

1 **Automated High-Throughput Light-Sheet Fluorescence**

2 **Microscopy of Larval Zebrafish**

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13 **ABSTRACT**

14 Light sheet fluorescence microscopy enables fast, minimally phototoxic, three-dimensional
15 imaging of live specimens, but is currently limited by low throughput and tedious sample
16 preparation. Here, we describe an automated high-throughput light sheet fluorescence
17 microscope in which specimens are positioned by and imaged within a fluidic system
18 integrated with the sheet excitation and detection optics. We demonstrate the ability of the
19 instrument to rapidly examine live specimens with minimal manual intervention by imaging
20 fluorescent neutrophils over a nearly 0.3 mm³ volume in dozens of larval zebrafish. In
21 addition to revealing considerable inter-individual variability in neutrophil number, known
22 previously from labor-intensive methods, three-dimensional imaging allows assessment of
23 the correlation between the bulk measure of total cellular fluorescence and the spatially
24 resolved measure of actual neutrophil number per animal. We suggest that our simple
25 experimental design should considerably expand the scope and impact of light sheet imaging
26 in the life sciences.

27

28 **INTRODUCTION**

29 Light sheet fluorescence microscopy (LSFM) is a powerful tool for examining the three-
30 dimensional structural and temporal dynamics of living systems. In LSFM, a thin sheet of
31 laser light excites fluorophores in a sample. Scanning the sample through the sheet enables
32 fast, three-dimensional imaging with low phototoxicity, high resolution, and a wide field of
33 view (1–5). Imaging with LSFM has enabled numerous studies of embryonic development
34 (6,7), neural activity (8), microbial dynamics (9–11), and other phenomena. A large body of
35 work has focused on improving the optical capabilities of light sheet imaging, for example
36 using structured illumination (12,13), multiple lens pairs (6), two-photon excitation (14), and

37 other techniques. However, current light sheet fluorescence microscopes have significant
38 constraints related to throughput and sample handling that, we argue, have placed much
39 greater limitations on their scientific utility than issues of spatial or temporal resolution. The
40 majority of existing light sheet fluorescence microscopes, both commercial and non-
41 commercial, are designed to hold a single specimen. A few instruments (including one from
42 the authors of this paper) can hold up to six specimens for sequential imaging. Moreover,
43 light sheet fluorescence microscopy typically requires extensive sample preparation and
44 manual sample mounting, most commonly by embedding specimens in agarose gels.
45 Examination of large numbers of specimens is therefore slow and difficult, which is
46 especially important given the high level of inter-individual variability found in many
47 complex biological processes. Increasing the pace of insights into developmental biology,
48 multicellular biophysics, or microbial community structure will require faster and simpler
49 acquisition of three-dimensional imaging datasets. To date, there exists only one report of an
50 automated light sheet microscope that makes use of fluidic positioning of live animals (15);
51 its throughput (specimens per hour) is not stated, and though its ability to image larval
52 zebrafish is clear, the total number of animals examined was only twelve. In contrast,
53 automated, high-throughput methods have been integrated with other types of microscopes,
54 including confocal microscopes (16–18). Our system adopts and builds upon these,
55 especially the confocal-based setup of Ref. 16.

56

57 To address the issues described above, we developed a light sheet fluorescence microscope
58 capable of automated, high-throughput imaging of live specimens. Our instrument uses
59 fluidic control and image-based registration to rapidly but precisely position specimens for
60 light sheet scans and subsequently remove them from the imaging area. We characterize the

61 optical quality of our instrument, and demonstrate its capabilities by rapidly imaging immune
62 cells in dozens of larval zebrafish. While the spatial resolution of our microscope does not
63 equal that of current “low-throughput” light sheet microscopes, it is more than sufficient for
64 determining cellular distributions. Moreover, we argue that the tradeoff of lower resolution
65 for higher throughput is worthwhile given the large variance in most biological datasets. We
66 find, for example, a high degree of variation between fish in total neutrophil number, and
67 clustering of neutrophils in two distinct regions near the swim bladder. While our instrument
68 is optimized for imaging of larval zebrafish, the design could easily be modified for rapid
69 imaging of a wide range of biological and non-biological samples, which should broaden the
70 impact of light sheet microscopy in a variety of fields.

71

72 **RESULTS**

73 **Instrument design**

74 The light sheet portion of the microscope closely follows the design of Keller et al (1); a
75 rapidly-scanned galvanometer creates a sheet of light for fluorescence excitation, and emitted
76 light is captured by a camera perpendicular to the plane of the sheet (Fig. 1A). In
77 conventional light sheet fluorescence microscopes, gel-mounted specimens are introduced
78 vertically in between horizontal lenses. To achieve high throughput, we use a continuous
79 fluidic path through plastic tubing and glass capillaries for transport as well as imaging,
80 detailed below. If the fluidic path were oriented vertically, specimens would gravitationally
81 drift during imaging. Therefore, we adopted a geometry in which specimens are transported
82 horizontally and the sheet plane is vertical (Fig. 1A,B). To allow this arrangement, we
83 designed an elongated sample chamber with windows oriented below and perpendicular to
84 the sample (Fig. 1B,C). Before entering the imaging chamber, specimens flow through a

85 system of 0.7 mm diameter plastic tubing at a typical flow rate of 1 ml/min, or 4 cm/sec
86 (Fig. 1). Flow speed and direction are controlled by a syringe pump (Fig. 1); see
87 Supplemental Methods for a parts list and descriptions.

88

89 **Figure 1. Instrument design.** (A) Schematic of the instrument design, with
90 labels corresponding to the parts list in Table 1. See also Supplementary
91 Movie 1. The excitation laser line is selected by an acousto-optic tunable
92 filter (AOTF₂), then directed to a galvanometer mirror (G₃) and objective
93 lens (L₄) to create a time-averaged sheet of light in the sample chamber (C)
94 via a prism (Pr₅). Specimens flow through a system of tubing controlled by a
95 syringe pump (Pu₉) and valves (V₁₀) and are automatically positioned in a
96 square-walled capillary (Cap₁₁) for imaging. Bright field images are used for
97 positioning the sample and are illuminated with an LED (LED₆). After
98 imaging, specimens are directed into a reservoir (R). (B) Schematic of the
99 imaging area. The 3D-printed sample chamber (C), prism (Pr₅), and imaging
100 capillary (Cap₁₁) are apparent. (C) Photograph of the imaging area
101 corresponding to the schematic in (B).

102

103 Inside the imaging chamber, specimens flow into a square-walled glass capillary in front of
104 the imaging objective where they are automatically detected by bright field microscopy.
105 Specimens are rapidly stopped using computer-controlled valves on either side of the
106 imaging chamber, with a precision of approximately 1 mm in position, comparable to the
107 length of a larval zebrafish. Fine positioning is performed by iterated movement of the
108 capillary by a computer-controlled stage, brightfield imaging, and cross-correlation of images

109 with a previously assembled image library (Fig. 2 A,B). The travel range of the capillary on
110 the stage allows movement of up to 30 mm in the x-direction. Like many studies, ours make
111 use of larval zebrafish as a model organism; strong features such as eyes and the swim
112 bladder enable straightforward correlation-based registration, with an estimated precision of
113 about 20 μm . This approach should be applicable to any specimen with a roughly
114 stereotypical anatomy.

115

116 Once positioned, specimens are automatically imaged using LSFM. The imaging chamber
117 has sufficient depth, 35 mm, that the capillary can be scanned through the sheet by a
118 motorized stage. In our setup, repeated scans with up to three excitation wavelengths are
119 possible; this is limited simply by the number of available laser lines. The precision of the
120 automated positioning enables scans to be taken of particular regions, for example the larval
121 gut, as shown below. After imaging, specimens flow into a collection reservoir and
122 subsequent specimens are automatically positioned for imaging. We provide a movie of the
123 instrument in operation as Supplementary Video S1. A complete parts list is provided as
124 Table 1, in Supplemental Methods.

125

126 **Optical quality**

127 Our instrument uses glass capillaries for specimen mounting, rather than more conventional
128 gel embedding. The square cross-section of these capillaries should lead to less distortion
129 than more common cylindrical capillaries. To assess the optical quality of our setup, we
130 measured the point spread function (PSF) by imaging 28 nm diameter fluorescent
131 microspheres dispersed in oil in these capillaries (see Methods for details). The diffraction-
132 limited width of the particles in the sheet plane (xy), assessed as the standard deviation of a

133 Gaussian function fit to the particle's intensity profile, is $0.6 \mu\text{m}$, and the width along the
134 detection axis (z) is $3.4 \pm 0.6 \mu\text{m}$, consistent with the expected sheet thickness of our setup
135 (Fig. 2 C,D).

136

137 **Figure 2. Specimen positioning and image quality.** (A) Composite
138 brightfield image of a larval zebrafish positioned in a glass capillary. Scale bar:
139 $50 \mu\text{m}$. (B) Normalized intensity averaged along the short axis of the
140 brightfield image, and the intensity of the template image that best matches
141 the fish in (A). Cross-correlation with the template is used to automatically
142 position the fish for light sheet fluorescence imaging. (C) Light sheet
143 fluorescence images of a 28 nm diameter fluorescent microsphere, showing
144 x - y and z - y planes centered on the particle. (D) Line-scan of intensity along
145 the detection axis (z) through a fluorescent microsphere, with a Gaussian fit
146 showing a width of approximately $3 \mu\text{m}$.

147

148 **Data collection capabilities**

149 Using this system, we can image approximately 30 larval zebrafish per hour, obtaining from
150 each a $666 \times 431 \times 1060 \mu\text{m}$ (x, y, z) three-dimensional scan, a marked improvement over
151 manual mounting and imaging that, even by a skilled researcher, is limited to about 5 fish per
152 hour. The triggering accuracy is about 90%, with roughly 10% of detected objects being
153 bubbles or debris that are easily identified after imaging. On average, 81% of larval fish are
154 automatically positioned correctly in front of the imaging objective. The remaining 19%
155 correspond to multiple fish being in the field of view, or other positioning errors.
156 Importantly, the majority of the run time of the instrument is spent obtaining light sheet

157 fluorescence images, and is not dominated by specimen positioning. In the batch of $N=41$
158 fish whose neutrophil distributions were imaged in experiments described below, for
159 example, the flow, detection, and positioning of the specimens occupied only approximately
160 30 seconds per fish. In the limit of zero imaging time (e.g. for very bright signals or small
161 regions of interest), the system could therefore record data from up to about 120 specimens
162 per hour in the absence of triggering or positioning errors, or about 90 specimens per hour
163 with the present system performance.

164

165 **Neutrophil distributions in larval zebrafish**

166 To demonstrate the capabilities of the instrument, we imaged fluorescent neutrophils in
167 larval zebrafish at 5 days post-fertilization (dpf), focusing especially on the number of these
168 immune cells and their distribution near the anterior of the intestine. Specifically, these were
169 fish engineered to express green fluorescent protein (GFP) under the promoter
170 myeloperoxidase, an enzyme primarily produced in neutrophils (19). In total, we imaged 41
171 fish, obtaining from each a single $666 \times 431 \times 1060 \mu\text{m}$ three-dimensional image in which
172 neutrophils were readily evident (Fig. 3A and Supplementary Movie S2). The strong GFP
173 signal enabled automated identification of neutrophils by standard segmentation methods.
174 Corroborating previous work done by manual dissection of zebrafish (20), we found a high
175 degree of variation in neutrophil number between specimens, with the standard deviation
176 being 30% of the mean (Fig. 3B). Furthermore, we found that neutrophils tend to cluster in
177 two distinct regions: adjacent to the swim bladder on both the anterior and posterior (Fig.
178 3C). Notably, the total GFP intensity in a fish is weakly correlated with the number of
179 neutrophils, with a coefficient of determination $R^2 = 0.4$, indicating that a simple measure of

180 overall brightness, as could be assessed without three-dimensional microscopy, would
181 provide a poor diagnostic of the actual abundance of immune cells (Fig. 3D).

182

183 **Figure 3. Imaging neutrophils in larval zebrafish.** (A) A maximum
184 intensity projection of a three-dimensional light sheet fluorescence image of
185 GFP-expressing neutrophils near the intestine of a 5 dpf larval zebrafish. The
186 3D scan is provided as Supplemental Movie 2. The intestine and swim
187 bladder are roughly outlined by the yellow dotted lines. Scale bar: 100 μm .
188 (B) The total number of neutrophils in each fish; the x-axis is ordered by
189 neutrophil count. (C) Neutrophil count along the anterior-posterior
190 dimension, summed over all fish examined ($N=41$). The x-axis corresponds
191 approximately to the horizontal range of (A). (D) The total intensity of the
192 detected neutrohils per fish vs the total number of neutrophils in that fish.
193 The two measures are weakly correlated with a coefficient of determination
194 $R^2 = 0.4$.

195

196 **DISCUSSION**

197 While our microscope is optimized for rapid imaging of larval zebrafish, the design is general
198 and opens numerous possibilities for imaging a wide range of specimens, such as organoids,
199 drosophila embryos, small marine invertebrates, and more, with the appropriate expansion
200 of the positioning image library. While the present design provides only a single three-
201 dimensional image of each specimen, we envision future integration of a closed-loop fluidic
202 circuit, with which specimens can be automatically loaded, imaged, and circulated repeatedly,
203 allowing for high-throughput acquisition of multiple snapshots of the same specimen over

204 time. In general, tackling the challenge of automated, high-throughput specimen handling
205 will allow the technique of light sheet fluorescence microscopy to maximize its impact on
206 the life sciences.

207

208 **METHODS**

209 **Hardware**

210 The majority of the instrument was constructed with off the shelf parts. Custom parts were
211 either laser cut from acrylic sheets or were 3D printed (see Parts List).

212

213 Fluorescence excitation is provided by various solid state lasers, selected by an acousto-optic
214 tunable filter (AOTF, AA Opto-electronic) for coupling into a fiber launch to a
215 galvanometer mirror (Cambridge Technology), which oscillates with a triangular waveform
216 at 1 kHz to sweep the beam into a sheet. An objective lens (Mitutoyo 5X) and a prism route
217 the sheet to the water-filled sample chamber where it intersects the specimen (Fig. 1).
218 Detection is provided by a 20x water immersion objective (Olympus 20XW), mounted in the
219 side of the chamber and sealed with an o-ring, its corresponding tube lens, and an sCMOS
220 camera (Hamamatsu Orca Flash 4.0). Exposure times were 25 ms for all experiments
221 presented here. Instrument control software was written in MATLAB.

222

223 **Imaging Procedure**

224 For live imaging, each larval zebrafish flows through 0.7 mm inner diameter silicone tubing
225 to a 50 mm long section of round 0.7 mm inner diameter, square exterior cross section, glass
226 tubing in the specimen chamber. The contrast between the specimen and background in
227 brightfield images is used to detect the specimen, stop the pump, and close off tubing using

228 pinch valves to prevent specimen drift. The xyz stage arm holding the glass capillary is
229 scanned along the capillary's length to locate the section to be imaged, as described in the
230 main text. Following positioning, bright field illumination is switched off and the desired
231 laser wavelength is selected by the AOTF. The xyz arm is then scanned perpendicular to the
232 sheet plane, generating a 3D image stack. The scan can be repeated for another region or
233 another wavelength before the pump is directed to send the specimen out of the chamber,
234 and bring in the next specimen.

235

236 **Point Spread Function**

237 For measurements of the point spread function, 28 nm diameter fluorescent carboxylate-
238 modified polystyrene spheres (Thermo Fisher cat. #F8787), with peak excitation and
239 emission wavelengths 505 and 515 nm, respectively, were dispersed in oil with a similar
240 index of refraction as water (Zeiss Immersol W 2010) inside the imaging capillaries. Three-
241 dimensional scans were taken, with a z-spacing of 0.5 μm .

242

243 **Ethics statement**

244 All zebrafish experiments were carried out in accordance with protocols approved by the
245 University of Oregon Institutional Animal Care and Use Committee (21).

246

247 **Zebrafish husbandry**

248 The zebrafish line *Tg[BACmpo:gfp]* (19) was used for neutrophil imaging. Larval zebrafish
249 were raised at a density of one embryo per milliliter and kept at a constant temperature of 28
250 °C. Embryos were not fed during experiments and were sustained by their yolks.

251

252 **Sample preparation**

253 Larval zebrafish were placed in dishes containing sterile embryo media with 0.05%
254 methylcellulose and anaesthetized using 240 µg/ml tricaine methanesulfonate. This
255 anaesthetic concentration is higher than the standard dosage, but was necessary likely
256 because of permeation through the plastic tubing. In preparation for imaging, larval
257 zebrafish were drawn into a system of tubing with a spacing of approximately 6 inches.
258 During experiments, larvae flowed through the tubing and were automatically stopped and
259 positioned by a syringe pump and a series of valves. After imaging, larvae flowed into a dish
260 containing sterile embryo medium.

261

262 **Image-based neutrophil quantification**

263 Neutrophil segmentation was performed using an iterative approach. First, an intensity
264 threshold was applied to the image stack creating a binary mask, after which connected
265 objects were detected and objects over a minimum size were kept. Then, each object larger
266 than a specified size was re-thresholded using Otsu thresholding followed by erosion.
267 Connected objects were then detected and this process was repeated until all objects were
268 either removed or retained.

269

270 **ACKNOWLEDGEMENTS**

271 We thank Rose Sockol, Kyleah Murphy, and the University of Oregon Zebrafish Facility
272 staff for fish husbandry, and Karen Guillemin and Brandon Schlomann for useful comments
273 and conversations. Research reported in this publication was supported in part by the
274 National Science Foundation (NSF) under award 1427957 (to RP), by the the M. J. Murdock
275 Charitable Trust, and by the National Institutes of Health (NIH) as follows: by the National

276 Institute of General Medical Sciences under award number P50GM098911 and by the
277 National Institute of Child Health and Human Development under award P01HD22486,
278 which provided support for the University of Oregon Zebrafish Facility. The content is
279 solely the responsibility of the authors and does not represent the official views of the NIH,
280 NSF, or other funding agencies.

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355

356

357 **SUPPORTING INFORMATION CAPTIONS:**

358

359 **Supplemental Movie 1.** Video of the automated, high-throughput light sheet microscope in
360 operation. After a larval zebrafish, flowing through tubing, is detected and positioned, 3D
361 scans in two colors in two different regions, are obtained. The final image (0:57-1:03) is of
362 tdTomato-labeled bacteria and GFP-labeled neutrophils.

363

364 **Supplemental Movie 2.** A light sheet fluorescence scan through the anterior intestine of a
365 larval zebrafish with GFP-expressing neutrophils evident as very bright objects above
366 autofluorescent background. The total image depth is 475 μm (2.5 μm /slice x 190 slices).
367 The maximum intensity projection of this scan is shown in Figure 2E.

368





