axes by 45° relative to the sample surface normal in objective-coupled planar illumination (OCPI) microscopy, inverted selective plane illumination microscopy (iSPIM), dual view iSPIM (diSPIM), and triple-view implementations [16–21], and the generation of illumination light sheets through the detection objectives themselves in swept confocally aligned planar excitation (SCAPE)/oblique plane microscopy (OPM) [22, 23]. The iSPIM/diSPIM approach does alleviate the limits on the lateral dimensions of imaging volumes, although at the cost of significant reduction in the usable working distance of the detection objective (Fig. 1a); therefore, it remains restricted to relatively low-numerical aperture (NA)/long-working distance (WD) detection objectives for imaging of large samples. The triple-view approach [21] incorporated an additional objective in the diSPIM implementation for simultaneously detecting the obliquely illuminated plane from the opposite side by rapid scanning with the piezo motors, resulting in enhanced spatial resolution for small samples (such as single cells). The SCAPE/OPM implementations use rotation optics to image an oblique plane illuminated using an oblique light sheet generated through the detection objective itself (Fig. 1a). The use of a single objective for detection as well as illumination is effective for fast volumetric imaging of small samples such as developing embryos. However, the imaging of an oblique (relative to the native detection focal plane) plane provides less-than-optimal image uniformity (across the imaged plane) and resolution. In summary, the scaling up of LSM imaging volumes, while maintaining uniform high imaging quality and speed, faces steep challenges. Here we address some of these challenges by developing a conceptually distinct microscopy framework, termed light sheet theta microscopy (LSTM), which builds upon the principles of LSM to allow high-speed quantitative imaging of large intact tissues with uniform high resolution. The LSTM uses two symmetrically arranged oblique light sheets, generated using independent illumination objectives, for rapid high-resolution imaging of large samples (Fig. 1). The oblique optical arrangement eliminates the restrictions on the sample lateral dimensions while ensuring high imaging speed and resolution and utilization of the entire available WD of high-NA detection objectives (Fig. 1a). We present a detailed characterization of the LSTM approach and demonstrate several real-world high-resolution imaging examples of very large samples including mouse and rat brain tissues, a large section of human brain, and a highly expanded ExM sample. The new capabilities of LSTM for high-speed quantitative imaging of larger samples at high resolution with low photo-bleaching may facilitate mapping of an entire post-mortem human brain (thick slab-by-slab) in a practical time-frame.

Results
LSTM implementation
LSTM includes a standard wide-field detection arm and two symmetrically arranged non-orthogonal (θ < 90°, relative to the detection axis) illumination arms for the generation of thin static light sheets that intersect at the detection focal plane (Fig. 1). The resulting thin line illumination profile is scanned along the detection focal plane in synchrony with the row-by-row rolling shutter imaging with a scientific complementary metal-oxide-semiconductor (sCMOS) camera (virtual slit effect [5]) to achieve thin optical sectioning (Fig. 1b). This is realized by simultaneously translating the light sheet along (using an electrically tunable lens (ETL)) and perpendicular (using a galvanometer (galvo) scanner) to its propagation direction such that only the thinnest part intersects the detection plane (Fig. 1c). Note that the translation of light sheets along their propagation direction may also be achieved using alternative implementations, including fast piezo motors for translating illumination objectives, or using an acoustic tunable lens such as the tunable acoustic gradient index of refraction (TAG) lens (TAG optics) [14, 24–28].

The LSTM illumination and detection arms were implemented as rigid assemblies (using a caging system from Thorlabs) which were connected to a vertically mounted breadboard via x-y manual translation stages for finer adjustments (Fig. 2, Additional file 1: Figure S1, Additional file 2: Figure S2, Additional file 3: Figure S3, Additional file 4: Video 1). An open-top sample mounting strategy was implemented by using a custom 3D-printed sample chamber (Fig. 2, Additional file 1: Figure S1) attached to a high-accuracy x-y-z motorized stage assembly (LNR50S, Thorlabs). The imaging samples can be mounted in a quartz glass cuvette of appropriate size. The illumination arm consists of a laser source (SOLE-6, 405, 488, 561, and 647 nm, Omicron-Laserage), a
Fig. 2 LSTM microscopy implementation. a LSTM optical path. Two symmetric light sheets are generated by using a cylindrical lens (CL), scan lens (SL), tube lens (TL), and illumination objectives. The galvo scanners are used to translate the light sheets perpendicular to their propagation direction, and the electrically tunable lens (ETL) for translating the thinnest part of the light sheets along the propagation direction. An input beam of ~10 mm diameter is then trimmed through an iris. A slit is placed after the ETL to control the effective numerical aperture of the illumination. An additional iris is placed between the SL and TL to control the light sheet width. The illumination axes are arranged at ~60° to the detection axis. A custom 3D-printed cap with a quartz coverslip is attached to the illumination objective to allow dipping in the immersion oil to ensure that the low NA illumination rays from an air illumination objective enter perpendicularly to the oil. The detection arm consists of a detection objective (Olympus 10x/0.6NA/8mmWD or 25x/1.0NA/8mmWD, both with correction collars), a tube lens, and an sCMOS camera. b 3D model of the LSTM microscope. A vertical breadboard was used to mount the caged optical assemblies via xy manual translation stages to allow fine adjustments. A sample chamber was attached to a 3-axis (x, y, z) motorized stage assembly. See also Additional files 1, 2, 3, and Additional file 4: Video 1 for further details and Table 1 for complete parts list.
collimator (~10-mm beam diameter, Omicron-Laserage), an ETL (Optotune), a cylindrical lens (LJ1695RM-A, f = 50 mm, Thorlabs), a galvo scanner (GVS001, Thorlabs), a scan lens (CLS-SL, Thorlabs), a tube lens (f = 200 mm, ITL200, Thorlabs), and an illumination objective (Macro 4×/0.28NA/28.5mmWD, Olympus; the arrangement of different components is summarized in Additional file 2: Figure S2, and the parts list is provided in Table 1). Note that even though the ETLs are mounted at an oblique angle (as opposed to vertical), this does not result in any significant observable aberrations because of the low NA of the illumination objectives. In addition, an iris is placed after the collimator to remove peripheral spreads of Gaussian beams, a one-dimensional (1D) slit is positioned before the cylindrical lens to control the effective NA (hence the light sheet thickness), and a second iris is placed at the conjugate plane between the scan lens and tube lens to control the light sheet width (i.e., the dimension of light sheet perpendicular to its propagation direction). The detection arm is composed of a detection objective (Olympus 10×/0.6NA/8mmWD or 25×/1.0NA/8mmWD, both with refractive index correction collars, from water to oil), a multi-band emission filter (FF01-432/515/595/730-50-D, Semrock), a tube lens (f = 200 mm, ITL200, Thorlabs), and an sCMOS camera (Orca Flash 4.0, Hamamatsu; 2048 × 2048 pixels, 6.5 μm × 6.5 μm pixel size). Note that since we used a 200-mm tube lens with Olympus objectives, the effective magnifications are 11.11 and 27.78 for the 10× and 25× objectives respectively. The LSTM assembly was optically aligned by placing a prism mirror (with scratches in the center; see Additional file 1: Figure S1 for mounting arrangements) in the focal plane of the detection optics, to visualize the location and cross section of the light sheet relative to the detection focal plane. The light sheet positioning parameters (i.e., galvo and ETL) were optimized such that the thinnest part was in alignment with the center of the field of view of the detection plane. Next, fluorescent beads, embedded in a high concentration (> 2%, to restrict the signal source to the gel surface) agarose, were used to find the optimal galvo and ETL parameters by examining the extent and focus quality. All imaging experiments were performed using 405 (for 4′-6-diamidino-2-phenylindole (DAPI)) or 488 nm (for eYFP detection) laser lines. LSTM parameters were adjusted to use a 2 to 5 μm effective light sheet thickness.

The angular separation of the illumination and detection arms is constrained by the physical dimensions and optical properties of the detection and illumination objectives. For instance, only an angular separation range of 43–62° is feasible for the specific combination of illumination (Macro 4×/0.28NA/29.5mmWD, Olympus) and detection objectives (10×/0.6NA/8mmWD or 25×/1.0NA/8mmWD, Olympus) used in this study (see Fig. 3a, Additional file 3: Figure S3). Note that the WD of the illumination objective is specified for use in air; therefore, an approximate effective WD in immersion oil was calculated as shown in Additional file 3: Figure S3.

**LSTM characterization**

We first characterized the LSTM point spread function (PSF) by imaging micrometer-size fluorescent microbeads. The same microbeads were also imaged with LSM as well as a non-optimal 1-axis LSTM scanning procedure (i.e., the light sheet is only translated perpendicular to its propagation; marked as 1-AS mode in Fig. 1c), using the exact same detection (10×/0.6NA) and illumination objectives (4×/0.28NA). The quantification and comparison of full width at half maximum (FWHM, Fig. 1d) revealed that LSTM indeed allows for uniform high axial resolution (~4 to 6 μm for the combination of these objectives) across the entire field of view, whereas as expected, both the LSM and the non-optimal 1-AS LSTM scan procedure resulted in lower axial resolution on the peripheries of the field of view (>11 μm) (Fig. 1d, Fig. 4 and Additional file 5: Video 2 [29]). Next, we compared the maximum illumination path length (MIPL), i.e., the maximum distance the illumination light sheets need to penetrate inside the tissue to achieve complete sample coverage. The shorter the illumination path length, the lesser the scattering, hence potentially better the imaging performance. In LSM, the MIPL depends on the sample width as the light sheet needs to penetrate the entire width (half width for two-sided illumination) of the sample to achieve complete coverage, whereas in LSTM, the MIPL depends on the angular separation of the illumination-detection arms and the tissue thickness (t): t/cos(θ) (Fig. 3b). The ratio of LSTM and LSM illumination path lengths was calculated for varying sample dimensions and the θ values (Fig. 3b shows θ = 0°, 10°, 20°, 30°, 40°, 50°, 60°, 70°, and 80°). This revealed that the LSTM illumination path length was smaller than that for LSM for wider samples, and larger for smaller samples, suggesting the complementary advantages of LSTM and LSM for high-resolution imaging of large and small samples respectively. Since the LSTM illumination path length decreases with decreasing angular separation (t/cos(θ)), minimizing the angular separation (θ) will result in potentially higher imaging quality; however, the angular separation also affects the effective light sheet thickness (approximated as b/sin(θ), Fig. 3c) in an inverse relationship. Therefore, we used the maximum allowed angular separation (~60°) to achieve better axial resolution.

Further, we sought to assess and compare the total energy loads in LSTM vs. LSM imaging (Fig. 3d–f, Additional file 7: Figure S5). Since LSTM employs only the thinnest part of the illumination light sheet for imaging, this results in significant redundancy in
### Table 1  Parts list of LSTM

| Vendor       | Number | Qty. | Description                                                                 |
|--------------|--------|------|-----------------------------------------------------------------------------|
| Detection unit |        |      |                                                                             |
| Thorlabs     | CX2    | 1    | 60-mm Cage System Translating Lens Mount for Ø2" Optics                   |
| Thorlabs     | LCP90F | 1    | 60-mm Removable Cage Plate                                                 |
| Thorlabs     | SM2A20 | 1    | SM2-M38 Adapter for Nikon Tube Lens                                       |
| Thorlabs     | SM2L30 | 2    | SM2 Lens Tube, 3" Thread Depth, One Retaining Ring Included               |
| Thorlabs     | LCP09  | 2    | 60-mm Cage Plate with Ø2.2" Double Bore for SM2 Lens Tube Mounting       |
| Thorlabs     | ER10   | 4    | Cage Assembly Rod, 10" Long, Ø6 mm                                        |
| Thorlabs     | SM1A1  | 1    | Adapter with External SM05 Threads and Internal SM1 Threads               |
| Thorlabs     | SM2A31 | 1    | Adapter with External C-Mount Threads and Internal SM2 Threads            |
| Hamamatsu   | C13440 | 1    | sCMOS Orca Flash 4.0 V3.0 camera                                           |
| / Objective Adapter |        |      |                                                                             |
| Thorlabs     | SM2V10 | 1    | Ø2" Adjustable Lens Tube, 0.81" Travel                                    |
| Olympus      | XPLN10XS/340 | 1 | 10x Long Working Distance Detection Objective                           |
| Olympus      | XSLPLN92XGMP | 1 | 25x Long Working Distance Detection Objective                           |
| Thorlabs     | LCP01B | 2    | 60-mm Cage Mounting Bracket                                               |
| Thorlabs     | RSH2   | 2    | Ø1" Post Holder with Flexure Lock, Pedestal Base, L = 2"                 |
| Thorlabs     | R52    | 2    | Ø1" Pillar Post, 1/4"-20 Taps, L = 2", 8-32 Adapter Included             |
| Thorlabs     | TB80606| 2    | Large-Area Translation Stage, 6" × 7.66"                                 |
| Thorlabs     | TTL200 | 1    | f = 200 mm Tube Lenses for Wide Field Imaging                            |
| Illumination unit |        |      |                                                                             |
| Thorlabs     | SM2V10 | 2    | Ø2" Adjustable Lens Tube, 0.81" Travel                                    |
| / Objective Adapter |        |      |                                                                             |
| Olympus      | XFLPLN4X/340 | 2 | 4x Air Objective                                                        |
| Thorlabs     | SM2A20 | 2    | SM2-M38 Adapter for Nikon Tube Lens                                       |
| Thorlabs     | CX2    | 2    | 60-mm Cage System Translating Lens Mount for Ø2" Optics                   |
| Thorlabs     | SM2A31 | 2    | Adapter with External C-Mount Threads and Internal SM2 Threads            |
| Thorlabs     | SM2V10 | 2    | Ø2" Adjustable Lens Tube, 0.81" Travel                                    |
| Thorlabs     | LCP09  | 2    | 60-mm Cage Plate with Ø2.2" Double Bore for SM2 Lens Tube Mounting       |
| Thorlabs     | LCP01B | 4    | 60-mm Cage Mounting Bracket                                               |
| Thorlabs     | R52    | 4    | Ø1" Pillar Post, 1/4"-20 Taps, L = 2", 8-32 Adapter Included             |
| Thorlabs     | RSH1.5 | 4    | Ø1" Post Holder with Flexure Lock, Pedestal Base, L = 1.5"               |
| Thorlabs     | TB80606| 4    | Large-Area Translation Stage, 6" × 7.66"                                 |
| Thorlabs     | ER05   | 8    | Cage Assembly Rod, 1/2" Long, Ø6 mm                                       |
| Thorlabs     | LCP02  | 6    | 30-mm to 60-mm Cage Plate Adapter, 8-32 Tap                              |
| Thorlabs     | LLI6958M-A | 2 | Ø1", N-BK7 Mounted Plano-Convex Round Cylindrical Lens                   |
| Thorlabs     | CRM1L  | 2    | Cage Rotation Mount for Ø1" Optics, Double Bored with Setscrew, 8-32 Tap |
| Thorlabs     | CP20S  | 2    | 30-mm Cage System Iris, Ø20.0-mm Maximum Aperture                        |
| Thorlabs     | CP90F  | 2    | 30-mm Removable Cage Plate, Front and Back Plate, Internal SM1 Threading |
| Thorlabs     | CXY1   | 2    | 30-mm Cage System, XY Translating Lens Mount for Ø1" Optics             |
| Thorlabs     | CP12   | 2    | 30-mm Cage Plate, Ø1.2" Double Bore for SM1 Lens Tube Mounting          |
| Thorlabs     | LCP01  | 4    | 60-mm Cage Plate, SM2 Threads, 0.5" Thick, 8-32 Tap (Two SM2RR Retaining Rings Included) |
| Thorlabs     | CLS-SL | 2    | Scan Lens with Large Field of View, 400 to 750 nm, EFL = 70 mm           |
illumination. Similarly, in LSM, the imaging of large samples (i.e., larger than a single field of view of detection) entails redundant illumination of the parts of samples (along the illumination) not being imaged. As summarized in the Fig. 3d schematics, the total redundant energy load depends on the sample thickness in LSTM (Fig. 3d, top row) and on the sample width in LSM (Fig. 3d, bottom row). Therefore, for a quantitative comparison of LSTM vs. LSM energy loads, we calculated the ratio of the total energy loads (the procedure is summarized in Additional file 7: Figure S5) as a function of the sample width, thickness, angular separation ($\theta$) of the LSTM, and the detection objective magnification. Note that we are comparing the LSTM energy load with the scanned light sheet microscopy, which is the commonly used implementation for imaging of large samples (e.g., COLM [5]) due to the reduced coherent illumination scattering and the synchronization possibilities with the rolling shutter detection of sCMOS cameras. Therefore, the dwell time of the illumination line profile is the same for both. As summarized in Fig. 3e, the LSTM indeed imparts a much higher energy load for imaging of smaller samples, but the ratio approaches 1 with increasing sample size and higher detection magnification. The LSTM energy load also depends on the angular separation ($\theta$); $\theta = 60^\circ$ was used for these calculations. To complement these calculations, we also performed empirical assessment of the signal photo-bleaching for the LSTM imaging datasets reported in this study. As shown in Fig. 3f, no significant trend is observed, suggesting minimal photo-bleaching consequences. In summary, the LSTM total energy load is much higher for smaller samples, but is comparable to LSM for high-resolution imaging of larger samples, further supporting the overall complementary advantages of LSTM and LSM for imaging of larger and smaller samples.
**Fig. 3** LSTM characterization. 

a) Geometric constraints in LSTM. Specific example of using Olympus 4×/0.28NA/29.5WD and 10×/0.6NA/8mmWD for illumination-detection. Note that the working distance of the air illumination objective is elongated in high refractive index immersion media (Additional file 3: Figure S3). Angular separation of ~ 60° was used for all experiments. 

b) Comparison of maximum illumination path length (MIPL) required for full sample coverage in LSTM and LSM. The illumination light sheets need to penetrate the entire width \(w\) of the sample (or half width for two-sided illumination) in LSM, whereas MIPL depends on the angular arrangement and the tissue thickness \(t\) in LSTM. 

Bottom left: dependence of LSTM MIPL on \(\theta\) and sample thickness \(t\); arrow indicates increasing \(t\). Bottom right: MIPL dependence on the sample width and thickness: magenta and cyan highlight LSTM < LSM and LSTM > LSM respectively, assuming \(\theta = 60°\) (see Additional file 6: Figure S4 for full \(\theta\) range). 

c) Effective planar illumination thickness can be approximated as \(b/\sin(\theta)\), where \(b\) is the actual light sheet thickness. The right graph plots the effective light sheet thickness as a function of \(\theta\) and \(b\) (arrow points to increasing \(b\)). 

d) Comparison of redundant illumination in LSTM and LSM for imaging of a single plane (top row) and an image stack (bottom row). 

e) Ratios of total illumination energy loads (LSTM/LSM) as a function of sample width \(w\), angular configuration (\(\theta\)), sample thickness \(t\), and objective magnification (10× and 25×). Illumination energy load is higher in LSTM for smaller samples and similar to LSM for larger samples. The energy load ratio also decreases with increased angular separation (60° is marked) and the magnification of detection objective. Additional file 7: Figure S5 provides details. 

f) The average signal of tiles in the order of acquisitions. Note that no significant photo-bleaching trend is observed.
rolling shutter exposure parameter). As expected, the axial resolution is better for smaller rolling shutter widths (Fig. 4a), analogous to the effect of the pinhole diameter in confocal microscopy.

Finally, a major advantage of LSM is the high imaging speed. Similar to standard LSM implementations (e.g., COLM [5, 14]), the LSTM imaging procedure involves synchronization of the illumination line with the rolling shutter detection of the scMOS camera. Therefore, a single image acquisition takes 20 ms (which is a property of the camera imaging speed in rolling shutter mode), yielding 50 full frames (2048 x 2048) per second. The LSTM also allows the synchronization of the two illumination line profiles independently with the bi-directional readout mode of the scMOS sensors (10 ms for full frame). However, the imaging of large volumes requires the use of long travel-range motorized stages, which are typically the rate-limiting step in the entire imaging procedure. For example, as reported previously [5], a typical sample stage takes more than 50 ms for a 5-μm z-step sample motion, resulting in ~60–70 ms acquisition time per z-plane, i.e., 10–15 Hz imaging speed for LSTM as well as state-of-the-art LSM implementations for large sample imaging. Alternatively, instead of using a step-wise motion, the sample can also be continuously scanned at a small uniform speed to allow overall higher imaging speeds, although at the cost of significant shearing artifacts. We would also like to highlight that, unlike LSM (e.g., the COLM system [5]), LSTM imaging did not require any pre-calibration/adaptive parameter correction steps because of the overall smaller illumination path lengths, resulting in a higher effective (about twofold) imaging speed.

Fig. 4 LSTM optical sectioning. a x-z maximum intensity projections of an image stack acquired from human brain tissue, shown in (b), stained with DAPI. The camera rolling shutter exposure time determines the effective slit (rolling shutter) width (0.1–1 ms, i.e., 66–665 μm on the scMOS sensor and 6–60 μm on the sample. The images were acquired using two different scanning modes: LSTM 1-axis scan (1-AS) and LSTM 2-axis scan (2-AS, default). Total frame exposure was 20 ms for all the images. As evident, the 2-AS mode allows for uniform planar illumination for achieving quantitative imaging, and the axial resolution decreases with increased rolling shutter exposure. All scale bars are 100 μm. b LSTM imaging of a large thick section of cleared human brain tissue (~ 10.5 mm x 14.1 mm x 3 mm) stained with DAPI. We used 0.5-ms rolling shutter exposure settings and 20 ms for entire frame exposure to acquire this dataset. Scale bar is 1 mm.
LSTM enables rapid quantitative imaging of large samples with uniform high resolution

To assess the performance of LSTM in real-world experimental scenarios, we performed high-resolution imaging of cleared samples of various sizes and shapes. First, we imaged a large thick section of cleared DAPI-stained human brain (~ 10.5 mm × 14.1 mm × 3 mm) and a cleared intact central nervous system (11.8 mm × 27.6 mm × 5.2 mm) of a Thy1-eYFP transgenic mouse using 405 nm and 488 nm illumination wavelengths respectively. As shown in Figs. 4b and 5a and Additional file 8: Video 3 [29], LSTM enabled rapid high-resolution imaging of these large samples with uniform imaging quality throughout, even for a sample with highly curved surfaces. We further imaged a thick (~ 9.6 mm × 13.5 mm × 5.34 mm; the sample expanded ~ 1.5- to 2-fold due to the immersion in glycerol solution [14]) coronal slice of a CLARITY-cleared Thy1-eYFP transgenic mouse brain, with 10×/0.6NA/8mmWD (Fig. 5b, Additional file 9: Video 4 [29]) as well as 25×/1.0NA/8mmWD (Fig. 5c) objectives. As demonstrated by zoom-in views at various locations in the samples, LSTM provides high uniform quality throughout the sample. To directly compare the performance of LSTM with LSM, we also imaged a very wide (~ 1.5 cm) and thick (> 5 mm, i.e., as thick as mouse brain) slice of highly cleared (see Fig. 6a) rat brain, which was stained for uniform distributed blood vessels (Fig. 6a; using tomato lectin, excitation wavelength 488 nm). Previous attempts of using LSM for the imaging of rat brain resulted in very poor image quality in most of the internal parts of the brain [15] because of the heavy scattering of illumination light sheets. As shown in Fig. 6 and Additional file 10: Video 5 [29], LSTM allowed uniform high-resolution imaging of the entire tissue, whereas LSM resulted in progressively poor image quality towards the center of the sample, similar to the previous report [15].

To further demonstrate the suitability of LSTM for imaging of very large samples, we performed rapid high-resolution imaging of a very large (3.32 cm × 1.93 cm × 1 mm) uniformly expanded (using the ExM approach [7]) brain slice of a transgenic Thy1-eYFP mouse using 488 nm excitation wavelength (Fig. 6c, d, Additional file 11: Video 6 [29]). The advent of tissue expansion approaches (ExM) is enabling higher effective imaging resolution, however at the cost of hugely increased imaging time and data sizes. For example, the imaging of this sample with state-of-the-art confocal or two-photon microscopy will likely take several weeks of continuous imaging, whereas LSTM took only ~ 22 h (using 10×/0.6NA), yielding 723,200 full frame (2048 × 2048 pixels) images. The resulting dataset reveals the finest details of brain neuronal architecture (e.g., dendritic spines, Fig. 6c, d and Additional file 11: Video 6 [29]). Taken together, these imaging examples clearly demonstrate the suitability of LSTM for rapid high-resolution imaging of very large samples of different shapes. Unlike LSM, LSTM provides high uniform imaging quality even for the interior parts of the samples. In summary, these examples clearly demonstrate the complementary advantages of LSTM over LSM for rapid high-resolution imaging of large samples.

LSTM enables rapid imaging of nervous system-wide neuronal dynamics of freely motile animal

Finally, we demonstrate the compatibility of LSTM in capturing the nervous system dynamics of a highly motile animal. Live samples often undergo substantial rearrangements in their body shape and cellular density which significantly alter their local optical properties. Although LSM-based imaging methods have been effective in capturing the cellular dynamics of developing embryos and the neuronal activity of immobilized zebrafish larvae, LSM remains susceptible to large changes in shape and density of motile samples, mainly because of the use of orthogonal illumination. This limitation has been partly addressed by utilizing a sophisticated array of hardware and software components that facilitate real-time adaptation of light sheet parameters [30]. LSTM, with its non-orthogonal illumination, provides a simpler and highly effective solution. We tested this hypothesis by performing rapid volumetric calcium imaging of a highly motile Hydra, which has been recently established as an effective model for exploring the role of neuronal circuit activity on behavior [31, 32]. We found that, indeed, LSTM enables aberration-free calcium imaging of freely behaving Hydra undergoing drastic changes in body shape and cellular density in the recordings (Fig. 7a and Additional file 12: Video 7, Additional file 13: Video 8 [29]). In a way, the large and non-isomorphic body deformation of Hydra represents the worst-case scenario for tracking the activity of neurons during behavior. We validated LSTM datasets by extracting and comparing neuronal traces with previous observations, finding excellent agreement [31]. Note that, for this demonstration, we used the relatively slow process of step-wise motion of the sample stage to acquire the image stacks. The LSTM mechanism can be straightforwardly combined with piezo motor-based synchronous rapid scanning of the detection objective and also with extended detection depth of field [14].

Discussion

We reported the development of LSTM, which addresses the lateral size limitation of state-of-the-art LSM approaches. LSTM employs two symmetrically arranged oblique static light sheets generated using independent illumination objectives, and their scanning using simultaneous two-dimensional (2D) translation along (using an electrically tunable lens (ETL)) and perpendicular (using galvo scanners)
to their propagation directions, resulting in uniform illumination and detection (using synchronized rolling shutter detection of sCMOS cameras) of thin optical sections. This optical configuration eliminates the fundamental restrictions of LSM on the lateral dimensions of the imaging volumes while ensuring high imaging speed and resolution and the utilization of the entire available WD of high-NA detection objectives. The use of two light sheets (as opposed to one) ensures better quality (e.g., if one of the sheets is obstructed by opaque objects) but is not necessary. To minimize optical aberrations, we used refractive index optimized detection objectives and used 3D-printed caps (with quartz coverslips) for illumination objectives (air, low NA, and long WD) to ensure a perpendicular incidence of light sheets to the mounting media.

The enhanced performance of LSTM entails an increased overall energy load for imaging of smaller samples, such as embryos, but remains comparably as low as LSM for high-resolution imaging of large samples, as supported by the simulations (Fig. 3d, e) and empirical calculations (Fig. 3f). Moreover, the use of only the thinnest part of the light sheets enables quantitatively uniform illumination of the entire detection plane, which is a foremost requirement for 3D quantitative imaging. Therefore, LSTM is most suitable for rapid high-resolution imaging of very large samples, whereas LSM provides better performance for smaller samples.

The use of ETLs in the illumination arm for translating light sheets along their propagation direction introduces an additional component which needs to match the camera acquisition speed. The high-resolution imaging of large samples essentially requires full camera frame acquisition (i.e., 20 ms or 50 Hz for current state-of-the-art sCMOS cameras in one-direction rolling shutter mode) and the use of long travel-range motorized sample stages for step-wise z-plane acquisition (typically > 50 ms motion and settling time, in addition to the camera exposure) which is generally the rate-limiting step. This results in a typical acquisition rate of 10–15 z-planes per second as reported previously [5]. ETLs are shown to easily achieve [33] > 30 Hz for the full-range coverage without any distortions, and they can achieve even higher speeds if run continuously using sinusoidal waveforms (given that the light sheets in LSTM have a significant confocal parameter, the synchronization of sinusoidal ETL waveforms with linear row-by-row detection is feasible). The ETLs can also be easily replaced with faster acoustic tunable lenses (e.g., TAG lenses from TAG Optics) for applications requiring much higher volumetric imaging speed (e.g., for the calcium imaging of functioning nervous systems). Hence, the introduction of ETLs in the LSTM imaging procedure has no consequences for the overall imaging speed vis-à-vis LSM for high-resolution imaging of large samples. We also found that, unlike LSM, LSTM does not require a pre-imaging calibration step for the estimation of sample-position-dependent alignment parameters, resulting in overall faster imaging speeds.

We demonstrated the LSTM performance by rapid high-resolution imaging of large samples of various sizes and shapes, including the entire intact mouse central nervous system, thick coronal sections of mouse and rat brains, a large chunk of human brain, uniformly expanded brain tissue, and a highly motile Hydra. Also, due to geometric advantage, LSTM is expected to enable volumetric calcium imaging in live rodent brains, similar to SCAPE [22], especially by combining with rapid de-focusing/focusing (e.g., using ETLs in the detection arm). We also performed direct comparative imaging of the same thick rat brain coronal section using LSTM and LSM (Fig. 6) to demonstrate that LSTM indeed eliminates the limit on lateral dimensions of imaging volumes while providing high imaging speed and uniform imaging resolution. Therefore, LSTM provides complementary advantages over LSM for rapid high-resolution imaging of very large samples. These capabilities of LSTM are expected to significantly accelerate our understanding of healthy and diseased tissue architectures. Future work will include integration of super-resolution approaches (such as structured illumination) and simultaneous multi-view imaging.
Fig. 6 LSTM enables rapid uniform high-resolution imaging of very large samples. 

a For an unbiased comparison of the imaging performance of LSTM and LSM a highly cleared large rat brain tissue (~2 cm wide and ~5 mm deep; vasculature stained with tomato lectin) was imaged using the exact same detection (10×/0.6NA/8mmWD, correction collar adjusted to 1.45 refractive index) and illumination objectives (4×/0.28NA/28.5WD). Maximum intensity projections are shown. The bottom graph profiles the mean intensity across the length of the specified (dashed rectangles) regions of interest. In LSM (cyan), the intensity signal is progressively degraded towards the interior of the sample, whereas LSTM (magenta) allows uniform quality across the entire sample. The scale bars are 1 mm.

b An image stack from the sample shown in a. Maximum intensity projections (50 μm) are shown at three different depths (orange). The bounding box is 1 mm × 1 mm × 5 mm. The scale bars are 100 μm. A detailed volume rendering is shown in Additional file 10: Video 5 [29].

c Uniformly expanded (~4-fold in all three dimensions) slice of Thy1-eYFP transgenic mouse was imaged using LSTM with 10×/0.6NA/8mmWD detection objective. A rolling shutter exposure window of 0.2 ms and a full frame exposure of 20 ms were used. The resulting dataset consists of 723,200 images (2048 × 2048 pixels) and required ~22 h of acquisition time. The volume rendering was performed with 8 × 8 fold down-sampled dataset. Zoomed-in images are marked.

Additional file 11: Video 6 [29].
Conclusions

We report the development of a distinct light sheet microscopy (LSM) approach, termed light sheet theta microscopy (LSTM), which addresses the fundamental limitation of LSM on the lateral dimensions of imaging volumes due to the orthogonal illumination-detection geometry. We have presented extensive characterization of the LSTM system properties and performed several real-world high-resolution imaging experiments of very large samples, including mouse and rat brains, a large section of human brain, a highly expanded expansion microscopy sample, and rapid volumetric calcium imaging of a highly motile Hydra. These new imaging capabilities will enable numerous novel applications, including imaging of an entire post-mortem human brain (thick slab-by-slab) in a practical time-frame and direct in situ transcriptomics of whole rodent brains.

Methods

LSTM implementation

The optical layout and physical implementation details are presented in Fig. 2, Additional file 1: Figure S1, and Additional file 2: Figure S2. A complete parts list is provided in Table 1. The illumination arms of the LSTM instrument consist of a laser source (Omicron-Laserage, Rodgau, Germany; SOLE-6 engine containing four wavelengths: 405, 488, 561, and 647 nm), a cylindrical lens (Thorlabs, Newton, NJ, USA; LJ1695RM-A, f = 50 mm), a vertical slit (Thorlabs, VA100C), an iris (Thorlabs, CP20S), an electrically tunable lens (Optotune, Dietikon, Switzerland), and a motorized XY table (Theta, USA) for sample positioning.

Fig. 7 LSTM enables rapid volumetric imaging of highly motile animals. Live samples can undergo substantial non-isomorphic rearrangements in their body shape and cellular density, resulting in continuously changing local optical properties. LSM is particularly susceptible to misalignments and other aberrations because of the use of orthogonal light sheet illumination. LSTM is uniquely suitable for rapid volumetric live imaging of such difficult samples, as demonstrated by imaging of highly motile Hydra. a Hydra image is shown at different time points to highlight the non-isomorphic changes in freely moving animal. b LSTM was used to perform long-term (>1 h demonstrated, Additional file 12: Video 7 [29]) high-resolution live imaging of an adult Hydra expressing GCaMP6s [31]. Each volume consists of 17 z-planes. Manual tracking and analyses of calcium signaling were performed for the first ~500 s of recording. Maximum intensity projections covering the two halves are shown. Representative neuronal traces are shown for cells marked in corresponding colors. As shown in Additional file 13: Video 8 [29], the neuronal traces correlate with the rapid longitudinal contraction behavior of Hydra, and the other two traces are part of rhythmic potential circuits, in excellent agreement with the observations reported recently [31]. Scale bars are 100 μm.
Switzerland; EL-16-40-TC), a galvo scanner (Thorlabs, GVS001), a scan lens (Thorlabs, CLS-SL), a tube lens (Thorlabs, TTL200), and the illumination objective (Olympus Macro 4×/0.28NA/28.5 mmWD air). Since the illumination objectives were air objectives, we used a 3D-printed cap (using Ultimaker 2+ extended, in polylactic acid (PLA)) containing a 1-in. diameter quartz coverslip, to allow dipping in immersion oil, ensuring that the low-numerical aperture (NA) light rays enter the media near perpendicularly. The detection arm is composed of a detection objective (Olympus, XLPLN10XSVMP/10× or XLSLPLN25XGMP/25×), a tube lens (Thorlabs, TTL200), a multi-band emission filter (Semrock, FF01-432/515/595/730-50-D), and an sCMOS camera (Hamamatsu Orca Flash 4.0 V3). The illumination arms were vertically mounted at an approximately 60° angle relative to the detection axis. To facilitate the optical alignment adjustments, all three optical arms were mounted on manual translation stages (Thorlabs, TBB0606) attached to the breadboard. We used a 3D-printed (using Ultimaker 2+ extended in PLA) open-top sample chamber that was filled with an immersion oil of refractive index 1.454 (Cargille Laboratories, Cedar Grove, NJ, USA). The CLARITY-cleared (refractive index = ~1.454) sample was mounted in a quartz cuvette (refractive index ~1.45, FireflySci, Staten Island, NY, USA) which was then affixed to appropriate grooves at the base of the sample chamber (Additional file 1: Figure S1). The 3D model of the LSTM microscope was made with Autodesk Inventor 2017.

**LSM imaging**

All LSM imaging experiments were performed using the COLM implementation as described previously [5]. Briefly, the optical components were the same as described for LSTM, i.e., detection objectives: Olympus 10×/0.6NA/8mmWD or Olympus 25/1.0NA/8mmWD with a correction collar for the refractive indices of water to oil; sCMOS camera: Hamamatsu Orca Flash 4.0 V3; low-NA illumination objective: Olympus Macro 4×/0.28 NA; tube lens: Thorlabs TTL200; and scan lens: Thorlabs, CLS-SL. The dynamic light sheets are generated by rapid scanning of Gaussian beams (Thorlabs, GVS102). Similar to LSTM, COLM uses synchronized illumination-detection to improve the imaging quality. Additionally, an automated-alignment parameter calibration (using linear adaptation) corrects for misalignment artifacts across the whole sample space.

**LSTM geometric constraint calculations**

The physical geometric constraints of the oblique arrangement of illumination and detection objectives were analyzed by calculating the upper and lower bounds (Fig. 3a, Additional file 3: Figure S3) with the following criteria: the illumination objective should not touch the detection objective (Fig. 3a top) and it should not extend below the physical extent of the detection objective (Fig. 3a middle).

The range of allowable angular positions was calculated by taking the effective working distances (WDs) and the objective diameters into account as shown in the schematics of Additional file 3: Figure S3, resulting in the following relationships:

\[ W_2 \sin(\theta_i) = \frac{D_1}{2} + D_2 \cos(\theta_i) \]

\[ W_2 \cos(\theta_f) = W_1 + D_2 \sin(\theta_f) \]

where \(W_1\) and \(W_2\) are the WDs of the detection and illumination objectives respectively, \(D_1\) and \(D_2\) are the diameters of the detection and illumination objectives respectively, and \(\theta_i\) and \(\theta_f\) are the boundary angular positions. Since we used a Macro 4×/0.28NA/29.5WD (Olympus) air objective for illumination, the approximate effective WDs were calculated as shown in Additional file 3: Figure S3. For most of the experiments we used a 10×/0.6NA/8mmWD objective (Olympus) in the detection arm with values of \(W_1 = 8\), and \(D_1 = 40\). For this combination, we found the allowable angular range to be 43.3° to 62.3°, which served as our initial guide for identifying the maximum possible angular positioning. We used ~60° as the final angle separation. All calculations were performed in MATLAB.

**Illumination depth calculations**

We used geometric calculations (Fig. 3b) to estimate the maximum illumination path lengths (MIPLs) of LSTM as \(t \cos(\theta)\), where \(t\) is the sample thickness to be imaged and \(\theta\) is the angle between the illumination propagation direction and the detection axis. The MIPL in LSM would be the same as the sample width (\(w\)). We calculated the ratio of these illumination path lengths, which was then converted into a binary representation by thresholding at 1 and plotted as a heat map, shown in Fig. 3b. The edge effects were approximated.

**LSTM effective light sheet thickness calculations**

Due to the non-orthogonal incidence of the light sheets on the detection plane, the effective light sheet thickness can be approximated as the projection of the original thickness onto the detection direction, resulting in \(b / \sin(\theta)\), where \(b\) is the actual light sheet thickness at the most focused position, and \(\theta\) is the angle of incidence relative to the detection axis. The relationship is plotted in Fig. 3c.

**Imaging experiments**

All the imaging experiments are summarized in Table 2. We used the passive CLARITY method (as described previously [5]) for all the tissue clarification experiments. The hydrogel monomer (HM) solution recipe consisted of 1–4% (wt/vol)
Table 2  Summary of imaging experiments reported in this study

| Samples                              | Fig. No. | Label                | Det. objective | Illum. objective | Imaging volume dimensions | No. of images/raw data | Imaging time |
|--------------------------------------|----------|----------------------|----------------|------------------|---------------------------|------------------------|--------------|
| Thick human brain tissue             | 4        | DAPI                 | 10×/0.6NA/8mmWD | 4×/0.28NA/28.9WD | ~ 10.5 mm × 14.1 mm × 3 mm | 116,736/~ 0.97 TB      | ~ 2.7 h      |
| Mouse brain with attached spinal cord| 5a       | Thy1-eYFP            | 10×/0.6NA/8mmWD | 4×/0.28NA/28.9WD | 11.8 mm × 276 mm × 5.2 mm | 388,687/~ 3.3 TB       | ~ 9 h        |
| Thick mouse brain slice              | 5b       | Thy1-eYFP            | 10×/0.6NA/8mmWD | 4×/0.28NA/28.9WD | ~ 9.6 mm × 13.5 mm × 5.34 mm | 256,560/~ 2.1 TB        | ~ 5.9 h      |
| Thick mouse brain slice              | 5c       | Thy1-eYFP            | 25×/1.0NA/8 mm/WD | 4×/0.28NA/28.9WD | 6 mm × 9.6 mm × 1.9 mm³  | 211,616/~ 1.8 TB       | ~ 4.9 h      |
| Thick rat brain slice expanded (~4x) | 6a, b    | Tomato lectin       | 10×/0.6NA/8mmWD | 4×/0.28NA/28.9WD | ~ 20 mm × 16.5 mm × 3.6 mm deepb | 285,821 slices/24 TB  | ~ 6.6 h      |
|                                     | 6c       | Thy1-eYFP            | 10×/0.6NA/8mmWD | 4×/0.28NA/28.9WD | 33.2 mm × 193 mm × 2 mm²  | 723,200/~ 6 TB         | ~ 22 h       |
| Hydro live imaging                  | 7        | GCaMP6s              | 10×/0.6NA/8mmWD | 4×/0.28NA/28.9WD | 1.2 mm × 1.2 mm × 0.136 mm | 23,001/~ 193 GB        | ~ 1 h live imaging |

Summary of the datasets reported in this report.

*The image volume acquired was ~ 6 mm × 9.6 mm × 1.9 mm; however, due to constraints of high-quality volume rendering, a smaller (0.5-mm-thick) subset was used for the rendering shown in Fig. 5c

bThe approximate imaging volume was ~ 20 mm × 16.5 mm × 3.6 mm, and a few ~ 5-mm-deep image stacks were acquired to demonstrate the imaging depth in Fig. 6b

cThe imaging volume acquired was ~ 33.2 mm × 193 mm × 2 mm to ensure complete coverage of ~ 1-mm-thick expanded non-rigid tissue

TB terabytes, GB gigabytes, h hours
acrylamide, 0.05% (wt/vol) bisacrylamide, 4% paraformaldehyde (PFA), 1x phosphate-buffered saline (PBS), deionized water, and 0.25% thermal initiation VA-044 (Wako Chemicals, Boston, MA, USA; NC0632395). All animal procedures were followed according to Institutional Animal Care and Use Committee (IACUC) guidelines. For whole brain clearing, transcardiac perfusion was performed with 20 mL HM solution, followed by overnight incubation at 4 °C. The rat brain was perfused with 4% PFA, post-fixed for 16 h, and then frozen in isopentane for storage. The frozen brain was thawed at room temperature in PBS buffer, then sliced and incubated in HM solution overnight at 4 °C. The human brain tissue was incubated in 4% PFA for ~2 days, followed by incubation in HM solution overnight at 4 °C. All the perfused tissues were de-gassed and then stored at 37 °C for 3–4 h for hydrogel polymerization. The tissues were cleared by incubating (with shaking) in clearing buffer (4% (wt/vol) sodium dodecyl sulfate (SDS), 0.2 M boric acid, pH 8.5) at 37 °C until clear (2–3 weeks). Afterwards, the tissues were washed with 0.2 M boric acid buffer (pH 8.5) with 0.1% Triton X-100 for up to 24 h. The cleared tissue was labeled with DAPI (1 μg/mL final concentration) and/or the blood vessel marker tomato lectin (Vector Labs, Burlingame, CA, USA; FL-1171) by incubating in the labeling solution for 3–4 days. After washing with the buffered solution (0.2 M boric acid buffer, pH 7.5, 0.1% Triton X-100), the tissue was transferred into 85–87% glycerol solution in graded fashion (i.e. 25%, 50%, 65%, and finally 87%) for final clearing and imaging. For uniform tissue expansion (4–4.5x uniformly), a Thy1-eYFP mouse brain slice (250 μm, perfused and fixed with 4% PFA and sliced with vibratome) was gelled and digested following the protein retention expansion microscopy (proExM) protocol [34]. The sample was stored in 1x PBS before changing the buffer to 65% glycerol (with 2.5 mg/mL 1,4-diazabicyclo[2.2.2]octane (DABCO)) for the LSTM imaging. All imaging experiments were performed with an effective light sheet thickness of 2–5 μm.

Image analyses
A TeraStitcher [35]-based pipeline [5] was used for the stitching of acquired image stack tiles of all the datasets. Maximum intensity projections and other linear image contrast adjustments were performed using Fiji [36, 37] and MATLAB. All volume renderings were performed using Amira (FEI, Lausanne, Switzerland). All the fluorescent bead image analyses were performed using Fiji. To calculate the axial full width at half maximum (FWHM), x-z projections of the bead image stacks were used. For individual beads a line intensity profile was calculated along the central axis, followed by manual calculations of FWHM intensity values.

Illumination energy load calculations
The procedure is summarized in Additional file 7: Figure S5. To calculate the total illumination energy load in LSTM, we performed a simulation of the step-wise scanning of the sample through the illuminating light sheet. A horizontal plane across an entire sample can be imaged with approximately non-overlapping thin sheets of light; therefore, the total energy is calculated by step-wise scanning of the sample through the illumination volume. All voxels receiving illumination are incremented by 1. The final energy load is calculated as the total sum of accumulated illuminations in all voxels. The procedure was implemented for a range of parameters and two detection objectives (10×0.6NA/8/mmWD and 25×1.0NA/8/mmWD). For LSM calculations, each voxel is illuminated (w/f) times, where w is the width of the sample, and f is the field-of-view size of the detection arm. Therefore, the total energy load is approximated as (w/f)*number of voxels. Note that the energy load in LSM as well as LSTM scales up by the same constant factor, which cancels out in the ratio.

Additional files

**Additional file 1: Figure S1.** LSTM microscopy implementation. (a) Image of the physical LSTM setup. (b) 3D model of LSTM and the sample mounting system. The 3D-printed sample chamber is designed to accommodate large biological samples of virtually any dimensions, while still allowing the objectives to be immersed in the immersion oil. Two transparent glass windows, located on the lateral sides, provide visual view of the sample for ease of positioning. An additional window is realized at the bottom part of the chamber to allow the illumination light to pass through. An additional adapter was designed to allow mounting a prism mirror at about approximately 10° from the normal surface to facilitate the optical alignment of the system. (PDF 596 kb)

**Additional file 2: Figure S2.** Detailed annotation of LSTM optical path shown in Fig. 2a. (PDF 571 kb)

**Additional file 3: Figure S3.** Physical constraints of positioning illumination and detection objects. (a) Schematics showing calculations of the elongated working distance (EWD) of the air illumination objective (Olympus Macro 4x/0.28NA/29.5WD Air) when used in immersion liquid (refractive index 1.454). Original working distance (OWD) is the working distance in air according to the objective specifications. A thin quartz coverslip and a 3D-printed cap were used to seal the illumination objectives. EWD was estimated to be 43.84 mm. (b) Geometric constraints calculation for the co-arrangement of the illumination and detection objectives. The two boundary conditions are shown in blue and green shading of the illumination objective. For the upper bound limit (blue), the relationship among different parameters is defined by the equation W2 * sin(θ) = ½ + 2D₂ * cos(θ) / W₁. For the lower bound limit (green), it is defined by W2 * cos(θ) = W₁ + 2D₂ * cos(θ) / W₁ and W₂ are the effective working distances of detection and illumination objectives respectively. D₁ and D₂ are the diameters of the detection and illumination objectives respectively. θ₁ and θ₂ are the angular positions of upper and lower bounds respectively. For the 4x/0.28NA/29.5 mmWD (as illumination objective) and 10x/0.6NA/8/mmWD (as detection objective), the calculated θ₁ and θ₂ are 43.32° and 63.37° respectively. This range served as a starting point during the optical alignment of the system. (PDF 596 kb)

**Additional file 4: Video 1.** 3D model of LSTM implementation. The 3D modeling and rendering was performed using Autodesk Inventor 2017, and the animation was performed using Autodesk Fusion 360 2017 and MATLAB. The components labeled are LS (laser source), collimator, ND.
Additional file 5: Video 2. Comparison of image volumes acquired with LSTM in 1-axis scan (1-AS) and 2-axis scan (2-AS) modes. The 3D rendering visualizes the image stacks acquired from the same sample (human brain section stained with DAPI) with LSTM in 1-AS and simultaneous 2-AS modes. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MP4 82944 kb)

Additional file 6: Figure S4. Comparison of maximum illumination path lengths in LSTM and LSM. The graphs plot the binarized ratios (w/(w+t)) of maximum illumination path lengths required for complete coverage of samples of various widths (w) and thicknesses (t) for different angular arrangements. Magenta and cyan regions mark the combinations of w and t for which the illumination path lengths were smaller in LSTM and LSM, respectively. (PDF 262 kb)

Additional file 7: Figure S5. Total illumination energy load in LSTM vs. LSM. The schematic summarizes the calculations of total energy loads imparted in LSTM and LSM for imaging of a sample of specific dimensions, imaged with a specific detection objective. (a) In LSTM, a horizontal plane across the entire sample is imaged with approximately non-overlapping thin sheets of light. Therefore, total energy load can be calculated by step-wise sampling of the sample (for each plane) through the illuminating light. For each of the steps, all voxels that receive light are incremented by 1. The procedure was implemented for a range of parameters and two detection objectives (10x/0.6NA/8mmWD and 25/1.0NA/8mmWD). (b) In LSM a stack (or tile) is acquired by approximately non-overlapping thin sheets of light. The total energy load is calculated by summing up the illumination for all tiles in a row along the sample width. Note that the dwell time of illumination line profile is similar for both LSTM and LSM (scanned light sheet implementation, e.g., COLUM). The energy load for tiles along the sample length scales up by the same constant factor in LSTM and LSM; therefore, we only simulated one row of tiles along the sample width. (PDF 556 kb)

Additional file 8: Video 3. High-resolution LSTM imaging of intact Thy1-eYFP mouse central nervous system. The bounding box for the entire sample is 11.8 mm x 27.6 mm x 5.2 mm, and for the subvolume shown is 5.1 mm x 3.1 mm x 3.5 mm. The raw data was down-sampled 2 x 2 fold (to make the volume rendering feasible) for the subvolume rendering. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MOV 143360 kb)

Additional file 9: Video 4. High-resolution LSTM imaging of a large tissue of Thy1-eYFP mouse brain. The bounding box is 9.6 mm x 13.5 mm x 5.34 mm. The raw data was down-sampled 4 x 4 fold to make the volume rendering feasible. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MOV 167936 kb)

Additional file 10: Video 5. Visualization of an image stack of vasculature-stained rat brain tissue. This video visualizes an image stack acquired from a large rat brain slice (stained for vasculature with tomato lectin) using LSTM in 2-AS mode. The bounding box is 1 mm x 1 mm x 5 mm. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MOV 143360 kb)

Additional file 11: Video 6. High-resolution LSTM imaging of a large expanded section of Thy1-eYFP mouse brain. A thin (250 μm) coronal section was expanded ~4-fold using proExM procedure and imaged using LSTM with 10x/0.6NA detection objective. The resulting dataset (~6 TB) consisted of 723,300 full frame images (2048 x 2048). The data was down-sampled 8 x 8 fold to allow high-quality volumetric rendering. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MOV 439296 kb)

Additional file 12: Video 7. Rapid volumetric calcium imaging of highly motile Hydra. CCAmP6 expressing Hydra was imaged using LSTM with 10x/0.6NA objective. Maximum intensity projections are shown for the two halves of the volume. First occurrences of longitudinal and radial contractions are annotated. The scale bar is 100 μm. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MOV 66662 kb)

Additional file 13: Video 8. Neuronal activity traces of representative neurons. A visualization of the neuronal traces shown in Fig. 7b. The high-resolution video is available in the figshare repository at https://doi.org/10.6084/m9.figshare.c.4072160. (MOV 11161 kb)

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Availability of data and materials
A complete computer-aided design (CAD) model of LSTM and other related resources are detailed in the additional files. A complete parts list is included as Table 1. High-resolution movies and figures will also be shared at the resource webpage: http://tomerlab.org/lstm/. The datasets generated and/or analyzed during the current study are available in the figshare repository (https://doi.org/10.6084/m9.figshare.c.4072160).

Authors’ contributions
RT conceived the project and designed the microscopes. BM, MSD, and RT built the microscopes. MSD, BM, and RT performed the imaging experiments. MCA assisted with the experiments. CD prepared the Hydra samples for the live imaging experiments. Ry conceived of the Hydra project in general, and supervised the Hydra sample preparations used in this work. SA, RG, and ESB generated the expanded tissue. OH supported BM, RT, BM, and MSD analyzed the data and wrote the paper with inputs from all authors. RT supervised all aspects of the work. All authors read and approved the final manuscript.

Ethics approval and consent to participate
Not applicable.

Competing interests
Columbia University has filed a patent application for LSTM.

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References

1. Richardson DS, Lichtman JW. Clarifying tissue clearing. Cell. 2015;162(2):246–57.
2. Romanov RA, Zeisel A, Bakker J, Girach F, Helluyaz A, Tomer R, Alpar A, Mulder J, Clotman F, Kimpener E, et al. Molecular interrogation of hypothalamic organization reveals distinct dopamine neuronal subtypes. Nat Neurosci. 2017;20(2):176–88.
3. Lemer TN, Shilyansky C, Davidson TJ, Evans KE, Beier KT, Zalocusky KA, Crow AK, Malenka RC, Luo L, Tomer R, et al. Intact-brain analyses reveal distinct information carried by Snc dopamine subcircuits. Cell. 2015;162(3):345–47.
4. Chung K, Wallace J, Kim SY, Kalyanasundaram S, Andalman AS, Davidson TJ, Mirzabezov JJ, Zalocusky KA, Mattis J, Denisin AK, et al. Structural and molecular interrogation of intact biological systems. Nature. 2013;497(7449):332–7.
5. Tomer R, Ye L, Hsieh B, Deisseroth K. Advanced CLARITY for rapid and high-resolution imaging of intact tissues. Nat Protoc. 2014;9(7):1682–97.
6. Tanaka N, Kanatani S, Tomer R, Sahligen C, Kronquist P, Pczynski D, Louhivouri L, Kis L, Lind C, Mitura P, et al. Whole-tissue biopsy phenotyping of three-dimensional tumours reveals patterns of cancer heterogeneity. Nat Biomed Eng. 2017;1(10):796–806.
7. Chen F, Tillberg PW, Boyden ES. Optical imaging. Expansion microscopy. Science. 2015;347(6221):543–8.
8. Chang JB, Chen F, Yoon YG, Jung EB, Babcock H, Kang JS, Asano S, Suk HJ, Pak N, Tillberg PW, et al. Iterative expansion microscopy. Nat Methods. 2017;14(6):593–9.
9. Mei E, Fonitochov PA, Graves R, Campion M. A line scanning confocal fluorescence microscope using a CMOS rolling shutter as an adjustable aperture. J Microsc. 2012;244(3):269–76.
10. Welschenkens R, Zimmermann B, Kempe M. High-speed confocal fluorescence imaging with a novel line scanning microscope. J Biomed Opt. 2006;11(6):040411.
11. Weber M, Mickoleit M, Huiskens J. Light sheet microscopy. Methods Cell Biol. 2014;123:193–215.
12. Stelber EH. Light-sheet fluorescence microscopy for quantitative biology. Nat Methods. 2015;12(1):23–6.
13. Migliori B, Datta MS, Tomer R. Advanced light microscopy enables rapid mapping of brain structure and function in high resolution. Laser Focus World. 2016;52(2):47–52.
14. Tomer R, Lovett-Barron M, Kauvar I, Andalman A, Burns VM, Sankaran S, Grosenick L, Broxton M, Yang S, Deisseroth K. SPED light sheet microscopy: fast mapping of biological system structure and function. 2015;13(7):796–806.
15. Stefaniuk M, Guzda EJ, Pawlowska M, Legutko D, Matryba P, Koza P, Konopka W, Owczarczak D, Wawrzynek M, Loza-Alvarez P, et al. Light-sheet microscopy imaging of a whole cleared rat brain with Thy1-GFP transgene. Sci Rep. 2016;6:28209.
16. Holekamp TF, Turaga D, Holy TE. Fast three-dimensional fluorescence imaging of activity in neural populations by objective-coupled planar illumination microscopy. Neuron. 2008;57(5):661–72.
17. Wu Y, Ghitani A, Christensen RS, Santella A, Du Z, Rondeau G, Bao Z, Colon-Ramos D, Shroff H. Inverted selective plane illumination microscopy (SPIM) enables coupled cell identity lineaging and neurodevelopmental imaging in Caenorhabditis elegans. Proc Natl Acad Sci U S A. 2011;108(24):976–82.
18. Wu Y, Wawrzusin P, Senseney J, Fischer RS, Christensen RS, Santella A, York AG, Winter PW, Waterman CM, et al. Spatially isotropic four-dimensional imaging with dual-view plane illumination microscopy. Nat Biotechnol. 2013;31(11):1032–8.
19. Strad P, Gunther S, Reichmann K, Kriz U, Balas B, de Medeiros G, Norlin N, Hiliarri T, Hufnagel L, Ellenberg J. Inverted light-sheet microscopy for imaging mouse pre-implantation development. Nat Methods. 2016;13(2):139–42.
20. Glaeser AK, Rieder NP, Chen Y, McCarty EF, Yin C, Wei L, Wang Y, True LD, Liu JTC. Light-sheet microscopy for slide-free non-destructive pathology of large clinical specimens. Nat Biomed Eng. 2017;1:0084.
21. Wu Y, Chandris P, Winter PW, Kim EY, Jaumeouille V, Kumar A, Guo M, Leung JM, Smith C, Rey-Suarez I, et al. Simultaneous multiview capture and fusion improves spatial resolution in wide-field and light-sheet microscopy. Optica. 2016;3(8):897–910.