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 threedimensional structural and temporal dynamics of living systems. In LSFM, a thin sheet of 30 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.

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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 (L_4) 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 (Pu_9) and valves (V_{10}) 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 sterotypical 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

Our instrument uses glass capillaries for specimen mounting, rather than more conventional gel embedding. The square cross-section of these capillaries should lead to less distortion than more common cylindrical capillaries. To assess the optical quality of our setup, we measured the point spread function (PSF) by imaging 28 nm diameter fluorescent microspheres dispersed in oil in these capillaries (see Methods for details). The diffractionlimited 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 μ m, and the width along the 134 detection axis (z) is 3.4 ± 0.6 μ 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 µm. (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 γ -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 $3 \,\mu m$.

147

148 Data collection capabilities

149 Using this system, we can image approximately 30 larval zebrafish per hour, obtaining from 150 each a 666 x 431 x 1060 μ 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 fluorescence images, and is not dominated by specimen positioning. In the batch of N=41fish whose neutrophil distributions were imaged in experiments described below, for example, the flow, detection, and positioning of the specimens occupied only approximately 30 seconds per fish. In the limit of zero imaging time (e.g. for very bright signals or small regions of interest), the system could therefore record data from up to about 120 specimens per hour in the absence of triggering or positioning errors, or about 90 specimens per hour 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 x 431 x 1060 µm 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 neutrophils, with a coefficient of determination $R^2 = 0.4$, indicating that a simple measure of 179

180 overall brightness, as could be assessed without three-dimensional microscopy, would181 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 $\mu m.$
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 drosophilia 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

time. In general, tackling the challenge of automated, high-throughput specimen handlingwill allow the technique of light sheet fluorescence microscopy to maximize its impact onthe life sciences.

207

208 Methods

209 Hardware

210 The majority of the instrument was constructed with off the shelf parts. Custom parts were211 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

For live imaging, each larval zebrafish flows through 0.7 mm inner diameter silicone tubing to a 50 mm long section of round 0.7 mm inner diameter, square exterior cross section, glass tubing in the specimen chamber. The contrast between the specimen and background in 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

For measurements of the point spread function, 28 nm diameter fluorescent carboxylatemodified polystyrene spheres (Thermo Fisher cat. #F8787), with peak excitation and emission wavelengths 505 and 515 nm, respectively, were dispersed in oil with a similar 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 μ m.

242

243 Ethics statement

All zebrafish experiments were carried out in accordance with protocols approved by theUniversity 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

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.

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

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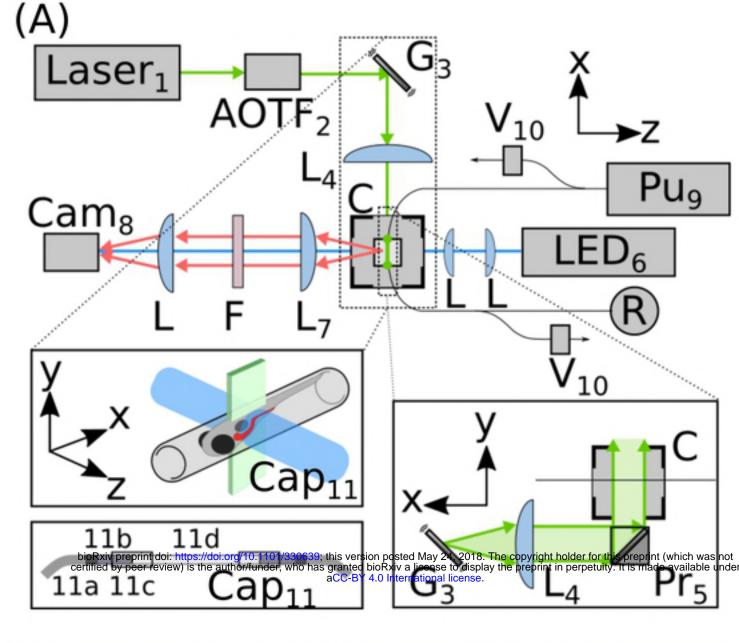
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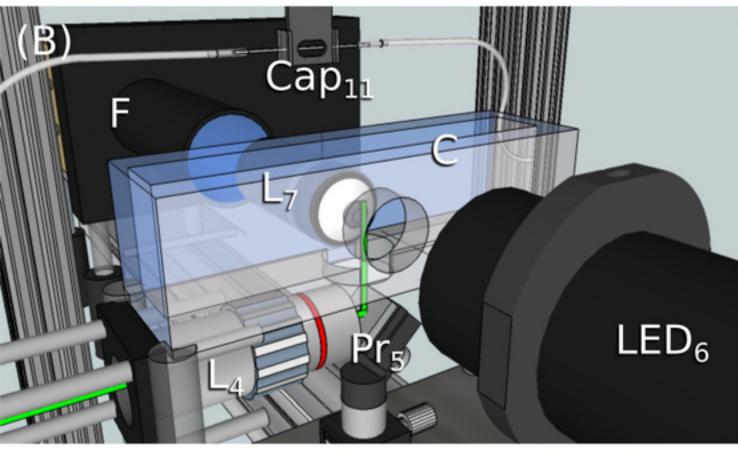
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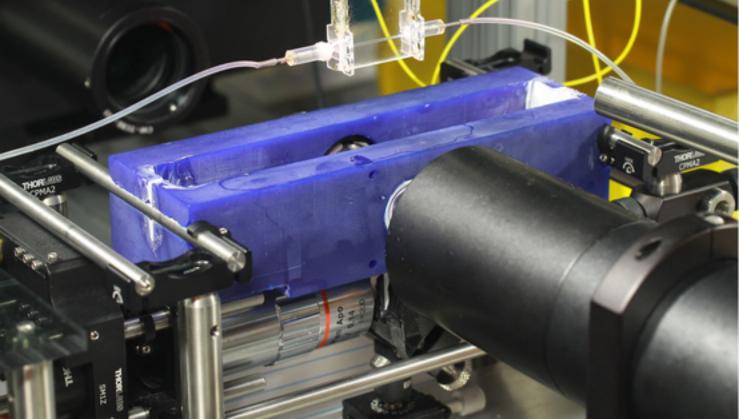
- **Supplemental Movie 1.** Video of the automated, high-throughput light sheet microscope in
- 360 operation. After a larval zebrafish, flowing through tubing, is detected and positioned, 3D
- 361 scans in two colors in two different regions, are obtained. The final image (0:57-1:03) is of
- 362 tdTomato-labeled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

363

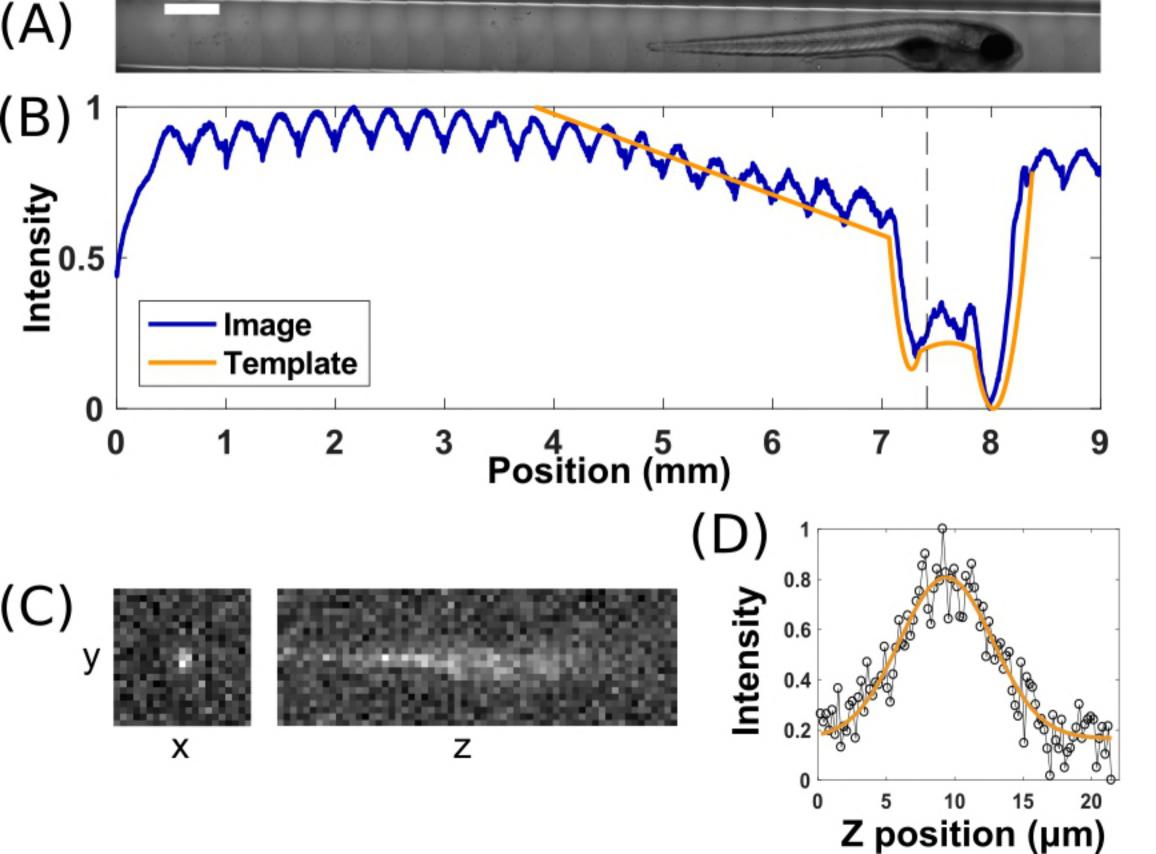
- **Supplemental Movie 2.** A light sheet fluorescence scan through the anterior intestine of a
- 365 larval zebrafish with GFP-expressing neutrophils evident as very bright objects above
- **366** autofluorescent background. The total image depth is $475 \,\mu m$ (2.5 μm /slice x 190 slices).
- **367**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

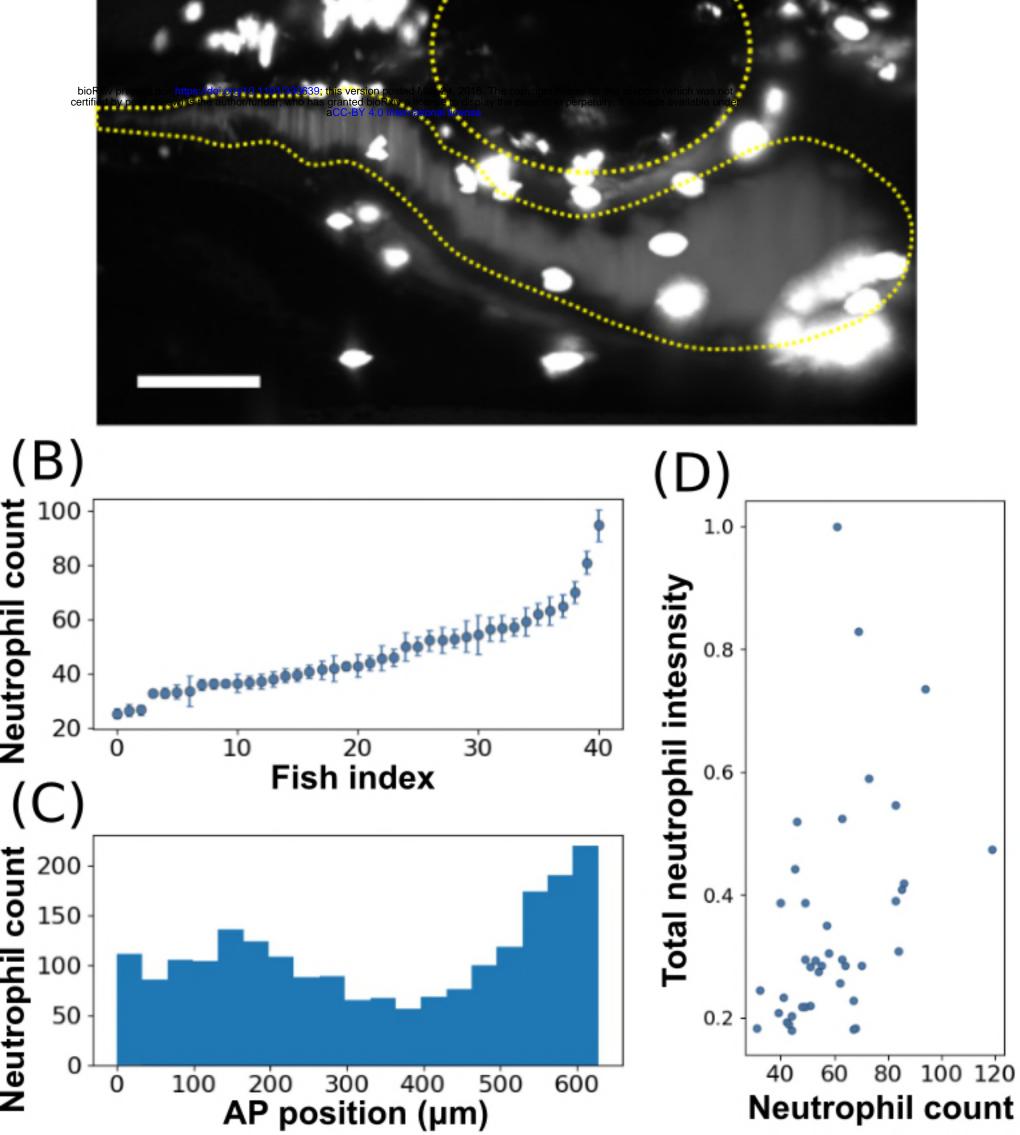






(C)





(A)