

# Label-free structural imaging of plant roots and microbes using third-harmonic generation microscopy

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## **ABSTRACT**

Root biology is pivotal in addressing global challenges including sustainable agriculture and climate change. However, roots have been relatively understudied among plant organs, partly due to the difficulties in imaging root structures in their natural environment. Here we used microfabricated ecosystems (EcoFABs) to establish imaging environments with optical access and employed nonlinear multimodal microscopy of third-harmonic generation (THG) and three-photon fluorescence (3PF) to achieve label-free, *in situ* imaging of live roots and microbes at high spatiotemporal resolution. THG enabled us to observe key plant root structures including the endodermis, Caspary strips, dividing meristematic cells, and root cap cells, as well as subcellular features including nuclear envelopes, nucleoli, starch granules, and putative stress granules. THG from the cell walls of bacteria and fungi also provides label-free contrast for visualizing these microbes in the root rhizosphere. With simultaneously recorded 3PF fluorescence signal, we demonstrated our ability to investigate root-microbe interactions by achieving single-bacterium tracking and subcellular imaging of fungal spores and hyphae in the rhizosphere.

## INTRODUCTION

Root biology research informs wide-ranging areas of intensifying global interest including soil remediation<sup>1</sup>, climate<sup>2</sup>, genetics<sup>3</sup>, and sustainable agriculture<sup>4,5</sup>. Optical microscopy, capable of visualizing plant morphology at sub-cellular resolution, is critical for understanding plant structure and function<sup>5</sup>. However,

29 compared to more extensively studied plant organs such as stems and flowers, a dearth of imaging studies on  
30 plant roots has resulted in roots being continually designated the “hidden half” of the plant body<sup>6</sup>.

31 The native environment of plant roots, composed of soil and organic matters, strongly scatters and  
32 absorbs light, severely limiting optical access to plant roots. As an alternative, microfabricated ecosystems such  
33 as EcoFABs<sup>7</sup> provide controllable and reproducible growth conditions to plants. The absence of soil in EcoFABs  
34 relieves the requirement of uprooting the plant from its growing environment and provides non-destructive  
35 optical access for *in situ* imaging of root structure and its immediate microenvironment<sup>8,9</sup>.

36 Single-photon fluorescence microscopy techniques such as confocal microscopy<sup>10</sup> and light sheet  
37 microscopy<sup>11-13</sup> have been used to image plant roots<sup>14,15</sup>. However, light scattering by root tissues has limited their  
38 applications to relatively transparent or cleared root samples<sup>16</sup>. They also often require the introduction of  
39 extrinsic fluorescent labels. Transgenic fluorescent markers are compatible with live root imaging but can only be  
40 implemented for plants with established transformation systems and even in the best cases are laborious,  
41 whereas vital stains can suffer from poor incorporation<sup>17</sup> or unwanted interference with cellular activity<sup>18</sup>. Thus,  
42 a label-free microscopy approach that can image structures at high resolution in opaque live root tissues would  
43 be highly desirable but has yet to be demonstrated.

44 Nonlinear optical microscopy methods utilizing near-infrared excitation light provide greater depth  
45 penetration in tissues than single-photon fluorescence techniques, and have been used to image both  
46 fluorescently labeled and unlabeled plant tissues<sup>19-23</sup>. Two nonlinear imaging modalities, three-photon  
47 fluorescence (3PF) microscopy and third-harmonic generation (THG) microscopy, have recently emerged as a  
48 powerful technique to image subcellular resolution at millimeter imaging depths in opaque tissues such as the  
49 mouse brain<sup>24,25</sup>. Here, we explored the potentials of using 3PF and THG microscopy for label-free *in situ* imaging  
50 at subcellular resolution of live plant roots in EcoFABs, including *Brachypodium distachyon*, a monocot grass  
51 known for its genetic tractability and relevance to agricultural crops<sup>26,27</sup>, and the dicot *Arabidopsis thaliana*, a  
52 central model organism in plant research<sup>28</sup>. We found that THG provided excellent optical resolution and label-  
53 free contrast for a variety of subcellular structures, while 3PF from the intrinsic fluorophores of root cells provided  
54 complementary structural information. We were able to visualize root hairs, vasculature, as well as subcellular  
55 components of mitotically active and border-like cells. In the optically opaque root of *B. distachyon*, THG  
56 microscopy imaged vasculature beyond 200 µm in depth in the mature zone and imaged through the entire 230-  
57 µm thickness of a root tip. Furthermore, we combined simultaneous label-free and fluorescence imaging to study  
58 root-microbe interactions, including real-time monitoring of *Pseudomonas simiae* dynamics in the vicinity of *A.*  
59 *thaliana* roots at single-bacterium resolution and visualizing the fungus *Trichoderma atroviride* adjacent to *B.*  
60 *distachyon* roots at subcellular resolution.

61 **RESULTS**

62 **THG microscopy provides label-free contrast for plant root structures at subcellular resolution**

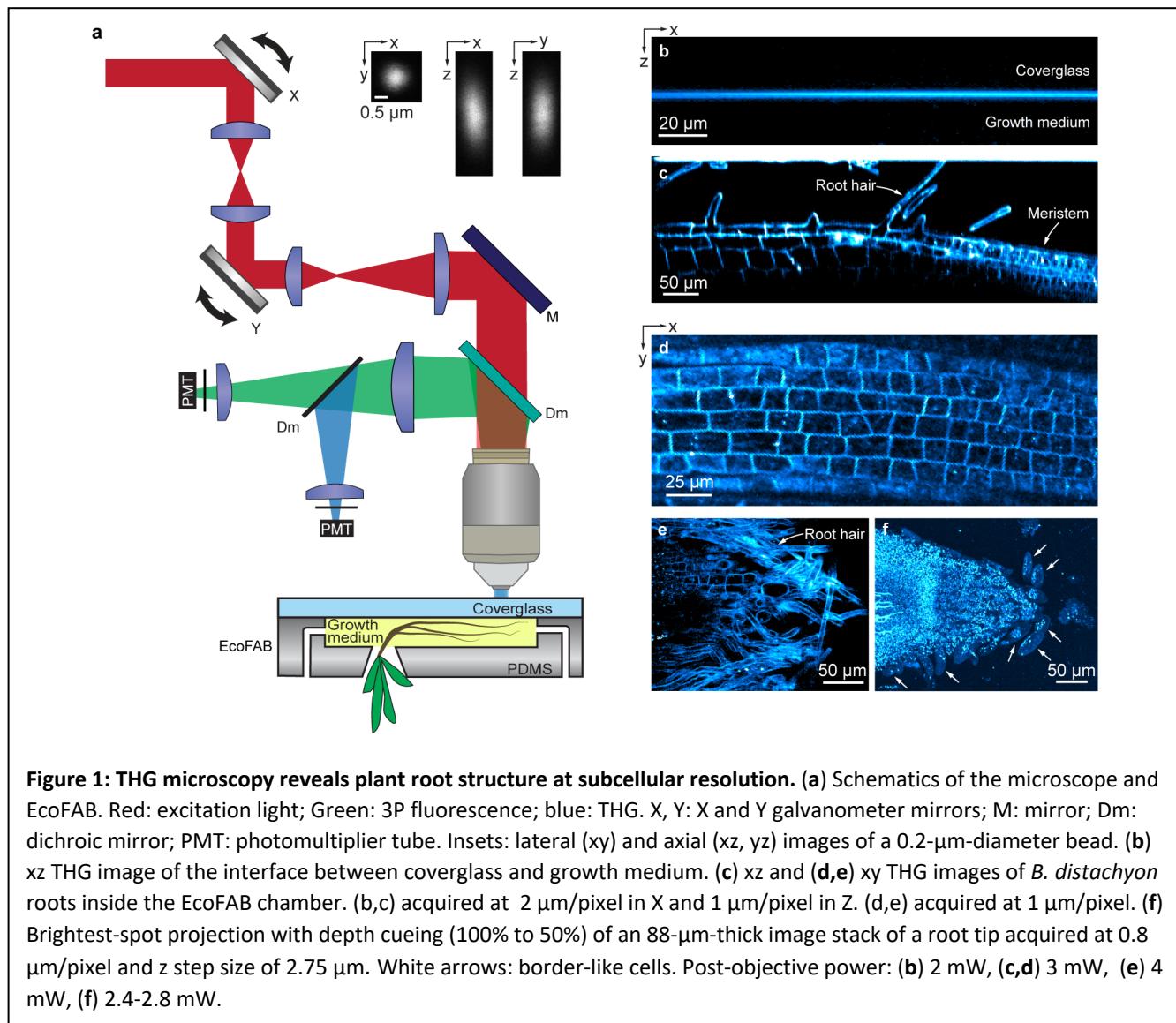
63 THG, a coherent optical process that can convert three near-infrared excitation photons to one visible photon  
64 having one-third the excitation wavelength, has its signal originating from the third-order nonlinear susceptibility<sup>29</sup>.  
65 Under tight focusing conditions in microscopy, its contrast derives from the local optical heterogeneities within  
66 the excitation focus<sup>19,30-32</sup>. When the absorption of three near-infrared photons also promotes a fluorophore to  
67 its excited electronic state, 3PF can be generated by the same excitation laser. At longer wavelengths than the  
68 third-harmonic signal, 3PF signal can be spectrally separated from and simultaneously detected with the THG  
69 signal. The third-order nonlinear excitation involved in THG and 3PF confines the signal generation to within the  
70 excitation focus, thus optically sections 3D samples.

71 We conducted all imaging experiments using a custom-built multimodal microscope<sup>25</sup> capable of  
72 simultaneously acquiring THG and 3PF signals (**Fig. 1a**). A near-infrared excitation laser beam ( $\lambda = 1300$  nm;  
73 Monaco-Opera-F system, Coherent Inc.) propagated through a laser-scanning assembly comprising galvanometric  
74 mirrors (X,Y) and scan lenses to overfill a high-numerical-aperture (NA) water-dipping objective (Olympus  
75 XLPLN25XWMP2, NA 1.05, 25 $\times$ ). THG at 433 nm and 3PF (500 – 550 nm) signals were separated from the excitation  
76 light by a dichroic mirror (Dm) and further separated into two paths by an additional dichroic mirror for detection  
77 by photomultiplier tubes. We measured the point spread function (PSF) of our microscope by 3PF imaging of 0.2-  
78  $\mu\text{m}$ -diameter fluorescent beads. The PSF had a lateral full width at half maximum (FWHM) of 0.65  $\mu\text{m}$  in the xy  
79 plane and an axial FWHM of 1.78  $\mu\text{m}$  (insets, **Fig. 1a**).

80 Seedlings of *B. distachyon* or *A. thaliana* were allowed to germinate within an EcoFAB growth chamber<sup>9</sup>  
81 made of coverglass and PDMS before loading into the microscope (Methods). The excitation light entered through  
82 the coverglass side of the chamber and the emitted THG and 3PF signals were collected and detected in the epi  
83 direction. When the excitation focus was entirely within the coverglass or the growth medium, there was no THG  
84 signal generated due to the optical uniformity of the material within the focus<sup>29,30</sup>. When the excitation focus  
85 bisected the coverglass-medium interface, a strong THG signal was generated due to the abrupt change of  
86 susceptibility from glass to growth medium (**Fig. 1b**).

87 When the excitation focus was scanned across *B. distachyon* roots, THG signal provided high-resolution  
88 label-free visualization of various structures in both axial (**Fig. 1c**) and lateral (**Figs. 1d-f**) planes. Within the root,  
89 plant cells appeared as individual compartments (**Figs. 1c-e**), with strong THG signal observed at their cell walls,  
90 likely due to the different optical properties of the cell wall and the cytoplasm<sup>33,34</sup>. We confirmed that the THG  
91 signal arose from cell walls by acquiring 3PF and THG images from root samples stained with Auramine O, a  
92 fluorescent dye known to stain plant cell walls. Compared to unstained roots, fluorescent staining substantially

93 reduced THG signal of the cell walls and severely disturbed intracellular structures (**Supplementary Fig. S1**),  
94 indicating that extrinsic fluorescent labelling led to unwanted perturbations on root structure.

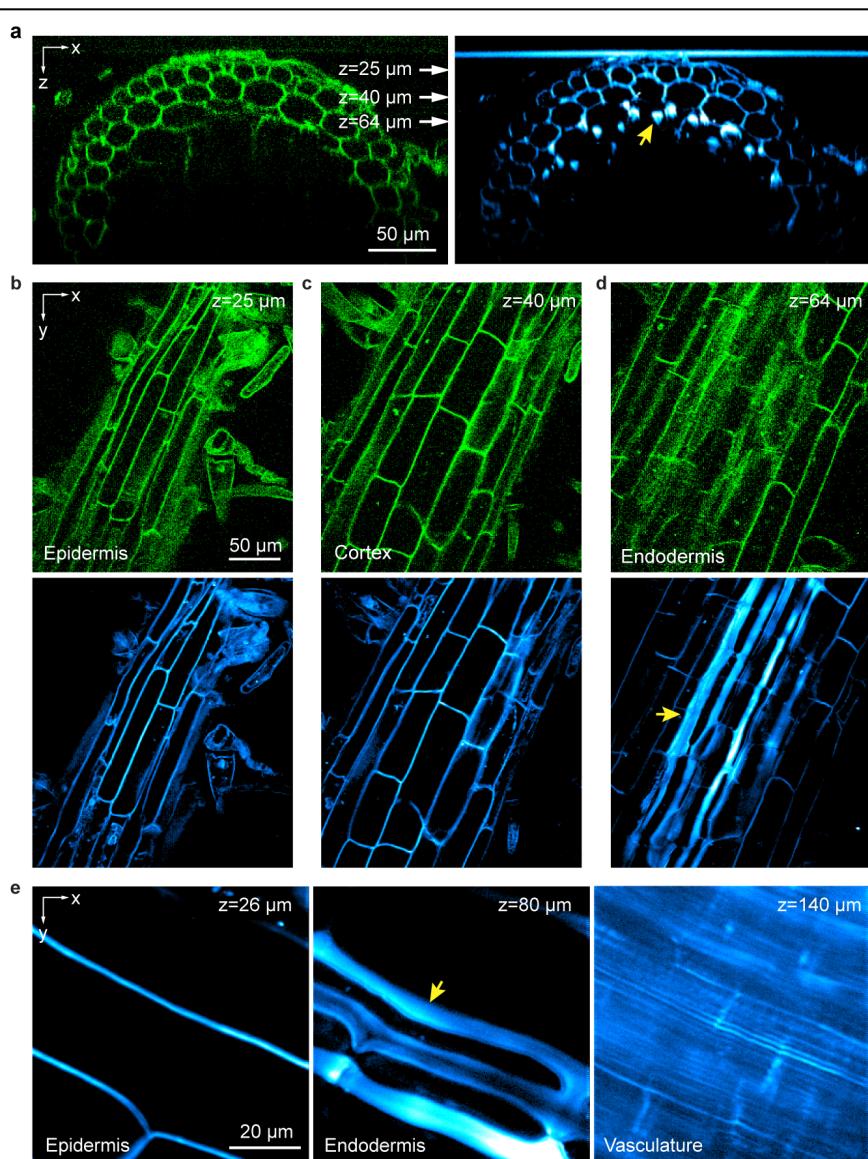


95 In an example axial (xz) image (**Fig. 1c**), we observed elongated cells with root hairs protruding from the  
96 root surface – a well-known characteristic of the mature root zone<sup>35</sup>. Here from left to right, cells progressively  
97 diminished in size, corresponding to the transition into the meristematic zone. A similar transition was observed  
98 in a lateral (xy) image (**Fig. 1d**). Another lateral image section of a slightly inclined root revealed root hairs  
99 enveloping the root's surface (**Fig. 1e**). Imaging the tip of a root, we observed cells that were detached from the  
100 primary body of the root near the root cap region (white arrows, **Fig. 1f**), reminiscent of border cells<sup>36,37</sup> and  
101 suggesting that roots grown in the EcoFAB system resemble roots grown in soil. These label-free high-resolution  
102 images motivated us to systematically explore the cellular and subcellular features revealed by THG in *B.*  
103 *distachyon* roots, as detailed below.

104 **THG and 3P autofluorescence microscopy imaging of the mature zone of *B. distachyon* roots**

105 In addition to THG, intrinsic contrast can arise from the autofluorescence of endogenous chromophores in  
106 biological specimens. Spectrally separating the THG and 3P autofluorescence signals, we simultaneously acquired  
107 THG and 3P autofluorescence images in the mature zone of *B. distachyon* roots (Fig. 2; Supplementary Video 1).

108 Roots in the mature zone contain concentric layers of epidermis, cortex, endodermis, and vasculature. In  
109 epidermis and cortex, THG and 3P autofluorescence signals co-localized at cell walls (Fig. 2a-c). Even though THG  
110 signal was ~3.4–8.8× stronger than 3P autofluorescence signal, cell walls were easily visible in both channels. The



**Figure 2: 3P autofluorescence and THG microscopy visualize epidermis, cortex, endodermis, and vasculature in the mature root zone of *B. distachyon* roots.** (a) 3P autofluorescence (green) and THG (cyan) xz images acquired at 1 μm/pixel showing a cross section of mature root zone. (b-d) xy images acquired at depths indicated by white arrows in a corresponding to (b) epidermis, (c) cortex and (d) endodermis tissues. Pixel size: 1 μm/pixel. (e) xy THG images of root tissues at 26 μm, 80 μm, and 140 μm depths, corresponding to epidermis, endodermis and vasculature, respectively. Pixel size: 0.3 μm/pixel. Post-objective powers: (a-d) 2.6 mW, (e) 1.2, 1.6, and 14 mW from left to right.

111 contrasting composition of the plant cell wall and the surrounding cytoplasm generated THG signal, while the  
112 phenolic compounds in the cell wall<sup>38-40</sup> were likely the source of autofluorescence.

113 Right before the THG and autofluorescence signals dropped off at larger depths, in the THG images we  
114 observed striation features of greater brightness and larger widths than the cell walls above (yellow arrows, **Fig.**  
115 **2a,d,e**). Interestingly, the corresponding autofluorescence image did not exhibit this large increase of brightness  
116 (e.g., 65 – 83  $\mu\text{m}$ , **Supplementary Video 1**), suggesting that THG signal was caused by a substantial change in the  
117 optical susceptibility. The location and morphology of these bright striated features of  $\sim 10 \mu\text{m}$  width were  
118 consistent with Caspary strips in endodermis, which surround the vascular cylinder and regulate the passage of  
119 water and other solutes between cortex and vasculature<sup>35</sup>. The distinct molecular composition<sup>41</sup> and thickness of  
120 Caspary strips from those of regular cell walls presumably led to its stronger THG contrast. Below the  
121 endodermis, vasculature structures were visualized as parallel channels  $\sim 1.5 \mu\text{m}$  apart (**Fig. 2e**). Although  $\sim 10 \times$   
122 higher excitation power was needed for vasculature than for cells in epidermis and Caspary strips in endodermis,  
123 we were able to observe structures more than 200  $\mu\text{m}$  deep into the mature root (**Supplementary Video 1**).

124 **THG and 3P autofluorescence microscopy image meristem and root tip of of *B. distachyon* roots at subcellular  
125 resolution**

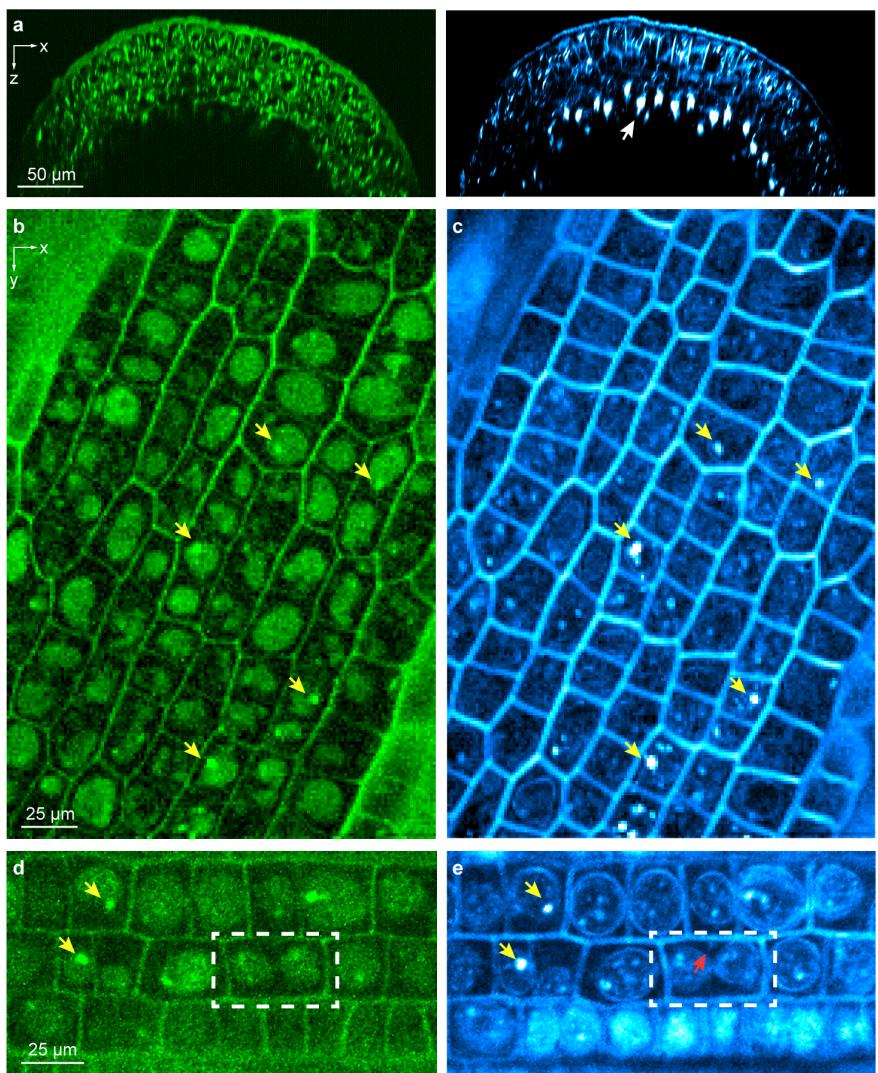
126 Compared with differentiated cells in the mature zone, cells in the meristem region contain more intracellular  
127 features required for root growth, which were visualized in the intracellular space in both 3P autofluorescence  
128 and THG cross-sectional images of a *B. distachyon* root (**Fig. 3a**). As in the mature zone, cell walls generated  
129 stronger THG signal than 3P autofluorescence signal and strong THG signal was observed in striated structures as  
130 those in the mature zone (white arrow, **Fig. 3a**).

131 In contrast to cells in the mature zone, cells in the meristem region were less elongated and approximately  
132 isodiametric (**Fig. 3b-e**). In 3P autofluorescence images, we observed ellipsoidal structures occupying most of the  
133 cell volume (**Fig. 3b,d**). These structures were likely enlarged nucleus, a characteristic feature of meristem cells,  
134 with their autofluorescence arising from the aromatic chemical structures of nucleic acid molecules themselves<sup>42</sup>.  
135 In several nuclei, we found bright and micron-sized autofluorescent aggregates (yellow arrows, **Fig. 3b,d**), which  
136 were consistent with nucleoli. The higher molecular density<sup>43</sup> could explain the brighter fluorescence observed.

137 Due to the coherent nature and symmetry requirements of THG, subcellular structures exhibited distinct  
138 features in THG images (**Fig. 3c,d**) from those of 3P autofluorescence. When the excitation focus was within the  
139 optically uniform portion of the cell (e.g., cytoplasm or the non-nucleolus, nucleoplasm part of nucleus)<sup>44</sup>, THG  
140 signal was minimal. The varying susceptibilities across cytoplasm and nucleoplasm, however, gave rise to strong  
141 THG signal at the nuclear envelope (**Fig. 3e**). Inside the nuclei, strong THG signal was observed from the putative  
142 nucleoli (yellow arrows, **Fig. 3c,e**) due to their distinct optical susceptibility from nucleoplasm<sup>44</sup>. In both

143 autofluorescence and THG images, nucleoli were often found near the nuclear envelopes, consistent with previous  
144 reports for plant cells<sup>45,46</sup>.

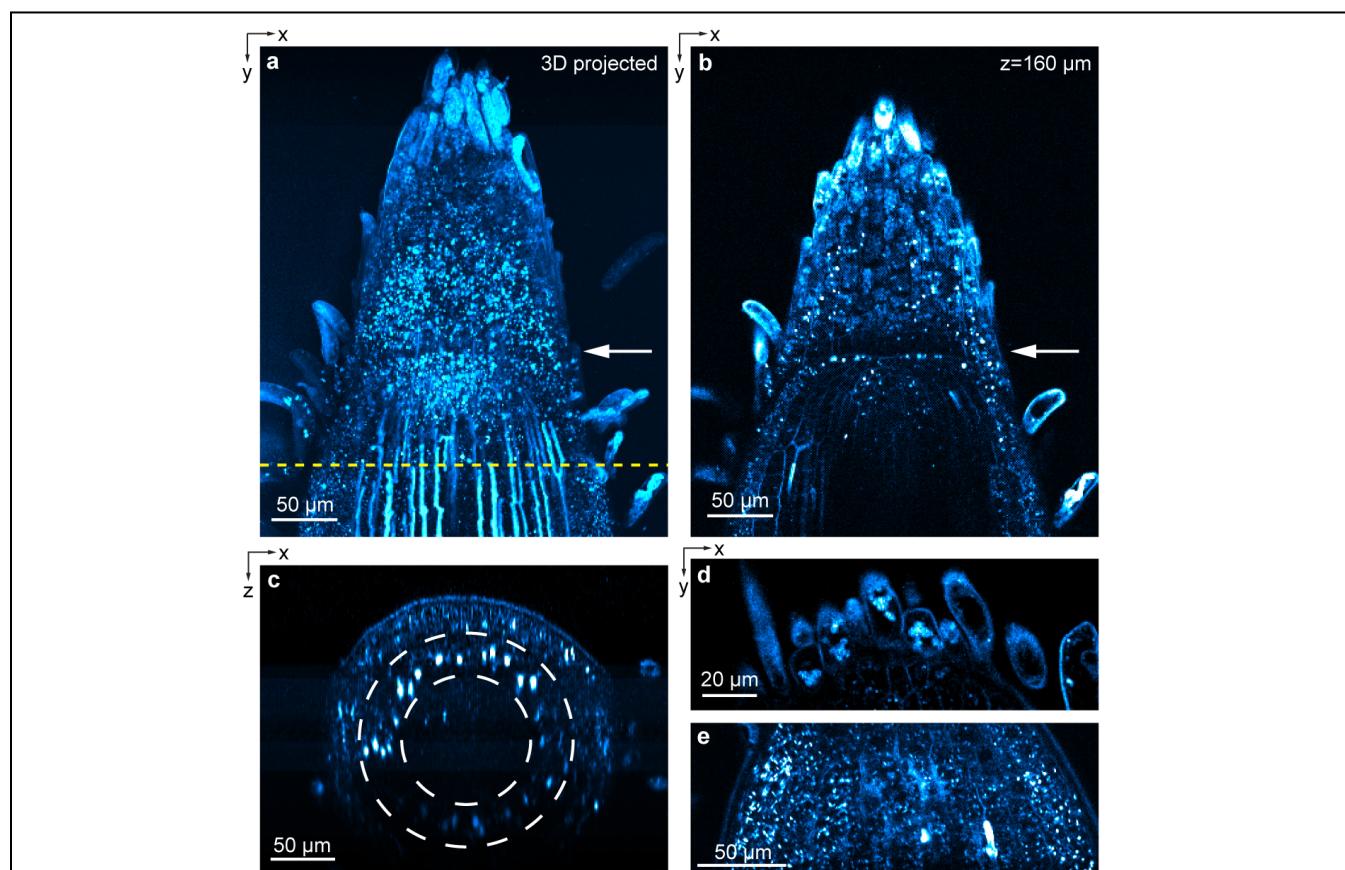
145 The distinctive features of meristem cell nuclei and the strong signal of cell walls in THG images provided  
146 us a label-free method to monitor cell division. For example, we observed a dividing cell with two daughter nuclei  
147 close to being separated (dashed box, **Fig. 3d,e**). In the equatorial plane<sup>47</sup>, a nascent cell wall could be detected in  
148 the THG image (red arrow, **Fig. 3e**).



**Figure 3: 3P autofluorescence and THG microscopy provide label-free imaging of the *B. distachyon* root meristem at subcellular resolution.** (a) 3P autofluorescence (green) and THG (cyan) xz images acquired at 0.3  $\mu\text{m}/\text{pixel}$  showing a cross section of the root meristem. White arrow: putative Caspary strip. (b-e) 3P autofluorescence (green) and THG (cyan) xy images acquired at (b,c) 0.5  $\mu\text{m}/\text{pixel}$  and (d,e) 0.3  $\mu\text{m}/\text{pixel}$ . Yellow arrows: putative nucleoli; Dashed white box: a mitotic cell; red arrow: nascent cell wall. Post-objective power: (a) 5.3 mW; (b,c) 5 mW; (d,e) 7 mW.

149 THG also provided label-free structural contrast at subcellular resolution at the root tip encompassing the  
150 apical meristem and the root cap (**Fig. 4**). Imaging through a 230- $\mu\text{m}$ -thick root tip (**Supplementary Video 2**), THG  
151 revealed a clear boundary between the meristem and the root cap<sup>35</sup> (white arrows in **Fig. 4a**, a projected image

152 of a  $373 \times 310 \times 230 \mu\text{m}^3$  volume, and **Fig. 4b**, a single image section). Bright striated structures as in the  
153 endodermis image in **Fig. 2** were found within the THG images of the meristem region but not the root cap region  
154 (**Fig. 4a; Supplementary Video 2**), consistent with known anatomy of Casparyan strips<sup>35</sup>. An axial cross-sectional  
155 view of the apical meristem region showed a complete encirclement of the central vasculature by the striated  
156 structures (**Fig. 4c**). Because light scattering and sample-induced aberration degraded focal intensity at deep  
157 depths, the Casparyan strips in the bottom half of the root were substantially dimmer than those above. In cells  
158 protruding from the tip of the root cap, we observed bright granules of 3-7  $\mu\text{m}$  in size (**Fig. 4d**), whose location  
159 and morphology were consistent with starch granules<sup>48</sup>. Throughout the apical meristem and root cap, we also  
160 saw smaller granules of 1-3  $\mu\text{m}$  in size (**Fig. 4e**). We speculated that they may be processing bodies or stress  
161 granules<sup>49</sup>.



**Figure 4: THG imaging of *B. distachyon* apical meristem and root cap.** (a) Brightest-spot projection with depth cueing (100% to 50%) of an image stack through a root tip. 230- $\mu\text{m}$ -thick image stack acquired at 1  $\mu\text{m}/\text{pixel}$  and z step size of 2  $\mu\text{m}$ . (b) xy image acquired at z=160  $\mu\text{m}$ . White arrows in a,b: boundary between meristem and root cap. (c) xz images acquired along dashed yellow line in a. (d,e) xy images from two other root samples acquired at 0.5  $\mu\text{m}/\text{pixel}$ . These roots were placed in between a microscope slide and a coverslip instead of inside EcoFAB to minimize sample motion. Post-objective power: (a-c) 3-15 mW; (d) 4 mW; (e) 3.4 mW.

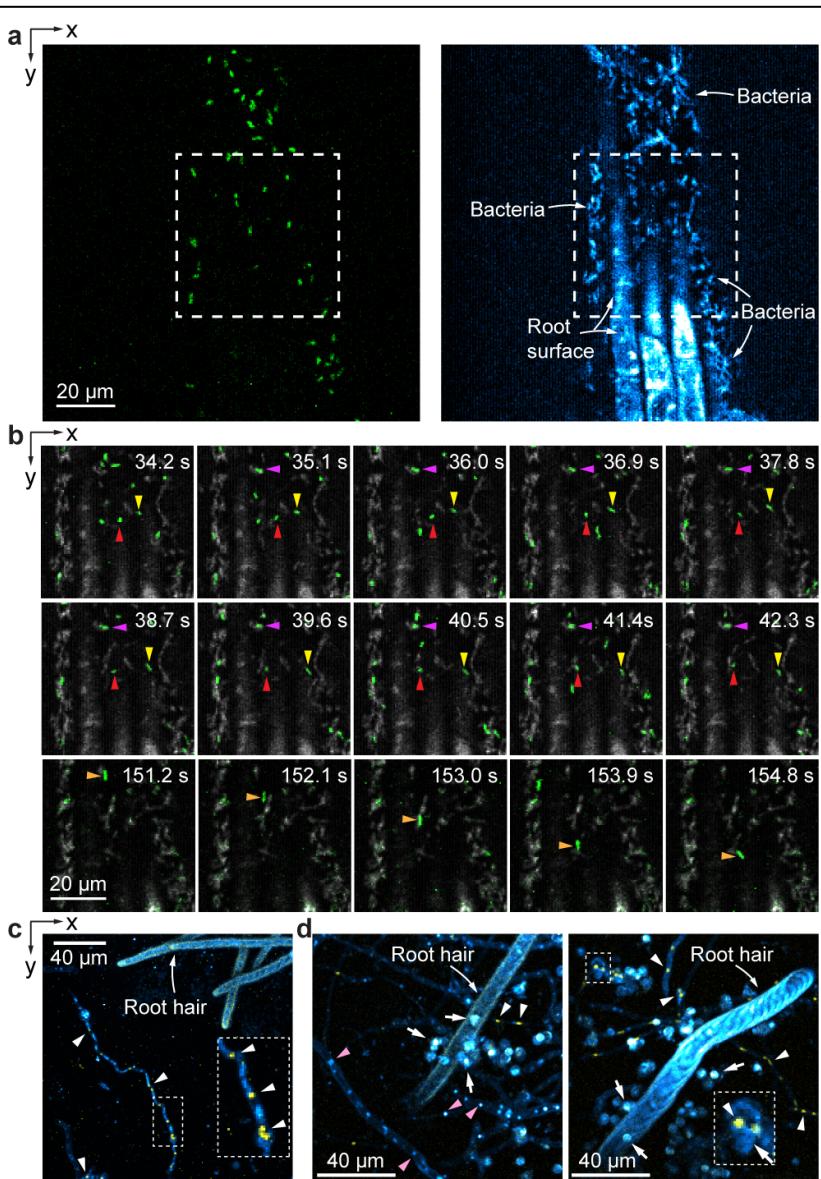
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## THG and 3PF microscopy enable simultaneous imaging of plant roots and microbes in the rhizosphere

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In addition to imaging plant roots themselves, the ability to simultaneously image microbes in the rhizosphere, the region in the vicinity of the roots where the microbiome interacts with the plant, would help understand the complex mechanisms through which root-microbe interactions impact plant growth<sup>50,51</sup>. We found



**Figure 5: Imaging root-microbe interactions at high spatial and temporal resolution.** (a) 3PF (green) and THG (cyan) xy images of *A. thaliana* root inoculated with two strains of *P. simiae* (wildtype *P. simiae* labeled with GFP, mutant *P. simiae* without GFP). (b) Consecutive frames of time-lapse imaging of the dashed box area in a with 3PF in green and THG in gray. Red/purple, yellow, and orange arrowheads: stationary, slowly-moving, fast-moving bacteria, respectively. (c,d) Maximal intensity projected 3PF (yellow) and THG (cyan) images of *B. distachyon* roots inoculated with *T. atroviride* strain IMI with GFP-labeled nuclei. (c) 50-μm-thick image stack acquired at 0.75 μm/pixel and z step size of 2.5 μm. (d) 70-μm-thick image stacks acquired at 0.5 μm/pixel and z step size of 2.5 μm. White arrowheads: GFP-labeled nuclei; pink arrowheads: unlabeled nuclei; white arrows: spores. Insets: zoomed-in views of white dashed boxes. Post-objective power: (a, b) 3.4 mW; (c) 5.6mW; (d) 5.3 mW.

169 that THG can be combined with 3P fluorescence to simultaneously image plant roots and bacteria as well as fungi  
170 in the rhizosphere *in situ*.

171 We first imaged two strains of *Pseudomonas simiae* bacteria<sup>52</sup> near the surface of an *A. thaliana* root,  
172 including a GFP-labeled wildtype strain and a mutant strain without fluorescent labeling. Both *P. simiae* strains  
173 appeared as rod-like structures in THG images, often forming aggregates around the root tissue (**Supplementary**  
174 **Videos 3,4**; right panel, **Fig. 5a**). The GFP-labeled wildtype *P. simiae* had much stronger 3P fluorescence signal than  
175 the autofluorescence from root tissue (left panel, **Fig. 5a**) and showed up in both 3PF and THG channels at  
176 comparable signal strengths. During time-lapse imaging over 160 s at 1.1 Hz frame rate (**Supplementary Video 4**,  
177 **Fig. 5b**), we observed stationary (red and purple arrowheads), slowly moving (yellow arrowheads), as well as fast  
178 moving (orange arrowheads) bacteria near the *A. thaliana* root. These results indicate that our microscope is  
179 capable of simultaneously imaging and tracking of bacteria in the root rhizosphere, and that with additional  
180 fluorescence labeling for 3PF, it can image multiple bacterial strains simultaneously.

181 We also investigated fungal colonization by imaging *B. distachyon* roots inoculated with *Trichoderma*  
182 *atroviride* strain IMI<sup>53,54</sup>, which had its nuclei labeled with GFP (H1-GFP<sup>55</sup>). Filamentous structures (**Fig. 5c**) with  
183 multiple fluorescent nuclei puncta of 1.5-2.5  $\mu\text{m}$  in size<sup>56</sup> (white arrowheads and inset, **Fig. 5c**) were identified as  
184 fungal hyphae. With their THG signal coming from fungal cell walls, these hyphae were observed near root hairs  
185 (**Fig. 5c**). In the THG image of another sample, we observed unlabeled punctate structures that are embedded  
186 within the hyphae and of similar size to the labeled nuclei, suggesting their identity as nuclei (pink arrowheads,  
187 **Fig. 5d**). In addition, spherical features with high THG signal were observed in close proximity to root hairs (white  
188 arrows, **Fig. 5d**). These spheres were 4-6  $\mu\text{m}$  in diameter and were consistent with being fungal spores, with  
189 several of them having colocalized, GFP-labeled nuclei (inset, **Fig. 5d**). Therefore, THG microscopy proved to be a  
190 valuable tool for visualizing fungal hyphal structures, spores, and nuclei, alongside root structures. When  
191 combined with fluorescent labeling, the simultaneous detection of 3PF and THG signals can provide more specific  
192 structural insights into the interactions between roots and fungi.

193 **DISCUSSION**

194 THG microscopy combined with microfabricated ecosystems allowed us to capture subcellular-resolution  
195 images of living plant roots without extrinsic fluorescent labels. Because THG signal originates from heterogeneity  
196 of optical susceptibilities within the excitation focal volume, it generates label-free visualization of cell walls. The  
197 1.3- $\mu\text{m}$  excitation light penetrated deep into the opaque tissues of *B. distachyon* roots, which are  $\sim 2.5\times$  thicker<sup>57</sup>  
198 than the more widely studied and optically transparent roots of *A. thaliana*, and enabled us to visualize the  
199 vasculature in mature roots and image through the entirety of a 230- $\mu\text{m}$ -thick root tip. Given that all cells in plant  
200 roots possess cell walls generating strong THG signal, THG microscopy can provide organ-scale views of root

201 structures at subcellular resolution. In contrast to electron microscopy, which also offer a view of subcellular  
202 features, THG microscopy can be applied to live roots without labeling. Furthermore, due to its distinct wavelength,  
203 THG signal can be combined with simultaneously acquired fluorescent signals, either from the autofluorescence  
204 of endogenous molecules or from exogenous fluorescent labels, to provide structural context for biological  
205 processes of interest.

206 The structural features we observed in THG images are consistent with the known anatomy of plant roots.  
207 These include root hairs and elongated cells in the mature root zone. In both mature and meristem roots, we  
208 observed the layered arrangement of epidermis, cortex, and endodermis. Within endodermis, we found  
209 longitudinal striation features with strong THG signal that terminated at the root cap and were consistent with  
210 the location and morphology of Caspary strips. In the root meristem, THG contrast allowed us to visualize  
211 nucleoli and nuclear envelopes, providing information on stages of cell division. These structural identifications  
212 were strengthened by simultaneously recorded 3P autofluorescence signals from cell walls, nuclei, and nucleoli.  
213 In both apical meristem and root cap, between which a clear boundary can be identified in their THG images, the  
214 subcellular resolution of our imaging system allowed us to visualize and differentiate granules of varying sizes.  
215 Whereas starch granules were observed in root cap cells and border-like cells, we speculated that the small and  
216 bright puncta throughout meristem and root cap were likely stress-related granules, whose identities need to be  
217 further confirmed with molecular labeling approaches.

218 In addition to providing global structural information throughout the plant roots at subcellular resolution,  
219 THG microscopy also allows one to image bacteria and fungi in the rhizosphere. Because both bacteria and fungi  
220 have cell walls, they could also be visualized in a label-free manner by THG microscopy. Transgenic bacteria and  
221 fungi with fluorescent protein labels further improve the specificity of structural imaging. With multimodal THG  
222 and 3P fluorescence imaging, we were able to observe dynamics of bacterial distribution and fungal spores and  
223 hyphae near roots *in situ*. With deep penetration depth and optical sectioning capability, THG and 3P fluorescence  
224 microscopy therefore enable the investigation of root-microbe interactions throughout rhizosphere and within  
225 plant roots at high spatial and temporal resolution.

226 It should be noted, however, that plant cells are susceptible to light- and/or heat-induced damages,  
227 especially during multiphoton excitation<sup>58,59</sup>. In our experiment, extended imaging of meristem zone at post-  
228 objective powers over 10 mW always induced damage in the form of increasing the amount of bright puncta.  
229 Mature zone, on the other hand, withstood prolonged imaging at tens of mW without exhibiting visible damage.  
230 Care should always be taken to ensure that the biological process of interest is not unduly perturbed by imaging.

231 Our combined EcoFAB and multimodal imaging approach provides a powerful tool for studying the cellular  
232 structure of the roots. The large imaging depth of THG and 3P fluorescence microscopy enables the study of root-

233 penetrating bacteria in opaque root tissues<sup>60</sup>. THG microscopy's ability to visualize dividing root cells will enable  
234 studies on cellular division, elongation, and differentiation during root growth. Growth conditions could be altered  
235 within the EcoFAB chamber – providing a testbed for investigating how roots respond to environmental conditions,  
236 such as salinity or nutrient levels<sup>61</sup>, at high spatiotemporal resolution. In summary, by integrating microfabricated  
237 systems with nonlinear optical microscopy for label-free imaging of plant roots, we expect that our approach will  
238 illuminate the “hidden half” of the plant, shedding light on numerous unexplored facets of root biology.

239 **METHODS**

240 **3PF and THG microscopy setup**

241 A simplified diagram of our multimodal (3PF and THG) microscopy is shown in **Fig. 1a**. The excitation source (not  
242 shown) consisted of an optical parametric amplifier (Opera-F, Coherent) pumped by a 40-W femtosecond laser  
243 (Monaco 1035-40-40, Coherent). Opera-F was tuned to generate 1,300 nm output at 1 MHz. A Pockels cell (M360-  
244 40, Conoptics) controlled the light power. A homebuilt single-prism compressor<sup>62</sup> was used to cancel out the group  
245 delay dispersion (GDD) of the excitation beam path. The excitation laser beam was reflected by two conjugated  
246 galvanometric scanning mirrors (6215H, Cambridge Technology) and relayed to the back-pupil plane of a high NA  
247 water-dipping objective (Olympus XLPLN25XWMP2, NA 1.05, 25×) by two pairs of scan lenses (SL50-3P and SL50-  
248 3P, SL50-3P and TTL200MP; Thorlabs). The objective was mounted on a piezoelectric stage (P-725.4CD PIFOC,  
249 Physik Instrumente) for axial translation of the excitation focus. The fluorescence and THG signals were collected  
250 by the same objective, reflected by a dichroic mirror (FF665-Di02-25x36, Semrock) and detected by two  
251 photomultiplier tubes (H10770PA-40, Hamamatsu). An additional dichroic mirror (Dm, FF458-Di02-25x36,  
252 Semrock) and two filters (FF03-525/50-25 for fluorescence, FF01-433/24-25 for THG; Semrock) were used to split  
253 and filter the 3PF and THG signals. Frame rates were 0.2 – 0.6 Hz except for **Fig. 4**, which was acquired at 0.03 –  
254 0.08 Hz, and **Fig. 5a,b**, which was acquired at 1.1 Hz.

255 **Bead sample**

256 Carboxylate-modified fluorescent microspheres (FluoSpheres<sup>TM</sup>, Invitrogen) were immobilized on poly(l-lysine)-  
257 coated microscope slides (12-550-12, Fisher Scientific).

258 **Imaging EcoFAB fabrication**

259 Imaging EcoFAB devices were fabricated as described previously<sup>9</sup>. Negative molds for imaging EcoFAB were 3D  
260 printed using a Form2 printer (Formlabs) with clear resin version 4 (Formlabs). EcoFAB design can be obtained  
261 from <https://eco-fab.org/device-design/>. Each EcoFAB device was housed in a magenta box with a vented lid (MK5,  
262 with vented lid, Caisson Labs) for autoclave sterilization.

263 ***Brachypodium distachyon* growth conditions for EcoFAB imaging**

264 *Brachypodium distachyon* line Bd21-3 seeds was used for this study<sup>63</sup>. Seeds were dehusked and surface sterilized  
265 in 70% ethanol for 30 s, followed by 50% v:v bleach (with 6.25% sodium hypochlorite chlorine) for 5 min, and  
266 rinsed 5 times with sterile milliQ water<sup>9</sup>. Seeds were then arranged on a sterile petri dish containing ½ Murashige  
267 and Skoog basal salt media (Caisson Labs) with 1% phytoGel (Sigma-Aldrich). Surface sterilized seeds were  
268 stratified in the dark at 4 °C for 3 days. Following stratification, seeds were allowed to germinate in a growth  
269 chamber at 25 °C at 200  $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$ , 16-hr light/8-hr dark. Three Days post germination, seedlings were  
270 transplanted into sterilized imaging EcoFABs containing 0.5x MS media with 0.8% phytoGel<sup>9</sup>. Following  
271 transplantation, plants were grown in the growth chamber for three more days before imaging. All root samples  
272 were imaged within the EcoFABs except for **Fig. 4**, for which the plant was taken out of the EcoFAB and imaged  
273 with its root between two glass coverslips to reduce root tip motion.

274 **Dissected *Brachypodium distachyon* root sample**

275 For dissected *B. distachyon* root samples imaged in **Supplementary Fig. S1**, staining was done using Bd21-3. *B. distachyon* seedlings 3 days post germination. Seedlings were dissected and fixed in 4% PFA in 1x PBS (Biotium)  
276 for at least 60 min with vacuum treatment. After fixation, seedlings were washed twice for 1 min in 1x PBS solution.  
277 For sectioned samples, primary root tissues were laterally hand sectioned using razor blades. Both hand sectioned  
278 and intact roots were then stained for lignin and suberin using 0.5% Auramine O solution (Sigma, CAS-No: 2465-  
279 27-2) adapted from Ursache et al., 2018<sup>66</sup> before imaging.

281 ***Trichoderma* culture conditions and Inoculation**

282 Three days post germination, sterile *B. distachyon* seedlings were inoculated with *Trichoderma atroviride* strain  
283 IMI<sup>54</sup> containing nuclear GFP label (H1-GFP) (generously provided by Drs. Catherine Adams and Louis Glass,  
284 University of California Berkeley, CA, USA). Fungal spores were grown on PDA plates at 28 °C for 7 days 12/12  
285 night/day cycle to induce sporulation. Spores were then harvested using sterile distilled water and separated from  
286 mycelia using a 0.4 micron filter (Pall). Spore concentration was determined using Neubauer chamber and then  
287 diluted to a spore suspension of 1x10<sup>6</sup> spores/ml. Seedlings were soaked in the spore suspension for 2 hours prior  
288 to transplanting onto Imaging EcoFABs<sup>64</sup>. Following transplantation, plants continued growing in the growth  
289 chamber for two more days before imaging.

290 ***Arabidopsis thaliana* and *P. simiae* sample preparation**

291 Seeds of *Arabidopsis thaliana* Col-0 (stock # CS66818) were obtained from the Arabidopsis Biological Resource  
292 Center (Ohio State University, Columbus, OH). Seeds were surface-sterilized by immersion in 70% (v/v) ethanol  
293 for 2 min, followed by immersion in 10% (v/v) household bleach containing 0.1% Triton X-100 (Roche Diagnostics  
294 GmbH) were stratified in distilled water at 4°C for 2 days. In this study, we utilized two strains of the root-

295 colonizing bacterium *Pseudomonas simiae*, the non-fluorescently labeled strain *P. simiae* WCS417r and the eGFP-  
296 expressing strain *P. simiae* SB642, which has been previously characterized<sup>52</sup>. Both strains were pre-cultured under  
297 kanamycin selection (150 µg·ml<sup>-1</sup>) in Luria-Bertani medium (Sigma-Aldrich) diluted in 0.5× MS medium containing  
298 2.15 g/L, 0.25 g/L of MES monohydrate (ChemCruz), and buffered to pH 5.7. The pre-cultured bacterial cells were  
299 washed twice with 0.5× MS medium and used to inoculate the stratified seeds of *A. thaliana* at an initial OD600  
300 of 0.01 for each strain. The inoculated seeds were sown into agar-filled imaging EcoFAB chamber. The growth  
301 medium contained 0.5× Murashige and Skoog basal salt mixture (Sigma-Aldrich), 2.5 mM of MES monohydrate  
302 (ChemCruz), and was buffered to pH 5.7 and solidified with 1 wt% SFR agarose (Electron Microscopy Sciences).  
303 *Arabidopsis* seedlings were grown under 16 h light (140 µmol·m<sup>-2</sup>·s<sup>-1</sup>) and 8 h dark regime at 23 °C for 10-14 days.

304 **Digital image processing**

305 Imaging data were processed with Fiji<sup>65</sup>. We used the 'Green' lookup table for 3PF images and the 'Cyan hot'  
306 lookup table for THG images. In **Fig. 5b**, the 3PF and THG signals were presented with 'Green' and 'Gray' lookup  
307 tables, respectively. For **Fig. 5c,d**, the 3PF and THG signals were presented with 'Yellow' and 'Cyan hot' lookup  
308 tables, respectively. To improve visibility, saturation and gamma of some images were adjusted. For **Fig. 5d**, we  
309 applied the 'RemoveOutliers' function in Fiji to eliminate hot pixels. We generated three-dimensional projection  
310 images (**Fig. 1f** and **Fig. 4a**) and a video (the second half of **Supplementary Video 2**) using Fiji's '3D Project' function  
311 with 'brightest-point projection' and depth cueing set at 100% to 50%.

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318 **Author contribution.** T.N., J.V., and N.J. conceived of the project; J.A.R. designed and built the microscope with  
319 help from C.R.; P.K. and M.M. prepared *B. distachyon* samples; D.P., J.A.R., P.K., and M.M. acquired *B. distachyon*  
320 images; M.M. prepared *B. distachyon* and *T. atroviride* samples; D.P. and M.M. acquired *B. distachyon* and *T.*  
321 *atroviride* images; T.T. and N.H.E. prepared *A. thaliana* and *P. simiae* samples; D.P. and T.T. acquired *A. thaliana*  
322 and *P. simiae* images; D.P., J.A.R. and N.J. wrote the manuscript with inputs from all authors.

323 **Data availability.** All data generated or analyzed during this study are included in this published article and its  
324 Supplementary Information files.

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