1 **Title:** Multi-immersion open-top light-sheet microscope for high-throughput imaging of cleared tissues

2 **Authors:** Adam K. Glaser[1][*], Nicholas P. Reder[2], Ye Chen[1], Chengbo Yin[1], Linpeng Wei[1], Soyoung

3 Kang[1], Lindsey A. Barner[1], Weisi Xie[1], Erin F. McCarty[2], Chenyi Mao[3], Aaron R. Halpern[3], Caleb R.

4 Stoltzfus[4], Jonathan S. Daniels[5], Michael Y. Gerner[4], Philip R. Nicovich[6], Joshua C. Vaughan[3,7],

5 Lawrence D. True[2], and Jonathan T.C. Liu[1,2][*]

6 [1] Department of Mechanical Engineering, University of Washington, Seattle, WA USA

7 [2] Department of Pathology, University of Washington, Seattle, WA USA

8 [3] Department of Chemistry, University of Washington, Seattle, WA USA

9 [4] Department of Immunology, University of Washington, Seattle, WA USA

10 [5] Applied Scientific Instrumentation, Eugene, OR USA

11 [6] Allen Institute for Brain Science, Seattle, WA USA

12 [7] Department of Physiology and Biophysics, University of Washington, Seattle, WA USA

13 [*]To whom correspondence should be addressed: akglaser@uw.edu or jonliu@uw.edu

14 **One Sentence Summary:** Glaser *et al.* describe a multi-immersion open-top light-sheet microscope that

15 enables simple and high-throughput imaging of large numbers of preclinical and clinical specimens

16 prepared with a variety of clearing protocols.

Abstract

Recent advances in optical clearing and light-sheet microscopy have provided unprecedented access to structural and molecular information from intact tissues. However, current light-sheet microscopes have imposed constraints on the size, shape, number of specimens, and compatibility with various clearing protocols. Here we present a multi-immersion open-top light-sheet microscope that enables simple mounting of multiple specimens processed with a variety of protocols, which will facilitate wider adoption by preclinical researchers and clinical laboratories.

Main Text

Recent advances in tissue clearing have provided unprecedented visual access to structural features and molecular targets within large intact specimens [1]. These clearing approaches have the potential to accelerate new discoveries across multiple fields of research, including neuroscience, developmental biology, and anatomic pathology [2-5] (**Supplementary Figure 1**). However, fully harnessing the benefits of tissue clearing requires user-friendly and versatile microscopes for three-dimensional (3D) imaging of a wide variety of content-rich specimens.

Over the past decade, light-sheet fluorescence microscopy (LSFM) has emerged as the technique of choice for fast and gentle 3D microscopy of relatively transparent specimens [6, 7]. In LSFM, a thin sheet of light illuminates a specimen such that fluorescence is selectively excited within a single "optical section". The fluorescence is imaged onto a high-speed camera in the direction perpendicular to the light sheet. Initial LSFM systems were purposefully designed for imaging small model organisms, often living, over unprecedented long timescales [8]. However, in recent years, the advantages of LSFM have also been harnessed for 3D imaging of large cleared tissues, where speed is the main advantage over alternative microscopy methods [9-11]. Unfortunately, most LSFM architectures impose constraints on the size, shape, number of specimens, and/or compatibility with various tissue clearing protocols (**Supplementary Discussion**). In addition, specimen mounting has typically been complex and slow, which has hampered the application of LSFM for high-throughput imaging (**Supplementary Figure 2**).

42   Here, we have developed an easy-to-use multi-immersion open-top light-sheet (OTLS)

43   microscope in which all optical components are positioned below a transparent holder so that specimens

44   may be conveniently placed on top in a manner similar to a "flatbed scanner" (**Fig. 1a** and

45   **Supplementary Video 1**) [5, 12, 13]. An air objective delivers an illumination light sheet through an

46   immersion chamber, holder, and into a specimen, where fluorescence is generated and then collected by

47   a multi-immersion objective in the direction orthogonal to the light sheet. The multi-immersion

48   capability of our OTLS system spans the refractive index range of all published tissue-clearing methods

49   (including expansion, aqueous, and solvent-based protocols). A key to achieving this is by precisely

50   matching the refractive index of the immersion medium, holder material, and specimen (**Fig. 1b**). A

51   plano-convex lens with a radius of curvature matched to the focus of the illumination beam acts as a

52   solid immersion lens (SIL) to prevent spherical and off-axis aberrations from occurring as the light sheet

53   enters the system. The point spread function (PSF) of the system is shown in **Fig. 1c**, in which sub-

54   micron resolution is achieved lateral to the collection axis. Resolution along the collection axis (~3.5

55   µm) is comparable to the thickness of a typical slide-mounted histology section. Note that the numerical

56   apertures (NA) of both the illumination and collection beams are proportional to the matched refractive

57   index of the immersion medium, holder material, and specimen (**Fig. 1d**). The open-top architecture

58   (**Fig. 1e**) enables fast and simple mounting of multiple specimens with diverse shapes such as human

59   biopsies, thin tissue slices, and whole mouse organs in modular holders (**Fig. 1f**). Large-volume imaging

60   at a speed of ~1 min/mm^3 is achieved entirely through stage scanning, in which a series of adjacent

61   volumetric image strips (scanned in the x dimension) are tiled in the lateral (y) and vertical (z) directions

62   (**Fig. 1g** and **Supplementary Video 2**). The working distance of the collection objective allows for an

63   imaging depth of up to 0.5 cm. In comparison to a previously published open-top prototype [5], our new

64   OTLS system exhibits (1) improved axial and lateral resolution (an order-of-magnitude reduction in the

65   focal volume), (2) ~20× greater imaging depth, (3) mitigation of shadowing artifacts, and (4) multi-

66   immersion capabilities (**Supplementary Figure 3**).

67   Since precise index matching of the immersion media, holder, and specimen are necessary for

68   aberration-free imaging, we performed optical simulations (**Fig. 2a**) to explore the tolerance of the

69   system to the optical path difference ($\Delta n \times t$) introduced by a holder with a refractive index mismatch,

70   Δn, and thickness, t. We quantified the Strehl Ratio, S, of the system as a function of $\Delta n \times t$, and

71   determined that $\Delta n \times t < 0.002$ is necessary for near-diffraction-limited imaging ($S > 0.8$). Based on these

72   findings, we surveyed glasses and monomers/polymers as potential holder materials, and determined the

73   maximum-allowed thickness, t_{max}, based upon the intrinsic mismatch, Δn, between these materials and

74   published clearing protocols (**Fig. 2b**). These chemical-reagent and material combinations are enabling

75   for our OTLS system, and are beneficial for other cleared tissue imaging systems. To validate the multi-

76   immersion capabilities of our system, we fabricated customized holders for a variety of clearing

77   protocols spanning solvent, aqueous, and expansion-based protocols, and performed high-throughput

78   imaging of diverse tissue types and shapes (a summary of the imaging parameters for all specimens are

79   shown in **Supplementary Table 3**).

80   Solvent-based protocols involve dehydration of tissue specimens and replace the water with

81   organic reagents of a higher refractive index ($n = 1.51 - 1.56$). These solvents are optically compatible

82   with several high-index glasses. Unfortunately, these higher-index glasses also have high Abbe numbers

83   (i.e. a low variation in refractive index versus wavelength) compared to organic solvents, which

84   typically have low Abbe numbers. In addition, glasses, which are brittle, must be relatively thick and are

85   not easily machined. Therefore, we explored the use of several monomers/polymers. Despite being

86   optically compatible, we observed that styrene-based polymers (e.g. PS, SMMA, and SAN) were all

87   destroyed after exposure to solvent-based clearing reagents. However, we identified three optically and

88   chemically compatible resin-based monomers (HIVEX, NK55, and KOC55, used for manufacturing

89   eyeglass lenses) that are ideal for many solvent-based clearing protocols, including DISCO, BABB, and

90   ECi [14-18]. We demonstrated the potential clinical utility of our system by imaging multiple human

91   prostate biopsies *in toto* (**Fig. 3a** and **Supplementary Videos 3 and 4**). We found that ECi-clearing ($n =$

92   1.56) is well-suited for this application due to its clearing efficacy, as well as low toxicity [19]. A

93   custom HIVEX holder ($n = 1.55$) was machined with channels to accommodate multiple biopsies (13 in

94   this case) and to keep them aligned and parallel. Due to the effective tissue clearing of ECi and precise

95   refractive index matching with the HIVEX holder, we were able to resolve sub-nuclear features in

96   benign and malignant prostate glands throughout the entire 1-mm diameter of all biopsies. As noted by

97   us and others, this ability to visualize the 3D structure of human cancers should improve prognostication

98   and treatment decisions [3, 5, 20].

99   Aqueous-based protocols typically involve removing lipids using detergents, followed by

100   immersion in diluted water-soluble reagents, resulting in a refractive index range of $n = 1.38$ to 1.49

101   [21]. For most aqueous-based protocols, there is an optically compatible glass such that $\Delta n \leq 0.01$. For

102   example, fused silica ($n = 1.46$) is well-suited for several protocols, including CLARITY [2, 22].

103   However, most transparent monomers/polymers (with the exception of PMMA) have either a high ($n >$

104   1.49) or low ($n < 1.38$) refractive index and are therefore not optically compatible with aqueous-based

105   protocols. To demonstrate compatibility of our system with aqueous protocols, we cleared adjacent

106   slices of a mouse brain (200 μm thick) using simple immersion in diluted TDE ($n = 1.46$) and imaged

107   them using a 1 mm thick fused-silica plate ($n = 1.46$) with a large 10×10 cm viewing area (**Fig. 3b** and

108   **Supplementary Video 5**). High-magnification views of our datasets enabled unambiguous visualization

109   of neuronal structures. We also cleared and labeled multiple mouse organs using the Ce3D protocol ($n =$

110   1.49) and imaged them in a single automated session using a customized 6-well plate with a 0.5 mm

111   thick PMMA bottom substrate ($n = 1.49$) (**Fig. 3c** and **Supplementary Video 6**). Lymph nodes, heart,

112   prostate, and lung tissues (1 to 4 mm thick) were mounted in separate wells and imaged *in toto* for 3D

113   visualization of immune cells in lymph nodes, the ventricles and valves within a heart, the glandular

114   network within the prostate, and the bronchial tree within the lung [23]. Note that the use of multi-well

115   plates (e.g. 6-well or 96-well plates) prevents contamination between samples, which is particularly

116   important for clinical assays and cell culture (a feature not possible with most other LSFM systems, see

117   **Supplementary Discussion**).

118   Expansion-based protocols provide a magnified view of structures that are otherwise too small to

119   resolve with a given microscope [24-26]. To date, these expanded hydrogel specimens have consisted

120   mostly of water and therefore have a refractive index close to that of pure water ($n = 1.33$). While there

121   are currently no glasses at this refractive index, fluoro-polymers (i.e., Teflons), including fluorinated

122   ethylene-propylene (FEP) and poly-tetra-fluoro-ethylene (PTFE) possess a compatible refractive index

123   ($n = 1.34$). These materials can be manufactured as thin sheets and stretched tight as "drumhead"

124   surfaces that are ideal holders for expanded specimens. Using a customized drumhead, we imaged a 4×

125   expanded 200 μm thick kidney section (**Fig. 3d** and **Supplementary Videos 7-9**). After expansion, the

126   physical size of the tissue was 2.1×3.2×0.1 cm. Representative zoomed-in views provide a highly

127   detailed view of 3D structures such as glomeruli, renal tubules, and blood vessels.

128   We have developed and characterized a multi-immersion OTLS microscope that enables high-

129   throughput automated imaging of optically cleared specimens with an ease of use that should facilitate

130   broader adoption of light-sheet-based 3D microscopy by both researchers and clinicians. Our system

131   imposes minimal constraints on specimen shape/size and allows for fast and convenient mounting of

132   multiple tissue specimens for automated imaging. The system provides an imaging depth of 0.5 cm over

133   a maximum lateral area of 10×10 cm at a speed of ~ 1 min/mm^3, which can be tailored for specific

134   research applications in future designs (see **Supplementary Figure 22**). We have shown that our system

135   can interface with a variety of modular specimen holders tailored for specific tissue types and clearing

136   protocols, with the ability to isolate different specimens in individual wells. Due to its open-top

137   geometry, our system also provides unsurpassed versatility to interface with future tissue-based

138   protocols and a wide range of potential accessory technologies such as microfluidic devices, single-cell

139   electrophysiology, and micro-aspiration [27-30].
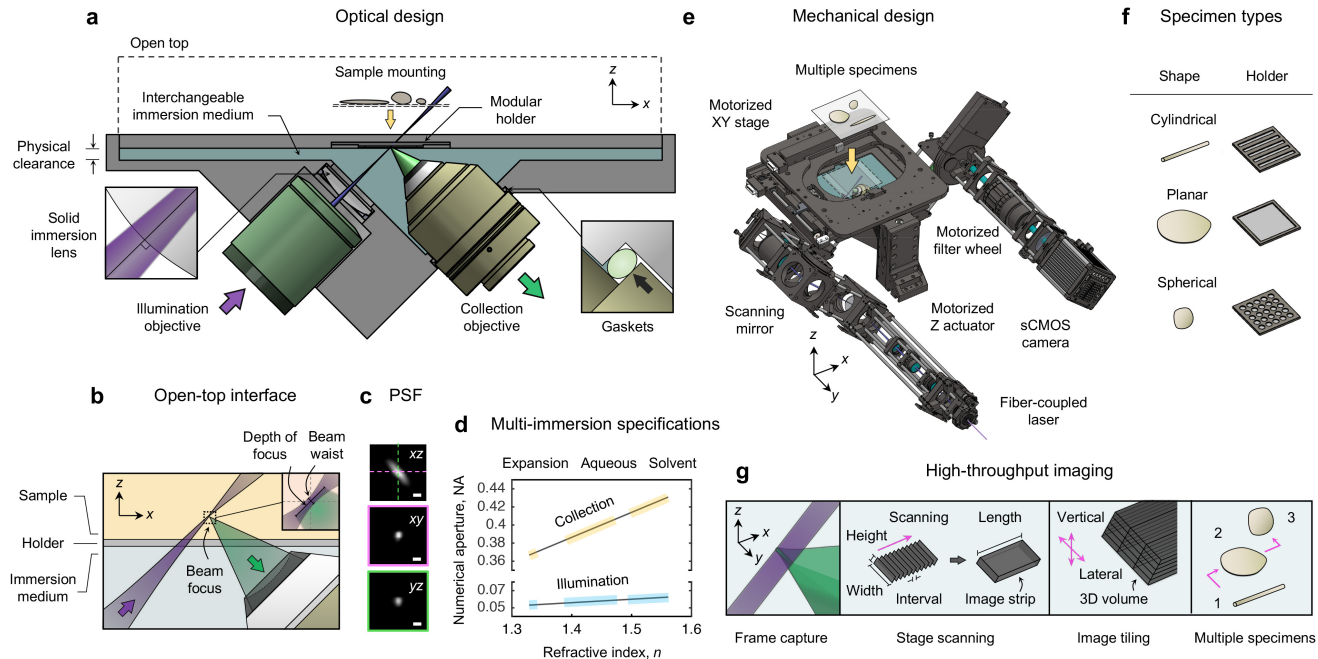
Main Figures

Figure 1. Multi-immersion open-top light-sheet (OTLS) microscope. (a) The system enables simple mounting of multiple specimens with modular transparent holders. Illumination and collection objectives are located underneath the specimen holders and are separated by a liquid reservoir filled with an interchangeable immersion medium. **(b)** The off-axis illumination light sheet and collected fluorescence travel obliquely through the immersion media, holder, and specimen. Aberrations are minimized by precisely matching the refractive index of all three materials, and by utilizing the wavefont-matching properties of a solid immersion lens (SIL) along the illumination path. The depth-of-focus and beam waist of the light sheet are depicted in the inset (upper right). The point spread function (PSF) of the system and refractive-index-dependent numerical aperture (NA) of the illumination and collection beams are shown in **(c)** and **(d)**. **(e)** The mechanical design of the system includes a motorized XY stage, motorized Z actuators, motorized filter wheel, scanning mirror, computer-controlled multi-wavelength fibre-coupled laser package, and sCMOS camera, all of which enable high-throughput automated imaging of multiple specimens simply placed on a flat plate, or placed within a diverse assortment of transparent holder designs **(f)**. **(g)** Volumetric imaging is achieved by using a combination of stage-scanning and lateral/vertical tiling. The scale bars in **(c)** denote 1 μm.
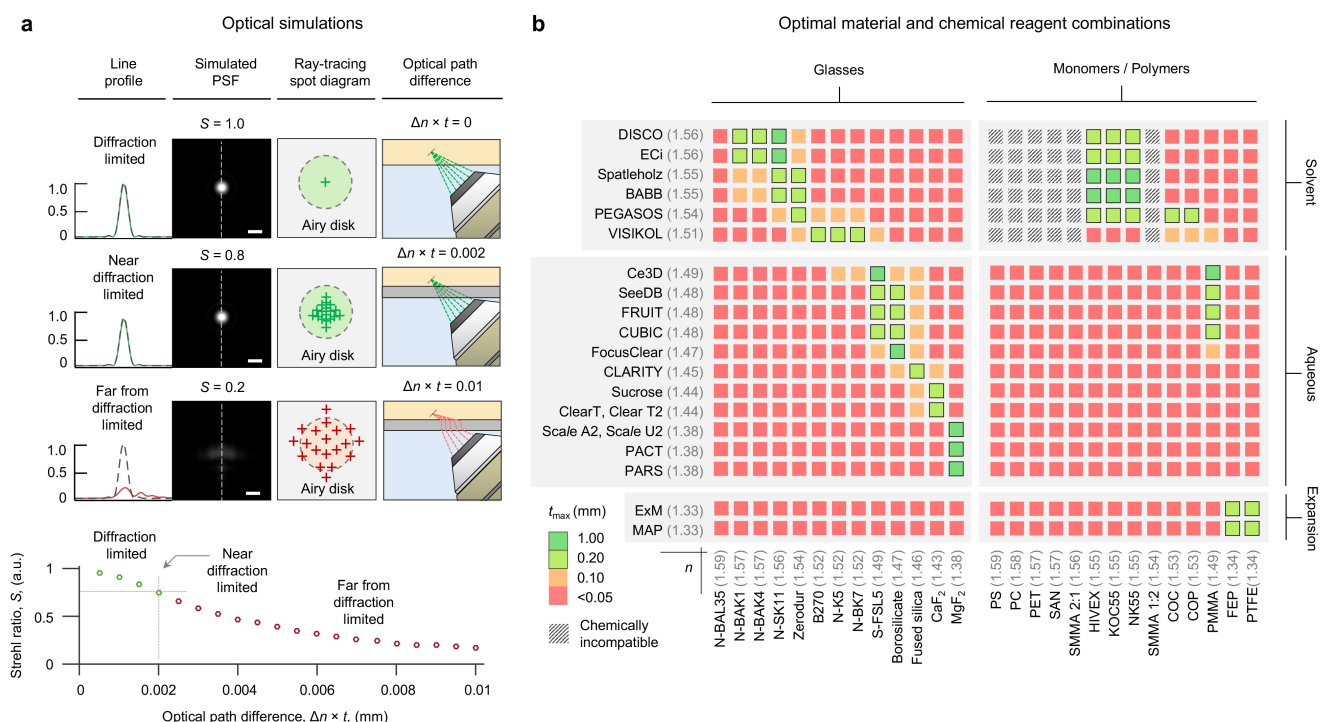
Figure 2. Holder design for OTLS imaging. (a) Optical simulations of the system's PSF, and ray-tracing spot diagrams, are shown for scenarios in which the optical path difference ($\Delta n \times t$) is negligible, small, or large, which results in diffraction-limited ($S \sim 1.0$), near-diffraction-limited ($S > 0.8$), or aberrated ($S < 0.8$) imaging performance, respectively. The dependence of the Strehl Ratio, S, as a function of $\Delta n \times t$ is plotted, indicating that for diffraction-limited imaging, the condition that $\Delta n \times t < 0.002$ should be maintained. Based on this condition, potential glass and monomer/polymer holder materials are shown in **(b)**. The color scale indicates the maximum material thickness, t_{max}, that is allowed based upon the intrinsic mismatch, Δn, of those materials with published clearing protocols. Chemically incompatible combinations of materials and chemical reagents are also indicated.
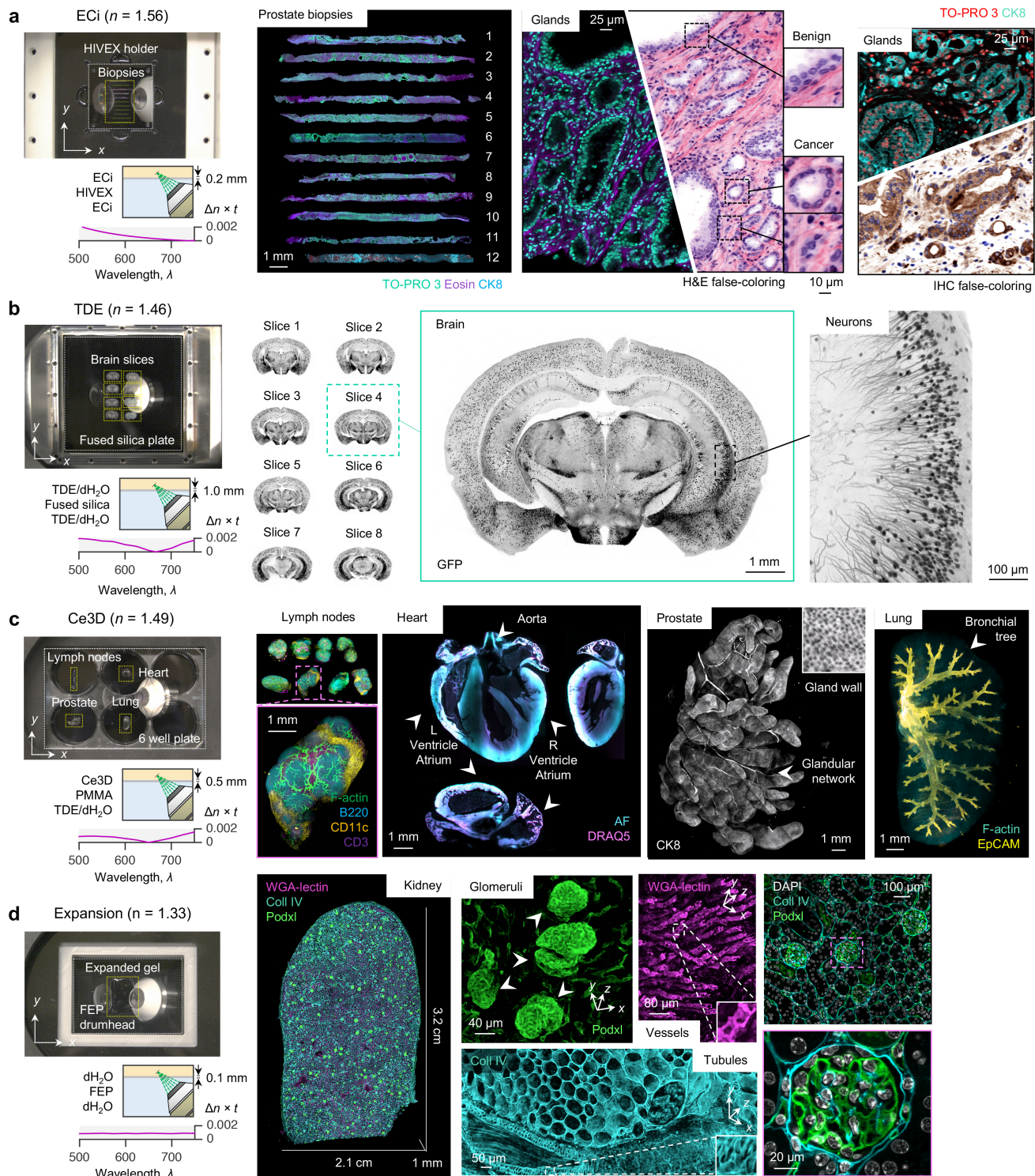
167

168 **Figure 3. Multi-immersion and multi-sample imaging. (a)** OTLS imaging of 12 ECi-cleared human
169 prostate biopsies placed within a multi-biopsy holder. Zoomed-in views illustrate the complex 3D
170 structure of benign and malignant glands. Images can be false-colored to mimic the appearance of
171 conventional chromogen-based (absorption-based) H&E and IHC histopathology. **(b)** High-throughput
172 imaging of 8 TDE-cleared mouse brain slices placed on a 10- by 10-cm glass plate. A higher-
173 magnification region of interest demonstrates the ability to visualize individual neurons. **(c)** Whole-
174 organ imaging of Ce3D-cleared mouse lymph nodes, heart, prostate, and lung placed within an multi-

175   well plate. **(d)** Large-scale imaging of an expanded thick-kidney slice placed on a teflon drumhead.
176   High-resolution regions of interest show individual glomeruli, vessels, and tubules.

177 ## Online Methods

178 *Multi-immersion open-top light-sheet microscope*

179 An optical schematic of the system is shown in **Supplementary Figure 4** and was modeled using

180 commercially available ray-tracing software (ZEMAX LLC) (**Supplementary Figs. 5** and **6**, available

181 as **Supplementary ZEMAX Files**). Illumination light is coupled into the system by a single-mode fiber

182 with a numerical aperture of 0.12 from a four-channel digitally controlled laser package (Skyra, Cobolt

183 Lasers). Light emanating from the fiber is collimated with a lens, L1 (AC1-128-019-A, f = 19 mm), and

184 then expanded along one axis using a 3× cylindrical telescope consisting of lenses, C1(ACY-254-50-A,

185 Thorlabs, f = 50 mm) and C2 (ACY-254-150-A, Thorlabs, f = 150 mm) to provide multi-directional

186 illumination (**Supplementary Figure 7**) [1]. The resulting elliptical Gaussian beam is then relayed to

187 the scanning galvanometer, GM (6210H, Cambridge Technology) using lenses R1 (AC-254-100-A,

188 Thorlabs, f = 100 mm) and R2 (AC-254-050-A, Thorlabs, f = 50 mm). The scanning mirror is driven by

189 a sinusoidal voltage from a waveform generator (PCI-6115, National Instruments) at a frequency of 800

190 Hz. The scanning beam is relayed to the back focal plan of the illumination objective

191 (XLFLUOR340/4× 0.28 NA, Olympus) using a scan lens, SL (CLS-SL, Thorlabs, f = 70 mm) and tube

192 lens, TL1 (TTL200, Thorlabs, f = 200 mm). Finally, the elliptical beam travels through a plano-convex

193 lens (LA4725, Thorlabs, R = 34.5 mm), immersion medium, holder, and finally specimen.

194   Fluorescence is collected by a multi-immersion objective (#54-10-12, Special Optics, distributed

195 by Applied Scientific Instrumentation). This provides <1 μm in-plane resolution for all immersion media

196 (**Supplementary Figure 8**). The fluorescence is filtered with a motorized filter wheel (FW102C,

197 Thorlabs) with band-pass filters for the 405 nm (FF02-447/60-25, Semrock), 488 nm (FF03-525/50-25,

198 Semrock), 561 nm (FF01-618/50-25, Semrock), and 638 nm (FF01-721/65-25, Semrock) excitation

199 wavelengths. The filtered fluorescence is focused onto a 2048×2048 pixel sCMOS camera (ORCA-

200 Flash4.0 V2, Hamamatsu) by a tube lens, TL2 (TTL165, Thorlabs, f = 165 mm). The tube lens provides

201 a Nyquist sampling of ~0.45 μm/pixel, which provides a horizontal field of view of ~0.9 mm over the

202 2048 pixels of the camera. The vertical field of view is reduced to 256 pixels to match the depth of focus

203   of the illumination light sheet (~110 μm). The 256 pixels are oriented parallel to the rolling shutter

204   readout direction of the camera, which provides an exposure time of 1.25 ms and a framerate of 800 Hz.

205   The maximum imaging depth is limited by the physical clearance of the holder and collection objective

206   (0.5 cm). The illumination objective, solid immersion lens, and collection objective interface with the

207   immersion chamber through customized aluminum mounts (**Supplementary Figure 9**) which are

208   available as **Supplementary CAD Files**.

209   Image strips are collected with a combination of stage-scanning and lateral/vertical tiling using a

210   motorized XY stage and Z actuators (FTP-2050-XYZ, Applied Scientific Instrumentation - ASI). The

211   stage-scanning firmware is used to send a TTL trigger signal from the XY stage to the sCMOS camera

212   for reproducible start positioning (<1 μm) of each image strip (**Supplementary Figure 10**). The spatial

213   interval between successive frames is set to ~0.32 μm, which, given the 800 Hz camera framerate,

214   corresponds to a constant stage velocity of ~0.25 mm/sec. For lateral tiling, an offset of 0.8 mm between

215   adjacent image strips is used (~11% overlap). For vertical tiling, the 110 μm depth of focus is oriented at

216   45 deg., which corresponds to an image strip height of ~80 μm. Therefore, a vertical tiling offset of 70

217   μm is used (~12% overlap). The laser power is increased with depth per a user defined attenuation

218   coefficient, $P = P_0 \times exp(z/\mu)$, to account for the attenuation of the illumination light sheet as it penetrates

219   deeper into the specimen. The entire image acquisition is controlled by a custom LabVIEW (National

220   Instruments) program. As shown in **Supplementary Figure 11**, the program consists of a series of

221   nested loops for imaging multiple specimens, collecting multiple color channels, and lateral/vertical

222   tiling. A complete list of components is available in **Supplementary Table 1**.

223   *Computer hardware*

224   During acquisition, the images are collected at the maximum data-transfer rate (~800 MB/sec) by a

225   dedicated workstation (Precision Tower 5810, Dell) equipped with a CameraLink interface (Firebird

226   PCI Express, Active Silicon). The data is streamed in real-time using the proprietary DCIMG

227   Hamamatsu format to a mapped network drive located on an in-lab server (X11-DPG-QT, SuperMicro)

228   running 64-bit Windows Server, equipped with 384 GB RAM and TitanXP (NVIDIA) and Quadro

229   P6000 (NVIDIA) GPUs. The server contains two high-speed RAID0 storage arrays of 4×2.0 TB SSDs,

230   as well as a larger direct-attached RAID6 storage array with 15×8.0 TB HDDs. All RAID arrays are

231   hardware-based, the RAID0 arrays are controlled by an internal 8-port controller (LSI MegaRaid 9361-

232   8i 1 GB cache) and the RAID6 array is controlled by an external 8-port controller (LSI MegaRaid 9380-

233   8e 1 GB cache). Both the server and acquisition workstation are equipped with 10G SFP+ network

234   cards, jumbo frames, and parallel send/receive processes matched to the number of computing cores on

235   the workstation (8 physical cores) and server (16 physical cores), which reliably enables >1.0 GB/sec

236   network transfer speeds (to accommodate the data-transfer rate of the sCMOS camera and enable

237   simultaneous data-processing routines). The hardware setup is shown in **Supplementary Figure 12**. The

238   complete hardware configuration is listed in **Supplementary Table 2**.

239   *Data processing and visualization*

240   Collected datasets undergo a Python pre-processing routine before being visualized in 2D and 3D by

241   several open-source and commercial packages. Each image strip is stored in a single DCIMG file. These

242   DCIMG files are read into RAM by a DLL compiled using the Hamamatsu DCIMG software

243   development kit (SDK) and first de-skewed at 45 deg. By precisely setting the interval between

244   successive frames, the de-skewing is quickly performed by simply shifting each plane of pixels in the

245   image strip by an integer pixel offset (**Supplementary Figure 13**). This operation is extremely fast

246   compared to alternative de-skewing approaches using computationally expensive affine transformations.

247   The data is then written from RAM to disk using the Hierarchical Data Format (HDF5) with the

248   metadata and XML file structured for subsequent analysis using BigStitcher [2]. A custom HDF5

249   compression filter (B3D) is used with default parameters to provide ~10× compression which is within

250   the noise limit of the sCMOS camera [3]. This pre-processing routine is applied to all DCIMG files,

251   ultimately resulting in a single HDF5/XML file for BigStitcher. The alignment of all image strips is

252   performed in BigStitcher, and finally fused to disk in either TIFF or HDF5 file formats. The resulting

253   TIFF and HDF5 files are then visualized using open-source and commercial packages, including

254   ImageJ, BigDataViewer, Aivia (DRVision), and Imaris (Bitplane) [4, 5]. To optionally provide false-

255   colored pseudo-H&E histology images, a Beer-Lambert coloring algorithm is applied using a Python

256   script [6]. The entire processing pipeline is shown in **Supplementary Figure 14** and available as

257   **Supplementary Code**.

258   *Specimen holders*

259   All holders were attached to the motorized XY stage using custom machined aluminum adapters plates

260   (HILLTOP21). For the mouse brain slices, a 1-mm thick fused silica window (Esco Optics) with a

261   10×10-cm cross-section was attached to a custom adapter plate using UV-curing glue (**Supplementary**

262   **Figure 15**). Mouse organs cleared using Ce3D were imaged on a customized 6-well plate. The bottom

263   of a conventional polystyrene 6-well plate (Cat:CLS3506, Sigma-Aldrich) was removed and replaced

264   with a 0.5 mm thick PMMA plate (Goodfellow USA) (**Supplementary Figure 16**). For the expanded

265   kidney specimen, a custom "drumhead" was fabricated and adapted for mounting to the microscope. The

266   drumhead tightens a 0.1 mm thick FEP film over an extruded opening, which is ideal for OTLS imaging

267   of expanded specimens (**Supplementary Figure 17**). To overcome the hydrophobic nature of the FEP

268   films (which cause drifting of expanded specimens), the upper surface of the FEP films were treated

269   with 0.1% (w/v) poly-lysine (Cat:P8920, Sigma-Aldrich) for charged-based adhesion of specimens to

270   the FEP surface. For the human prostate biopsies, HIVEX lens blanks (Conant Optical) were purchased

271   and custom machined using an in-house desktop mill (OtherMill, Bantam Tools) (**Supplementary**

272   **Figure 18**). The 1/8 inch, 1/16 inch, and 1/32 inch drill bits were used, and the feed rates and drill

273   speeds were optimized for the HIVEX material. CAD files for all sample holders are available as

274   **Supplementary CAD Files**. The system can also be used as a whole-slide scanner for conventional

275   fluorescently labeled histology slides using a commercially available slide holder (MLS203-P2,

276   Thorlabs) (**Supplementary Figure 19**). Dispersion curves for the various holder materials and clearing

277   reagent combinations are shown in **Supplementary Figure 20**.

278 *Optical simulations*

279 Optical simulations were performed using commercially available ray-tracing software (ZEMAX, LLC)

280 with a "blackbox" model of the multi-immersion objective (provided by the manufacturer, Special

281 Optics). For the simulations shown in **Fig. 2**, the base refractive index of the immersion medium and

282 specimen was assumed to be $n = 1.45$, and the optical path difference was varied. For all scenarios, the

283 imaging depth was set to 1 mm, and the PSF was measured at the center of the imaging field of view.

284 The same relationship between Strehl Ratio and optical path difference was observed for other base

285 refractive-indices and imaging depths, under the assumption that the optical properties of the immersion

286 medium and specimen were the same. The ZEMAX files for the OTLS system are available as

287 **Supplementary ZEMAX Files**.

288 *Collection and processing of mouse brain slices*

289 A mouse of line Sst-IRES-Cre;Ai139(TIT2L-GFP-ICL-TPT), characterized previously [7] was used for

290 imaging experiments. Genotyping confirmed expression of Cre and tdTomato for this individual. The

291 mouse was sacrificed at age P96 by trans-cardial perfusion with 4% paraformaldehyde. The brain was

292 dissected and post-fixed in 4% paraformaldehyde at room temperature for 3-6 hr followed by overnight

293 fixation at 4 deg. C. The brain was rinsed with $1 \times$ PBS and stored in $1 \times$ PBS with 0.1% sodium azide

294 prior (Cat:S2002, Sigma-Aldrich) prior to sectioning. 200-µm thick cortical sections were cut on a

295 vibratome and stored in $1 \times$ PBS. Prior to OTLS imaging, brain slices were incubated in a mixture of

296 68% 2,2′-thiodiethanol (TDE) (Cat:166782, Sigma-Aldrich) and 32% $1 \times$ PBS for clearing. The

297 refractive index of the solution ($n \sim 1.46$) was verified using a refractometer (PA202, Misco).

298 Procedures involving mice were approved by the Institutional Animal Care and Use Committee of the

299 Allen Institute for Brain Science in accordance with NIH guidelines.

300 *Collection and processing of heart, lung, prostate, and lymph nodes*

301 Lung, heart, prostate, and lymph nodes were collected from a CD11-YFP, Actin-dsRed expressing

302 mouse. Tissues were fixed for 24 hr at 4 deg. C in 1 part fixative (Cat:554655, BD Biosciences) and 2

303   parts 1× PBS and incubated in blocking buffer for 24 hr at 37 deg. C. The buffer consisted of 30 mL

304   Tris (Cat:252859, Sigma-Aldrich), 0.3 mL NMS (Cat:SML1128, Sigma-Aldrich), 0.3 mL BSA

305   (Cat:A2058, Sigma-Aldrich), and 0.09 mL TritonX100 (Cat:T8787, Sigma-Aldrich). Lymph nodes were

306   stained for 4 days at 37 deg. C in 400 μL blocking buffer, 2 μL CD3-BV421 (Cat: 100228, BioLegend),

307   and 2 μL B220-e660 (Cat: 50-0452-82, ThermoFisher). Lung tissue was stained for 3 days at 37 deg. C

308   in 500 μL blocking buffer and 2.5 μL Epcam-APC (Cat: 17-5791-82, ThermoFisher). Heart tissue was

309   stained for 1 day with 1 mM DRAQ5. Prostate tissue was incubated with fluorophore-conjugated anti-

310   CK8-18 (Cat:MS743S0, Fisher) conjugated to Alexa-Fluor 488 (Cat:A20181, Invitrogen) (1:100

311   dilution) in PBS/1% non-fat dry milk/0.2% Triton X-100 at 37 deg. C for 7 days with gentle agitation.

312   All tissues were then cleared with the Ce3D solution, consisting of 14 mL of 40% N-methyl-acetamide

313   (Cat:M26305, Sigma-Aldrich), 25 μL Triton X-100 (Cat:T8787, Sigma-Aldrich), 20 g Histodenz

314   (Cat:D2158, Sigma-Aldrich), and 125 μL Thioglycerol (Cat:88640, Sigma-Aldrich) for 1 day at room

315   temperature. Procedures involving mice were approved by the Institutional Animal Care and Use

316   Committee of the University of Washington in accordance with NIH guidelines.

317   *Collection and processing of expanded mouse kidney*

318   4% PFA fixed mouse kidney was sliced to 200 μm and processed using a previously described protocol

319   [8]. The tissue was then incubated in blocking/permeabilization buffer for 6 hr at 4 deg. C. Primary

320   antibodies goat anti-podocalyxin (cat: AF1556, R&D Sys. Inc., 1:50) and rabbit anti-collagen IV (cat:

321   ab6586, abcam, 1:50) were diluted with blocking/permeabilization buffer and used to stain the tissue for

322   2 days at 4 deg. C. The tissue was then washed with 1× PBS three times at room temperature (1 hr

323   each). Fluorescently-labeled secondary antibodies, Alexa 488 conjugated WGA (cat: W11261, Thermo

324   Fisher Scientific, 1:25), and Hoechst 33342 were then diluted in blocking/permeabilization buffer to

325   stain the tissue for 2 days at 4 deg. C. The tissue was washed with 1xPBS three times at room

326   temperature (1 hr each) followed by incubating in 1 mM MA-NHS (cat:730300, Sigma-Aldrich) for 1 hr

327   at room temperature. The tissue was then incubated in monomer solution for 1 hr at 4 deg. C and then

328   gelled in a humidified environment at 37 deg. C for 2 hr. Excess gel was removed and the specimen was

329   digested by proteinase K (cat: EO0491, Thermo Fisher Scientific) at 37 deg. C for two days and then

330   collagenase (cat; C7926, Sigma-Aldrich) at 37 deg. C for two days refreshing the solution daily. After

331   digestion, the specimen was incubated in DI water for at least 2 hr and the expansion factor was

332   determined through measuring the dimensions of the gel. The expanded specimen was mounted on poly-

333   lysine coated film for imaging.

334   *Collection and processing of human prostate biopsies*

335   All specimens were obtained from an IRB-approved genitourinary biorepository with patient consent.

336   Core-needle biopsy specimens were obtained from fresh *ex vivo* prostatectomy specimens using an 18-

337   gauge (approximately 1 mm inner diameter) needle biopsy device (Bard Max Core, Bard Biopsy). The

338   biopsy was immediately placed in 10% neutral buffered formalin, where it was maintained at room

339   temperature for 24 hrs. In contrast to mouse tissues, we found that human tissues require more

340   aggressive solvent-clearing approaches (**Supplementary Figure 21**). Due to its clearing efficacy and

341   non-toxic nature, we used ECi-clearing, which we observed does not interfere with downstream

342   histology or immunohistochemistry (**Supplementary Figure 22**).

343   Biopsies were then washed in $1 \times$ PBS with 0.1% Triton X-100 (Cat:T8787, Sigma-Aldrich), and

344   each biopsy was stained for 4 hr in a 1:2000 dilution of TO-PRO3 Iodide (Cat:T3605, Thermo-Fischer)

345   at room temperature with light shaking. Each biopsy was then dehydrated in ethanol for with 25/75,

346   50/50, 75/25, and 100/0 grades. The dehydration time for each grade was 1 hr, and the 100% ethanol

347   grade was performed twice to ensure removal of any excess water. Biopsies were then stained in 1:2000

348   dilution of Eosin-Y (Cat:3801615, Leica Biosystems) for 4 hr at room temperature with light shaking.

349   Finally, biopsies were cleared in ethyl-cinnamate (Cat:112372, Sigma-Aldrich) for 1 hr. Biopsy #12 was

350   stained with anti-CK8. The biopsy issue was incubated simultaneously with fluorophore-conjugated

351   anti-CK8-18 (Cat:MS743S0, Fisher) conjugated to Alexa-Fluor 488 (Cat:A20181, Invitrogen) (1:100

352   dilution) in PBS/1% non-fat dry milk/0.2% Triton X-100 at 37 deg. C for 7 days with gentle agitation.

Data Availability

All raw and processed data generated in this work, including the images provided in the manuscript and supplementary material are available from the authors upon request. The customized CAD and ZEMAX files are available as **Supplementary CAD Files** and **Supplementary ZEMAX Files**.

Code Availability

The computer code used to acquire, process, and generate the images in this study is available as **Supplementary Code Files**. **Supplementary Data** and instructions for installing and using the computer code are available at https://figshare.com/articles/Supplementary_Data/7685597.

Acknowledgements

375 **Contributions**

376 A.G., N.R., P.N., M.G., J.V., L.T. and J.L. designed the studies. A.G. and J.L. designed the multi-

377 immersion open-top light-sheet microscope. J.D. designed the multi-immersion collection objective and

378 provided input on the microscope design. A.G., Y.C., C.Y, L.B., W.X., and L.W fabricated the

379 microscope. A.G. and P.N. prepared and imaged the mouse brain slices. C.S., A.G., and E.M. prepared

380 and imaged the mouse organs. A.H., C.M., and J.V. prepared the mouse kidney tissues. A.G., A.H., and

381 C.M. imaged the expanded mouse kidney tissues. N.R., S.K., L.T., and E.M. prepared the human

382 prostate tissues. A.G., S.K., and N.R. imaged the human prostate tissues. E.M. prepared all downstream

383 histology for the study. N.R. and L.T. histologically characterized all human prostate tissues. All authors

384 prepared the manuscript.

385 **Competing Interests**

386 A.G., N.R., L.T., and J.L. are co-founders and shareholders of Lightspeed Microscopy Inc.

387 **Corresponding Author**

388 Correspondence to Adam K. Glaser or Jonathan T.C. Liu

References

1. Richardson, D.S. and J.W. Lichtman, *Clarifying Tissue Clearing.* Cell, 2015. **162**(2): p. 246-57.

2. Chung, K., et al., *Structural and molecular interrogation of intact biological systems.* Nature, 2013. **497**(7449): p. 332-7.

3. Tanaka, N., et al., *Whole-tissue biopsy phenotyping of three-dimensional tumours reveals patterns of cancer heterogeneity.* Nature Biomedical Engineering, 2017. **1**(10): p. 796.

4. Susaki, E.A., et al., *Advanced CUBIC protocols for whole-brain and whole-body clearing and imaging.* Nature Protocols, 2015. **10**: p. 1709-1727.

5. Glaser, A.K., et al., *Light-sheet microscopy for slide-free non-destructive pathology of large clinical specimens.* Nature Biomedical Engineering, 2017. **1**(7): p. 0084.

6. Power, R.M. and J. Huisken, *A guide to light-sheet fluorescence microscopy for multiscale imaging.* Nature Methods, 2017. **14**(4): p. 360.

7. Chen, B.-C., et al., *Lattice light-sheet microscopy: Imaging molecules to embryos at high spatiotemporal resolution.* Science, 2014. **346**(6208): p. 1257998.

8. Huisken, J., et al., *Optical sectioning deep inside live embryos by selective plane illumination microscopy.* Science, 2004. **305**(5686): p. 1007-9.

9. Dodt, H.-U., et al., *Ultramicroscopy: three-dimensional visualization of neuronal networks in the whole mouse brain.* Nature Methods, 2007. **4**(4): p. 331-336.

10. Tomer, R., et al., *Advanced CLARITY for rapid and high-resolution imaging of intact tissues.* Nature Protocols, 2014. **9**(7): p. 1682-97.

11. Migliori, B., et al., *Light sheet theta microscopy for rapid high-resolution imaging of large biological samples.* BMC Biology, 2018. **16**(1): p. 57.

12. McGorty, R., et al., *Open-top selective plane illumination microscope for conventionally mounted specimens.* Optics Express, Vol. 23, Issue 12, pp. 16142-16153, 2015.

13. Strnad, P., et al., *Inverted light-sheet microscope for imaging mouse pre-implantation development.* Nature Methods, 2015. **13**: p. 139-142.

14. Renier, N., et al., *iDISCO: a simple, rapid method to immunolabel large tissue samples for volume imaging.* Cell, 2014. **159**(4): p. 896-910.

15. Becker, K., et al., *Chemical Clearing and Dehydration of GFP Expressing Mouse Brains.* PLoS One, 2012. **7**(3): p. e33916.

16. Klingberg, A., et al., *Fully Automated Evaluation of Total Glomerular Number and Capillary Tuft Size in Nephritic Kidneys Using Lightsheet Microscopy.* Journal of the American Society of Nephrology, 2016. **28**(2): p. 452-459.

17. Ertürk, A., et al., *Three-dimensional imaging of solvent-cleared organs using 3DISCO.* Nature Protocols, 2012. **7**: p. 1983-1995.

18. Pan, C., et al., *Shrinkage-mediated imaging of entire organs and organisms using uDISCO.* Nature Methods, 2016. **13**: p. 859-867.

19. Klingberg, A., et al., *Fully Automated Evaluation of Total Glomerular Number and Capillary Tuft Size in Nephritic Kidneys Using Lightsheet Microscopy.* 2016.

20. Nojima, S., et al., *CUBIC pathology: three-dimensional imaging for pathological diagnosis.* Scientific Reports, 2017. **7**(1): p. 9269.

21. Tainaka, K., et al., *Chemical Landscape for Tissue Clearing Based on Hydrophilic Reagents.* Cell Reports, 2018. **24**(8): p. 2196-2210.e9.

22. Chung, K. and K. Deisseroth, *CLARITY for mapping the nervous system.* Nature Methods, 2013. **10**: p. 508-513.

23. Li, W., R.N. Germain, and M.Y. Gerner, *Multiplex, quantitative cellular analysis in large tissue volumes with clearing-enhanced 3D microscopy (Ce3D).* Proceedings of the National Academy of Sciences of the United States of America, 2017. **114**(35): p. E7321-e7330.

24. Chen, F., P.W. Tillberg, and E.S. Boyden, *Optical imaging. Expansion microscopy.* Science, 2015. **347**(6221): p. 543-8.

25. Chozinski, T.J., et al., *Expansion microscopy with conventional antibodies and fluorescent proteins.* Nature Methods, 2016. **13**(6): p. 485-8.

26. Ku, T., et al., *Multiplexed and scalable super-resolution imaging of three-dimensional protein localization in size-adjustable tissues.* Nature Biotechnology, 2016. **34**(9): p. 973.

27. Park, Y.G., et al., *Protection of tissue physicochemical properties using polyfunctional crosslinkers.* Nature Biotechnology, 2018.

28. Goltsev, Y., et al., *Deep Profiling of Mouse Splenic Architecture with CODEX Multiplexed Imaging.* Cell, 2018. **174**(4): p. 968-981.e15.

29. Cadwell, C.R., et al., *Multimodal profiling of single-cell morphology, electrophysiology, and gene expression using Patch-seq.* Nature Protocols, 2017. **12**(12): p. 2531.

30. Wang, X., et al., *Three-dimensional intact-tissue sequencing of single-cell transcriptional states.* Science, 2018. **361**(6400).

31. Glaser, A.K., et al., *Multidirectional digital scanned light-sheet microscopy enables uniform fluorescence excitation and contrast-enhanced imaging.* Scientific Reports, 2018. **8**(1): p. 13878.

32. Hörl, D., et al., *BigStitcher: Reconstructing high-resolution image datasets of cleared and expanded samples.* bioRxiv, 2018.

33. Balazs, B., et al., *A real-time compression library for microscopy images.* bioRxiv, 2017.

34. Pietzsch, T., et al., *BigDataViewer: visualization and processing for large image data sets.* Nature Methods, 2015. **12**(6): p. 481-3.

35. Schneider, C.A., W.S. Rasband, and K.W. Eliceiri, *NIH Image to ImageJ: 25 years of image analysis.* Nature Methods, 2012. **9**(7): p. 671.

460   36.   Giacomelli, M.G., et al., *Virtual Hematoxylin and Eosin Transillumination Microscopy Using*
461   *Epi-Fluorescence Imaging.* PLoS One, 2016. **11**(8): p. e0159337.

462   37.   Daigle, T.L., et al., *A Suite of Transgenic Driver and Reporter Mouse Lines with Enhanced*
463   *Brain-Cell-Type Targeting and Functionality.* Cell, 2018. **174**(2): p. 465-480.e22.

464   38.   Chozinski, T.J., et al., *Volumetric, Nanoscale Optical Imaging of Mouse and Human Kidney via*
465   *Expansion Microscopy.* Scientific Reports, 2018. **8**(1): p. 10396.

466