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1 **A mouse ear skin model to study the dynamics of innate immune**
2 **responses against *Staphylococcus aureus* biofilms**

3

4 **Running title:** Biofilm-dependent modulation of physiological inflammation

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17

18 **ABSTRACT**

19 Background: *Staphylococcus aureus* is a human pathogen that is a common cause of
20 nosocomial infections and infections on indwelling medical devices, mainly due to its ability
21 to shift between the planktonic and the biofilm/sessile lifestyle. Biofilm infections present a
22 serious problem in human medicine as they often lead to bacterial persistence and thus to
23 chronic infections. The immune responses elicited by biofilms have been described as **specific**
24 and ineffective. In the few experiments performed *in vivo*, the importance of neutrophils and
25 macrophages as a first line of defence against biofilm infections was clearly established.
26 However, the bilateral interactions between biofilms and myeloid cells remain poorly studied
27 and analysis of the dynamic processes at the cellular level in tissues inoculated with biofilm
28 bacteria is still an unexplored field. It is urgent, therefore, to develop biologically sound
29 experimental approaches *in vivo* designed to extract specific immune signatures from the
30 planktonic and biofilm forms of bacteria.

31 Results: We propose an *in vivo* transgenic mouse model, used in conjunction with intravital
32 confocal microscopy to study the dynamics of host inflammatory responses to bacteria.
33 Culture conditions were created to prepare calibrated inocula of fluorescent planktonic and
34 biofilm forms of bacteria. A confocal imaging acquisition and analysis protocol was then
35 drawn up to study the recruitment of innate immune cells in the skin of LysM-EGFP
36 transgenic mice. Using the mouse ear pinna model, we showed that inflammatory responses to
37 *S. aureus* can be quantified over time and that the dynamics of innate immune cells after
38 injection of either the planktonic or biofilm form can be characterized. First results showed
39 that the ability of phagocytic cells to infiltrate the injection site and their motility is not the
40 same in planktonic and biofilm forms of bacteria despite the cells being considerably recruited
41 in both cases.

42 Conclusion: We developed a mouse model of infection to compare the dynamics of the
43 inflammatory responses to planktonic and biofilm bacteria at the tissue and cellular levels.
44 The mouse ear pinna model is a powerful imaging system to analyse the mechanisms of
45 biofilm tolerance to immune attacks.

46 **Key words:** *Staphylococcus aureus*, Biofilm, Planktonic form, Inflammation, Mouse,
47 Intravital imaging

48

49 **BACKGROUND**

50 *Staphylococcus aureus* (*S. aureus*) is a common commensal Gram-positive bacterium that
51 colonizes the skin and mucous membranes of humans. It can also shift between planktonic
52 and biofilm lifestyles and colonize abiotic surfaces such as indwelling medical devices and
53 prosthetic implants [1]. Inside biofilms, bacteria are embedded in an extracellular matrix and
54 are more tolerant to antibiotics and to host immune attacks [2]. The resulting impact on
55 human health is enormous since biofilm infections account for more than 80 percent of
56 microbial infections in otherwise sterile tissue(s) and often become chronic [3].

57 The immune responses elicited by biofilms have been described as specific and ineffective
58 thus promoting bacterial persistence and the establishment of chronic infections [4]. Different
59 immune evasion mechanisms have been proposed to be involved, including phagocyte direct
60 killing (macrophages, neutrophils), specific recruitment of myeloid-derived suppressor cells
61 (MDSCs) and macrophage polarization towards an anti-inflammatory phenotype [4,5]. These
62 results were mostly obtained during experiments performed *in vitro* in which biofilms were
63 exposed to monocytes or neutrophils, or both [6]. In the few experiments performed *in vivo*
64 with different rodent models, several parameters vary, such as the presence of a biomedical
65 device, the tissue(s) that were inoculated or implanted with a bacteria-free or loaded device,

66 the bacteria delivery mode and the inoculum dose [7]. These studies illustrate the importance
67 of both neutrophils and monocytes/macrophages as a first line of defence against biofilm
68 infections. However, the bilateral interactions between biofilms and myeloid cells remain
69 poorly studied and analysis of the dynamic processes at the cellular level in tissues inoculated
70 with biofilms is still an unexplored field. The mouse ear pinna is currently one of the most
71 frequently used tissues to perform intravital confocal live imaging. In particular, it allows the
72 analysis of cellular behaviour in an inflamed tissue [8]. We previously developed a concept
73 that was potentially able to extract the biologically relevant features of the host and invasive
74 bacteria after injection of either the planktonic or biofilm form of bacteria in the ear pinna [7].
75 In the present study, it was decided to use the transgenic fluorescent reporter laboratory mice
76 line LysM-EGFP. Owing to the relative thinness of the ear pinna, the model enabled us to
77 perform live imaging on recruited enhanced green fluorescent protein (EGFP) fluorescent
78 leukocytes, in particular neutrophils and monocytes/macrophages. When the LysM-EGFP
79 mouse ear pinna dermis was loaded with either planktonic or biofilm bacteria, the first results
80 showed that the inflammatory response to *S. aureus* can be quantified in the skin.

81 Both bacterial forms induced a considerable inflammatory response at the injection site.
82 However, real-time analysis showed different cellular dynamics with a limited access of
83 recruited phagocytes to bacteria inside biofilms, resulting in less efficient phagocytosis. We
84 also investigated the motility of resident or recruited phagocytes and observed that cells arrest
85 at the injection site to interact with planktonic or biofilm bacteria. At early time points,
86 biofilms slowed down phagocytes and modified their trajectory. Finally, the nature of the
87 inoculum (planktonic or biofilm) influenced speed and straightness parameters differently,
88 independently of cell-bacteria interactions at the injection site.

89 We therefore developed a mouse model of infection to compare the inflammatory response to
90 planktonic and biofilm bacteria at the tissue and cellular levels. Our novel findings show that
91 the dynamics of the inflammatory responses against the two bacterial forms are different.

92

93 **RESULTS**

94 **Preparation and characterization of calibrated inocula of *Staphylococcus aureus* biofilm** 95 **and planktonic cultures**

96 A reproducible protocol of biofilm preparation was created to obtain a calibrated bacterial
97 inoculum of 10^7 colony-forming units (CFUs) in 3.8 μ L of biofilm suspension (injection
98 volume). As shown in Additional file 1: Figure S1A, titres of different aliquots of 24 h-old
99 biofilms collected in the same well or in different wells for three independent experiments
100 were comparable (Additional file 2: Table S1). To compare host immune responses to
101 planktonic and biofilm forms of *S. aureus* LYO-S2 bacteria, calibrated inocula of planktonic
102 bacteria were also prepared. The titres of the inocula were comparable for both bacterial
103 forms and contained the expected quantity of bacteria (Additional file 1: Figure S1B and
104 Additional file 2: Table S1). However, the morphological characteristics of the two inocula
105 were different, even after passing through the 34-gauge (34G) needle used for micro-injection
106 into the mouse ear tissue. Scanning electron microscopy (SEM) ultrastructural analysis
107 showed that planktonic bacteria were either dispersed or organized in small clusters (Fig. 1A
108 and Additional file 1: Figure S1C). In contrast, biofilms were organized in aggregates of
109 $29.43 \pm 7.06 \mu\text{m}$ across (Additional file 1: Figure S1D). When zoomed in, the extracellular
110 matrix is clearly observed inside these aggregates (Fig. 1B, red arrows and Additional file 1:
111 Figures S1E-H). However, the homogenization technique used to prepare biofilm inocula
112 results in an inoculum containing mainly biofilm aggregates but also detached bacteria and

113 planktonic bacteria. Future use of the term “biofilm inoculum” or “biofilm” will be in
114 reference to this type of inoculum. Using the fluorescent probe CDy11, which targets amyloid
115 fibrils, we observed that this biofilm matrix component was detected more abundantly in our
116 biofilm preparations than in the samples of planktonic bacteria (Figs. 1C-D) [9].

117

118 **Micro-injection of calibrated inocula of planktonic or biofilm forms of *Staphylococcus***
119 ***aureus* in the mouse ear pinna induces a strong inflammatory response**

120 LysM-EGFP transgenic mice were inoculated intradermally into the ear pinna with 10^7 CFUs
121 of either planktonic or biofilm mCherry-LYO-S2 fluorescent bacteria, or Trypticase Soja (TS)
122 culture medium, which was used as a control. Inflammatory responses were followed at early
123 (4-7 hours post-injection [hpi]) and late time points (after 24 hpi) by measuring the intensity
124 of the EGFP signal for each group (Figs. 2A-C). The image of the ear pinna enabled us to
125 analyse overall inflammation in the entire tissue (mosaic acquisition). To quantify this signal,
126 we created the following protocol. A region of interest (ROI) was drawn on late time point
127 images, where the EGFP signal was more easily detectable, and applied to early time point
128 images. The ratio of the sum of EGFP fluorescence intensities to ROI areas was calculated
129 with this protocol and the inflammatory response was compared at early and late time points
130 in the two groups of infected mice (Fig. 2D and Additional file 3: Table S2). We used the
131 same protocol for the control group and observed a non-specific recruitment of EGFP+ innate
132 immune cells due to the physical trauma from injection and the introduction of TS culture
133 medium (Additional File 4: Movie 1). At early time points, both bacterial forms induced an
134 inflammatory response, with a statistically significant increase only in the group of mice
135 inoculated with planktonic bacteria. Thus, planktonic bacteria induced a greater response than
136 biofilms after 4 hpi. Between the early and late time points, the inflammatory response was

137 significantly greater for both bacterial forms. At late time points, the response was
138 considerable in both groups of challenged mice compared to control mice, with no significant
139 difference between mice inoculated with planktonic or biofilm bacteria (Fig. 2D).

140

141 **Dynamics of immune cell recruitment after the micro-injection of either planktonic or**
142 **biofilm forms of *Staphylococcus aureus* in the mouse ear pinna are different**

143 LysM-EGFP transgenic mice were inoculated intradermally into the ear pinna with 10^7 CFUs
144 of either planktonic or biofilm mCherry-LYO-S2 fluorescent bacteria, or with TS culture
145 medium. We created a confocal acquisition protocol to analyse the dynamics of recruited
146 EGFP+ cells at the inoculation sites by real-time imaging. A red autofluorescence signal is
147 emitted by mice hairs and could not be prevented by shaving the ear pinna. Indeed, this
148 operation would have induced a non-specific inflammatory response. In control mice,
149 recruitment was low owing to injection trauma (Additional File 4: Movie 1). In mice
150 inoculated with bacteria, an influx of phagocytic cells was observed at early (3 to 6 hpi) (Figs.
151 3A-B, white circles) and late time points (after 24 hpi) (Figs. 3C-D) for both bacterial forms.
152 At early time points, immune cells were present over the entire surface of planktonic bacteria
153 injection sites and multiple contact points between cells and bacteria were observed. In
154 addition, numerous immune cells infiltrated the injection sites (Fig. 3A and Additional file 5:
155 Figures S2A-B, white arrowheads; see also Additional File 6: Movie 2). In biofilms, the
156 contact points were less numerous and were mainly located at the periphery of the injection
157 site. In contrast to planktonic inocula, a small number of cells succeeded in infiltrating the
158 biofilm (Fig. 3B and Additional file 5: Figures S2C-D, white arrowheads; see also Additional
159 File 7: Movie 3). The fluorescent signal was less detectable for planktonic bacteria after 24 h,
160 suggesting that bacterial lysis after phagocytosis had occurred (Fig. 3C, white empty

161 arrowhead). For biofilms, phagocytosis seemed to be less effective, since a fluorescent signal
162 was still clearly visible after 24 h (Fig. 3D). Overall, this real-time analysis using an intravital
163 imaging approach shows that the dynamics of the inflammatory responses against planktonic
164 and biofilm bacteria are different.

165

166 **Motility of recruited innate immune cells is different after injection of planktonic or** 167 **biofilm forms of *Staphylococcus aureus* in the mouse ear pinna**

168 Using Imaris software, we created an analysis protocol to track the motility properties
169 (average speed and straightness) of EGFP⁺ cells recruited at the injection zone from the
170 previously acquired time-lapse videos (Fig. 4). Using the “Spots” tool we attributed a sphere
171 to a number of immune cells observed in the acquisition field (Figs. 4A-B, white spheres).
172 This enabled us to establish a trajectory (Figs. 4A-4B, multicoloured lines) for each sphere
173 corresponding to the path taken by each cell over time in the ear tissue. We then compared the
174 average speed and straightness of the trajectories of phagocytic cells in response to the two
175 bacterial forms. In different zones of the injection site, cells interacted with bacteria (Fig. 4A)
176 or not (Fig. 4B). We first analysed the motility of all cells in response to bacteria (planktonic
177 or biofilm) or to TS culture medium without distinction between cells that interacted with
178 bacteria and cells that did not. At early time points, only biofilms induced a significant
179 decrease in cell speed, compared to control mice and mice inoculated with planktonic
180 bacteria. Thus, biofilms slowed down recruited cells, an effect that was maintained 24 hpi. In
181 contrast, planktonic bacteria significantly increased cell speed compared to the control group
182 (Fig. 4C and Additional file 8: Table S3). This differential response induced by the two
183 bacterial forms was also seen for cell trajectory straightness, which was significantly
184 decreased at early time points only by biofilms (less straight trajectory of EGFP⁺ recruited

185 cells) compared to the control group. At late time points, we observed an opposite effect, as
186 both bacterial forms significantly increased straightness compared to the control group, with a
187 more pronounced effect for planktonic bacteria (Fig. 4D and Additional file 9: Table S4). We
188 further analysed the motility of cells interacting with bacteria (bacteria contact) or not (no
189 bacteria contact) in different zones of the cutaneous injection site for the same time point
190 (Figs. 4E-H). Cell motility was compared after inoculation of biofilm or planktonic bacteria.
191 At early time points, the presence of the two forms of bacteria (bacteria contact) induced a
192 significant decrease in both speed and straightness (Figs. 4E-F, Additional file 10: Table S5
193 and Additional file 11: Table S6). This indicates that cells arrest at the injection site to interact
194 with inoculated bacteria. At early and late time points, cell speed was reduced in biofilms
195 compared to planktonic cells, independently of the presence of bacteria (Figs. 4E and 4G,
196 Additional file 12: Table S7). Finally, at late time points, straightness was reduced for cells
197 interacting with biofilms, compared to planktonic inocula (Fig. 4H and Additional file 13:
198 Table S8). Taken together, these results demonstrate that the cell dynamics of the
199 inflammatory response are different after inoculation of biofilm or planktonic bacteria. The
200 mouse ear pinna model evidences an inflammatory response specific to biofilms that is
201 probably one mechanism of its tolerance to immune attacks.

202

203 **DISCUSSION**

204 The dynamics of the implementation of immune responses during *S. aureus* infections *in vivo*
205 is a key event. It is a determinant factor especially during planktonic-to-biofilm transition, as
206 the bacterial persistence associated with the “chronicization” process of biofilm infection
207 often depends on it. However, it is difficult to follow these events in mammals over time and

208 this hinders the clear understanding of the immune evasion mechanisms of *S. aureus* biofilms
209 and therefore the design of preventive strategies against biofilm infections.

210 In the present study, we compared for the first time the dynamics of early innate immune
211 responses to planktonic and biofilm *S. aureus* in the skin. The skin is a common target tissue
212 for *S. aureus* infections and the mouse ear pinna is frequently used as a cutaneous imaging
213 site. This accessible tissue can be rapidly and easily prepared for imaging over long periods of
214 time. Calibrated inocula were injected intradermally in a very small volume with a 34G
215 needle to limit inflammation resulting from injection trauma. The maturation state of the
216 biofilm culture and the bacterial inocula were two major criteria in the finalization of the
217 protocol. We prepared “young” biofilms (24 h-old) as previous studies reported that immature
218 biofilms are more susceptible to neutrophil attack than mature biofilms [10]. We inoculated a
219 high number of bacteria (10^7 CFUs) into the ear tissue of LysM-EGFP transgenic mice, as in
220 previously published mouse models of *S. aureus* skin infections such as the chronic diabetic
221 wound model and the air pouch model [11,12]. Once drawn up, our protocol enabled us to
222 compare qualitatively and quantitatively the innate immune responses induced by a
223 comparable dose of planktonic or biofilm bacteria. As described above, our biofilm inoculum
224 contained bacterial aggregates and also planktonic and detached bacteria. The phenotype of
225 the latter is similar to that of biofilm bacteria. Indeed, previous studies have described
226 differential gene expression profiles for *S. aureus* in its planktonic or biofilm form [13,14].
227 One major difference between the two inocula was the presence of the extracellular matrix in
228 the biofilm aggregates. Among the components of the LYO-S2 *S. aureus* biofilm matrix
229 inoculated into mice, we detected amyloid fibrils. Small peptides called phenol-soluble
230 modulins (PSMs) produce amyloids in *S. aureus* biofilms and notably contribute to *S. aureus*
231 biofilm stability. They are also described as key virulence factors capable of stimulating

232 inflammatory responses or affecting leukocyte viability or functions [15,16], and could
233 contribute to the specific innate immune responses observed with biofilm bacteria.

234 After inoculation, we used the mouse ear pinna model to follow the inflammatory response to
235 *S. aureus* over time at the tissue level. Imaging analysis showed that bacterial inocula induced
236 an early inflammatory response at the cutaneous injection site in LysM-EGFP transgenic
237 mice, consisting of recruited EGFP⁺ phagocytes, in particular neutrophils and
238 monocytes/macrophages. We quantified this response for the first time and showed that it is
239 significantly increased with the planktonic form after 4 h, compared to that in control mice.
240 The absence of significant differences between the control and biofilm conditions could have
241 been due to early phagocyte killing, as previously reported [4,5,17]. After 24 h, the
242 inflammatory response was considerable and comparable for the two bacterial forms. In
243 rodent models, neutrophils are usually the most rapidly recruited, and therefore most
244 abundant, cells in the proximity of biofilms [7]. In a catheter-related model, however,
245 monocytes were the first cells to be recruited [18]. Although overall responses are comparable
246 at late time points, we postulate that the phenotype of recruited cells in our model differs
247 according to whether planktonic or biofilm bacteria are injected with, for example, the
248 specific recruitment of myeloid-derived suppressor cells (MDSCs) for the biofilm, as
249 described previously [19].

250 The mouse ear pinna model then enabled us to follow the inflammatory responses to
251 planktonic or biofilm *S. aureus* over time at the cellular level. Analysis of the dynamics of
252 recruited EGFP⁺ cells at the inoculation sites by real-time imaging showed that the dynamics
253 differed between planktonic and biofilm bacteria. Our results show that biofilm acts as a
254 physical barrier [20]. Few cells infiltrate the biofilm, with most recruited cells being present at
255 the periphery of the inoculum. After 24 h, when innate immune responses have been
256 considered as set up, phagocytosis seemed to be limited. Indeed, the bacterial signal was still

257 intense, compared to the low signal observed with planktonic bacteria at the same time point.
258 This impairment of phagocytosis observed with biofilms is commonly known as “frustrated
259 phagocytosis” [20]. Complementary experiments are required to quantify the bacterial load in
260 the ear tissue over time.

261 The mouse ear pinna model further enabled us to obtain reproducible quantitative
262 measurements of the speed and straightness of recruited innate immune cells. To obtain the
263 most accurate representation of these motility parameters, stringent algorithm settings were
264 used. Then, manual corrections were applied to cell tracks. For example, only tracks lasting
265 three or more frames were considered during the time that cells were visible in the
266 observation field. Tracks that converged into one were eliminated to avoid any uncertainty
267 about the resulting cell trajectory. Likewise, cells near or exiting the border of the image
268 volume were carefully checked to ensure that the same cell was not counted twice with two
269 different tracks. Analysis of innate immune cell migration showed that cells behaved
270 differently in presence of planktonic and biofilm bacteria. Study of the entire population of
271 cells in the tissue (cells interacting or not with bacteria) showed that biofilms generally
272 decreased cell speed and straightness. In addition, when immune cells interacted with bacteria
273 at the injection site, biofilms generally decreased cell speed more significantly than did
274 planktonic bacteria. A possible correlation could be made with previous observations
275 describing immobilized neutrophils on *Pseudomonas aeruginosa* biofilms *in vitro* after loss of
276 their pseudopodia [21]. Interestingly, biofilms also induced a remote effect on cell speed, as
277 cells with no visible contact with bacteria moved more slowly when the inoculum was in the
278 biofilm form. This result suggests the potential diffusion of small molecules from the biofilm
279 capable of influencing the behaviour of proximal recruited cells [5]. We thus provide
280 evidence that cell motility is affected differently by planktonic and biofilm bacteria. Notably,
281 the latter has a greater effect on speed and straightness. Further work is needed on the fine

282 interactions between cells and bacteria in order to study phagocytic cell arrest and subsequent
283 phagocytosis (or lack of).

284

285 **CONCLUSIONS**

286 The mouse ear skin model proposed here detects and measures the inflammatory responses
287 induced by biofilm and planktonic bacterial challenge over time. It has great potential to
288 elucidate the specific mechanisms used by biofilms to circumvent host innate immune
289 responses and therefore to develop new preventive strategies specifically targeting host
290 immune responses during biofilm infections.

291

292 **MATERIAL AND METHODS**

293 **Mice and ethical statement**

294 LysM-EGFP transgenic mice (6- to 8-week-old males and females) were obtained from the
295 bacteria-cell interactions unit, Pasteur Institute (Paris, France), and bred in the animal care
296 facility at Université Clermont Auvergne (Clermont-Ferrand, France). All experiments were
297 approved by the Ethics Committee on Animal Experimentation of Auvergne C2E2A,
298 Clermont-Ferrand, France (agreement number: 1725) and were carried out in accordance with
299 the applicable guidelines and regulations.

300

301 **mCherry-tagged strain construction**

302 The *S. aureus* LYO-S2 mCherry-tagged strain was constructed after insertion of pAH9
303 plasmid [22] into the LYO-S2 clinical strain [23] by electroporation, as described previously

304 [24]. The *S. aureus* LYO-S2 mCherry-tagged fluorescent strain, named *S. aureus* mCherry-
305 LYO-S2, was selected onto Luria-Bertani (LB) agar containing erythromycin (10 µg/mL).
306 The plasmid was maintained by growing the strain in TS culture medium containing
307 erythromycin (10 µg/mL). Fluorescence was detected in bacterial suspensions by fluorescence
308 microscopy.

309

310 **Bacterial growth conditions**

311 *S. aureus* LYO-S2 or the mCherry-LYO-S2 fluorescent strain were grown in TS culture
312 medium at 37°C with shaking and stored at -80°C in the same medium containing 15%
313 glycerol. Planktonic bacteria were cultured at 37°C in TS culture medium under aerobic
314 conditions and harvested after overnight growth (stationary phase). For biofilm preparations,
315 overnight cultures were adjusted to $2 \cdot 10^7$ CFUs/mL of TS culture medium and added to 24-
316 well cell culture plates (1 mL per well). Twenty-four-hour-old biofilms were obtained after
317 incubation of plates at 37°C without shaking.

318

319 **Preparation of bacterial inocula**

320 Before injection, *S. aureus* mCherry-LYO-S2 planktonic inocula were prepared from the
321 overnight growth, which was first homogenized. Bacterial concentration was then deduced by
322 measuring the OD₆₀₀ and using the known bacterial titre of the strain at $6.5 \cdot 10^8$ CFU/OD unit.
323 A specific volume of the overnight growth containing 10^7 CFUs was then withdrawn and
324 centrifuged at 3000 x g for 5 min. The supernatant was eliminated and bacteria were
325 resuspended in TS culture medium to obtain a final concentration of 10^7 CFUs per 3.8 µL of
326 culture medium. For *S. aureus* mCherry-LYO-S2 biofilms, inocula were obtained by carefully

327 eliminating 700 μL of the supernatant from each well in the cell culture plate. The remaining
328 biofilm volume was then delicately homogenized and 3.8 μL , corresponding to 10^7 CFUs, was
329 collected for further inoculation to mice. Serial dilutions of both planktonic and biofilm
330 inocula were plated on LB agar plates for titration. Biofilm inocula were sonicated three times
331 for 5 min each before dilution (Fisher Scientific, 80W, 37kHz). CFUs were counted after 24 h
332 at 37°C.

333

334 **Inoculation of bacteria into mice**

335 Mice were anesthetized by intraperitoneal injection of a mixture of ketamine (50 mg/kg) and
336 xylazine (5 mg/kg). A small volume (3.8 μL) of planktonic or biofilm inocula or TS culture
337 medium were injected into the dorsal ear dermis of anesthetized mice with a 34G needle fitted
338 to a NanoFil syringe (World Precision Instruments) [25]. A characteristic papule was
339 observable at the injection site, evidence of an intradermal injection.

340

341 **Scanning electron microscopy observation of bacterial preparations**

342 For electron microscopy observations, biofilms and planktonic inocula were prepared as
343 described above and deposited on SEM Pore (Jeol filters) with a 34G needle fitted to a
344 NanoFil syringe. After absorption, bacteria were fixed overnight at 4°C with glutaraldehyde
345 1.6% in 0.2 M cacodylate buffer at pH 7.4, supplemented with ruthenium red at 0.15%. They
346 were then rinsed in the same buffer. After post-fixation for 1 h with 1% osmium tetroxide in
347 cacodylate buffer at room temperature, samples were washed for 20 minutes in distilled water
348 and dehydrated by graded ethanol from 25° to 100° (10 minutes each) to finish in
349 hexamethyldisilazane (HMDS) evaporated overnight. After drying, samples were sputter-

350 coated with gold-palladium (JFC-1300, JEOL, Japan). Morphology analyses were made with
351 a scanning electron microscope JSM-6060LV (Jeol, Japan) at 5 kV in high-vacuum mode.

352

353 **Detection of amyloid fibrils in biofilm preparations**

354 Planktonic suspensions and biofilms of *S. aureus* LYO-S2 were prepared as described
355 previously. For planktonic bacteria, $5.6 \cdot 10^8$ CFU were withdrawn from the overnight culture.
356 The suspension was then centrifuged as before and bacteria were resuspended in 200 μ L of
357 TS culture medium. For 24 h-old biofilm cultures, 700 μ L of supernatant were carefully
358 withdrawn from the cell culture plates before homogenization of the remaining suspension. A
359 10 μ M stock solution of the fluorescent probe CDy11 [9] was prepared in dimethyl sulfoxide
360 (DMSO). The solution was diluted in phosphate-buffered saline solution (PBS) to prepare a
361 100 μ M solution. Ten μ L of the diluted probe was then added to each bacterial preparation
362 and incubated for 45 min in the dark at room temperature. TS culture medium (800 μ L) and 2
363 μ L of the live cell fluorescent label SYTO9 from the LIVE/DEAD BackLight Bacterial
364 Viability Kit (Molecular probes) were then added to each preparation and left to incubate for
365 15 min in the dark at room temperature. Ten μ L of planktonic and biofilm preparation
366 samples were deposited on glass slides for further observation by fluorescence microscopy.
367 Image acquisition was carried out on a ZEISS Cell Observer Spinning Disk Confocal
368 Microscope (Carl Zeiss Microscopy, Germany), with two different lasers to observe
369 fluorescence emitted from SYTO9 and CDy11 (excitation at 488 and 590 nm, emission at 509
370 and 612, respectively, with exposure times set at 100 ms for both channels). Acquisition was
371 performed with 20X (dry) objectives. Each image corresponds to the Z-projected average
372 intensity signal for each channel.

373

374 ***In vivo* confocal imaging: acquisition**

375 **Time-lapse video acquisition.** Three to 6 hpi, mice were anesthetized by intraperitoneal
376 injection of a mixture of ketamine (50 mg/kg) and xylazine (5 mg/kg). Infected ears were
377 prepared as described previously [26] and imaged on a ZEISS Cell Observer Spinning Disk
378 Confocal Microscope (Carl Zeiss Microscopy, Germany). Video acquisition was carried out
379 with two different lasers to observe EGFP and mCherry fluorescence (excitation at 488 and
380 590 nm, emission at 509 and 612, with exposure times set at 100 and 300 ms, respectively).
381 Acquisition was performed with 10X (dry) and 20X (dry) objectives for periods of 20 to 30
382 min. With the 10X objective, multiple fields of observation were required as the entire
383 injection site was imaged. Z-stacks and intervals between images were adjusted according to
384 the thickness of the ear tissue. Acquisition was repeated 24 hpi. Ear tissues of control mice
385 were inoculated with TS culture medium and imaged at the same time points.

386 **Mosaic acquisition.** Infected ears were also imaged on a ZEISS LSM 800 (Carl Zeiss
387 Microscopy, Germany) confocal microscope with a 10X objective (dry). Multiple fields of
388 observation covering the entirety of the tissue surface were imaged to get a reconstructed
389 image of the ear. To set up acquisition parameters, multiple focal points
390 distributed homogenously over the acquisition zone were chosen. EGFP fluorescence signal
391 was detected in six Z-stacks spanning 75 μm of tissue, with an exposure time of 9.5 ms. The
392 bright-field signal was also detected on a central stack, with an exposure time of 10 ms.
393 Acquisition was repeated after 24 h, with imaging sessions typically lasting 30 to 45 min. Ear
394 tissues of control mice injected with TS culture medium were also imaged with the same
395 protocol.

396

397 ***In vivo* confocal imaging: analysis**

398 **Time-lapse video analysis.** Videos acquired with the 10X objective were first stitched
399 together using ZEN software. Each image extracted from time-lapse videos corresponds to the
400 Z-projected average intensity signal for each channel at the corresponding time point. Time-
401 lapse videos at 20X and 10X were then analysed with Imaris software using the “Spots” tool.
402 For each cell, a track was generated by the software and manually corrected according to
403 specific criteria: number of frames superior to three and elimination of converging tracks
404 between two different cells. Two different parameters (average speed and straightness) of
405 immune cell dynamics were then extracted. For each time point, both parameters were
406 analysed in different zones of the cutaneous injection site, where cells were in contact or not
407 with the bacterial inoculum.

408

409 **Mosaic analysis.** Images acquired on the ZEISS LSM 800 confocal were stitched together
410 using ZEN software to reconstitute an entire image of the ear tissue at early and late time
411 points. A maximum intensity projection image was created from image Z-stacks. A ROI was
412 then drawn manually around the EGFP fluorescent zone of the 24 h image to obtain the sum
413 of EGFP fluorescence intensities of each pixel in the ROI. The shape of the ROI was
414 conserved and applied to the early time point image. The ratio of the sum of intensities of
415 EGFP fluorescence to the area of the ROI was then calculated for both time points. The
416 images shown represent the Z-projected maximal intensity signal of a reconstituted image of
417 the ear tissue for the EGFP channel.

418

419 **Statistical analysis**

420 Prism 5 software (GraphPad Software, Inc.) was used to analyse the statistical significance of
421 data sets by the Mann-Whitney two-tailed test. $p \leq 0.05$ was considered statistically significant
422 (symbols: **** $p \leq 0.0001$; *** $p \leq 0.001$; ** $p \leq 0.01$; * $p \leq 0.05$; ns = non-significant).

423

424

425 **Abbreviations**

426 CFU: Colony Forming Unit; DMSO: Dimethyl sulfoxide; EGFP: Enhanced Green
427 Fluorescent Protein; EGFP+: Enhanced Green Fluorescent Protein-positive; hpi: hours post-
428 injection; LB: Luria-Bertani; PBS: Phosphate Buffered Saline; PSM: Phenol Soluble
429 Modulins; ROI: Region Of Interest; SEM: Scanning Electron Microscopy; TS: Trypticase
430 Soja.

431

432 **Ethics approval and consent to participate**

433 LysM-EGFP transgenic mice (6-8 weeks-old male and female) were obtained from the
434 bacteria cell interactions Unit, Pasteur Institute, Paris, France, and bred in the SPF animal care
435 facility at University Clermont Auvergne (Clermont-Ferrand, France). All experiments were
436 approved by the local Ethics Committee on Animal Experimentation of Auvergne C2E2A,
437 Clermont-Ferrand, France (agreement number: 1725) and were carried out in agreement with
438 the applicable guidelines and regulations.

439

440 **Consent for publication**

441 Not applicable for that section.

442

443 **Availability of data and materials**

444 All data generated or analysed during this study are included in this published article and its
445 supplementary information files.

446

447 **Competing interests**

448 The authors declare that no competing interests exist.

449

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453 the design of the study and collection, in analysis and interpretation of data, and in writing the
454 manuscript.

455

456 **Author contributions**

457 AIAH, CM, LN and PG performed technical experiments and statistical analysis. MG and
458 YTC participated in the design of the amyloid fibrils' labelling method. AIAH and PG
459 participated in the design of the study and data analysis. PG conceived the study and
460 coordinated it. AIAH and PG wrote the initial draft of the manuscript. All authors read and
461 approved the final manuscript.

462

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553

554 **Figure legends**

555 **Figure 1. Characterization of calibrated inocula of *Staphylococcus aureus* biofilm and**
556 **planktonic cultures**

557 **(A and B)** SEM micrographs of *S. aureus* LYO-S2 planktonic **(A)** and 24 h biofilm **(B)**
558 inocula after passing through the 34G needle used for micro-injections. Red arrows in panel B
559 indicate the biofilm extracellular matrix. Scale bar: 5 μ m. **(C and D)** Fluorescence
560 microscopy images of *S. aureus* biofilm **(C)** and planktonic **(D)** cultures stained with the
561 green live cell fluorescent label SYTO9 and incubated with CDy11 red fluorescent probe.
562 Scale bar: 50 μ m.

563

564 **Figure 2. Micro-injection of calibrated inocula of *Staphylococcus aureus* in the mouse**
565 **ear pinna**

566 **(A to C)** Reconstituted confocal images of the mouse ear pinna tissue showing the maximal
567 projection intensities of the EGFP signal. LyM-EGFP transgenic mice were micro-injected
568 with TS culture medium **(A)** or *S. aureus* mCherry-LYO-S2 in its planktonic **(B)** or biofilm
569 **(C)** form at early (4-7 hpi) and late time points (after 24 hpi). The EGFP fluorescence (green)
570 signal corresponds to phagocytic cells (neutrophils and macrophages). The yellow line
571 indicates the ROI where the “Sum of EGFP fluorescence intensities” was measured. Scale
572 bar: 2 mm. One representative experiment is shown for each group of mice from four
573 independent experiments. **(D)** Ratio of the sum of EGFP fluorescence intensities to ROI area.
574 Data are expressed as median and interquartile ranges for four mice per group.

575

576

577 **Figure 3. Dynamics of recruited EGFP+ cells in the mouse ear pinna after inoculation of**
578 *Staphylococcus aureus*

579 **(A and B)** Live confocal imaging after micro-injection of *S. aureus* mCherry-LYO-S2 in its
580 planktonic **(A)** or biofilm **(B)** form in the ear pinna of LysM-EGFP transgenic mice at early
581 time points. Innate immune cell recruitment towards planktonic bacteria and biofilms was
582 observed between 3.20 to 3.50 hpi and 4.20 to 4.40 hpi, respectively. A progressive
583 recruitment of EGFP+ innate immune cells was observed at the injection site with cell-
584 bacteria contact areas (filled white arrowheads). White empty circles show cell accumulation
585 over time for the planktonic or biofilm inoculum at early time points. *: autofluorescent hair
586 (also in magenta). Scale bar: 100 μ m. **(C and D)** Live confocal imaging at late time points
587 after micro-injection of planktonic **(C)** or biofilm **(D)** bacteria, at 24.20 hpi and 26.20 hpi,
588 respectively. Empty white arrowhead indicates the presence of remaining planktonic form
589 after 24 h (low magenta signal) whereas biofilms were still easily detectable. Scale bar: 100
590 μ m. **(A to D)** Images show average intensity projections of green (innate immune cells) and
591 magenta (bacteria) fluorescence. One representative experiment is shown for each group of
592 mice from three independent experiments.

593

594 **Figure 4. Motility of recruited EGFP+ cells in the mouse ear pinna after micro-injection**
595 *of Staphylococcus aureus*

596 **(A and B)** Illustration of immune cell tracking with Imaris software using the “Spots” tool to
597 analyse the motility of recruited immune cells. The analysis was carried out in different zones
598 of the injection site where cells were either in contact with visible bacteria **(A)** or not **(B)**.
599 Each cell is represented by a white sphere and its trajectory in the thickness of the tissue by a

600 multicoloured line. Images shown were taken at 4.45 hpi (**A**) and 26 hpi (**B**). *: base of hair
601 follicles. Scale bar: 50 μ m.

602 (**C to H**) Average speed and straightness of EGFP⁺ cells recruited to injection sites at early
603 and late time points after inoculation of TS culture medium (control), planktonic bacteria
604 (planktonic form) or biofilms (biofilm form). Data are expressed as median and interquartile
605 ranges pooled from three different mice in three independent experiments for each group.

606 Average speed (**C**) and straightness (**D**) of all cells (in contact with visible bacteria or not) in
607 infected and control mice. Number of cells (N) analysed for each group at early and late time
608 points, respectively: Control: N = 90 and 94 cells; Planktonic form: N = 315 and 433 cells;
609 Biofilm form: N = 254 and 518 cells.

610 Average speed (**E and G**) and straightness (**F and H**) of cells either in contact (bacteria
611 contact) or not (no bacteria contact) with planktonic or biofilm bacteria at early (**E and F**) and
612 late (**G and H**) time points. Number of cells (N) analysed at early time points that were in
613 contact or not in contact with bacteria, respectively: Planktonic form: N = 157 and 158 cells;
614 Biofilm form: N = 142 and 112 cells. Number of cells (N) analysed at late time points that
615 were in contact or not in contact with bacteria, respectively: Planktonic form: N = 298 and
616 135 cells; Biofilm form: N = 98 and 420 cells.

617

618 **Additional material**

619 **Additional file 1: Figure S1**

620 **Preparation and characterization of calibrated inocula of *Staphylococcus aureus* biofilm**
621 **and planktonic cultures (pdf 5.18Mb)**

622 (A) Titration of 3.8 μ L aliquots of 24 h-old biofilms of *S. aureus* LYO-S2. Data represent
623 mean \pm SD of three samples per well collected from three different wells and prepared in
624 three independent experiments. (B) Titration of *S. aureus* LYO-S2 planktonic and 24 h
625 biofilm inocula on agar plates. Results are expressed as CFU numbers $\times 10^7$ in 3.8 μ L
626 (injection volume). Data represent mean \pm SD from 17 experiments for the planktonic form
627 and from 27 experiments for biofilms. (C) Planktonic inocula after passing through a 34G
628 needle. Scale bar: 10 μ m. (D to H) Biofilm inocula after passing through a 34G needle. Red
629 arrows indicate the biofilm extracellular matrix. Scale bar: 10 μ m (D), 5 μ m (E and F), 2 μ m
630 (G and H).

631 **Additional file 2: Table S1**

632 **Raw data used for Additional file 1: Figure S1A-B. (xls 67 Kb)**

633 Tables presenting raw data used for the preparation of calibrated *Staphylococcus aureus*
634 biofilm and planktonic inocula.

635 **Additional file 3: Table S2**

636 **Raw data used for Figure 3D (xls 67 Kb)**

637 Table presenting raw data used to measure the ratio of the sum of EGFP fluorescence
638 intensities to ROI areas.

639 **Additional file 4: Movie 1**

640 **Immune cells are recruited to injection sites even in the absence of bacterial challenge**
641 **(mp4 9 kb)**

642 *In vivo* confocal time-lapse imaging of immune cell migration in LysM-EGFP transgenic
643 mice ear tissue injected with TS culture medium from 4 hpi to 4.20 hpi. Average projections
644 of time-lapse images. Z-stacks collected 41.76 seconds apart. Scale bar: 100 μ m.

645

646 **Additional file 5: Figure S2**

647 **Dynamics of recruited EGFP+ cells in the mouse ear pinna after micro-injection of**
648 ***Staphylococcus aureus* (pdf 8.04 Mb)**

649 **(A and B)** Confocal images of injection sites after micro-injection of *S. aureus* mCherry-
650 LYO-S2 in its planktonic form in the ear pinna of LysM-EGFP transgenic mice at early time
651 points for two independent experiments. Images of innate immune cell recruitment towards
652 planktonic bacteria were acquired at 5.15 hpi **(A)** and 3.05 hpi **(B)**. **(C and D)** Confocal
653 images of injection sites after micro-injection of *S. aureus* mCherry-LYO-S2 in its biofilm
654 form in the ear pinna of LysM-EGFP transgenic mice at early time points for two independent
655 experiments. Images of innate immune cell recruitment towards biofilms were acquired at
656 4.20 hpi **(C)** and 3.30 hpi **(D)**. Images show average intensity projections of green (innate
657 immune cells) and magenta (bacteria) fluorescence. Filled white arrowheads indicate cell-
658 bacteria contact areas. *: autofluorescent hair (also in magenta). Scale bar: 100 μm .

659 **Additional file 6: Movie 2**

660 **Numerous immune cells penetrate the injection site and interact with planktonic**
661 **bacteria (mp4 1.14 Mb)**

662 *In vivo* confocal time-lapse imaging of immune cell migration in LysM-EGFP transgenic
663 mice ear tissue injected with planktonic bacteria from 3.20 hpi to 3.50 hpi. Average
664 projections of time-lapse images. Z-stacks collected 41.73 seconds apart. Scale bar: 100 μm .

665 **Additional file 7: Movie 3**

666 **Most immune cells arrest at the periphery of injected biofilms (mp4 9kb)**

667 *In vivo* confocal time-lapse imaging of immune cell migration in LysM-EGFP transgenic
668 mice ear tissue injected with planktonic bacteria from 4.20 hpi to 4.40 hpi. Average
669 projections of time-lapse images. Z-stacks collected 45.15 seconds apart. Scale bar: 100 μm

670 **Additional file 8: Table S3**

671 **Raw data used for Figure 4C (xls 164 Kb)**

672 Table presenting the average speed of all cells in infected and control mice at early and late
673 time points. Raw data extracted from Imaris software.

674 **Additional file 9: Table S4**

675 **Raw data used for Figure 4D (xls 224 Kb)**

676 Table presenting the straightness of all cells in infected and control mice at early and late time
677 points. Raw data extracted from Imaris software.

678 **Additional file 10: Table S5**

679 **Raw data used for Figure 4E (xls 125 Kb)**

680 Table presenting the average speed of cells in contact with bacteria or not in infected mice at
681 early time points. Raw data extracted from Imaris software.

682 **Additional file 11: Table S6**

683 **Raw data used for Figure 4F (xls 132 Kb)**

684 Table presenting the straightness of cells in contact with bacteria or not in infected mice at
685 early time points. Raw data extracted from Imaris software.

686 **Additional file 12: Table S7**

687 **Raw data used for Figure 4G (xls 133 Kb)**

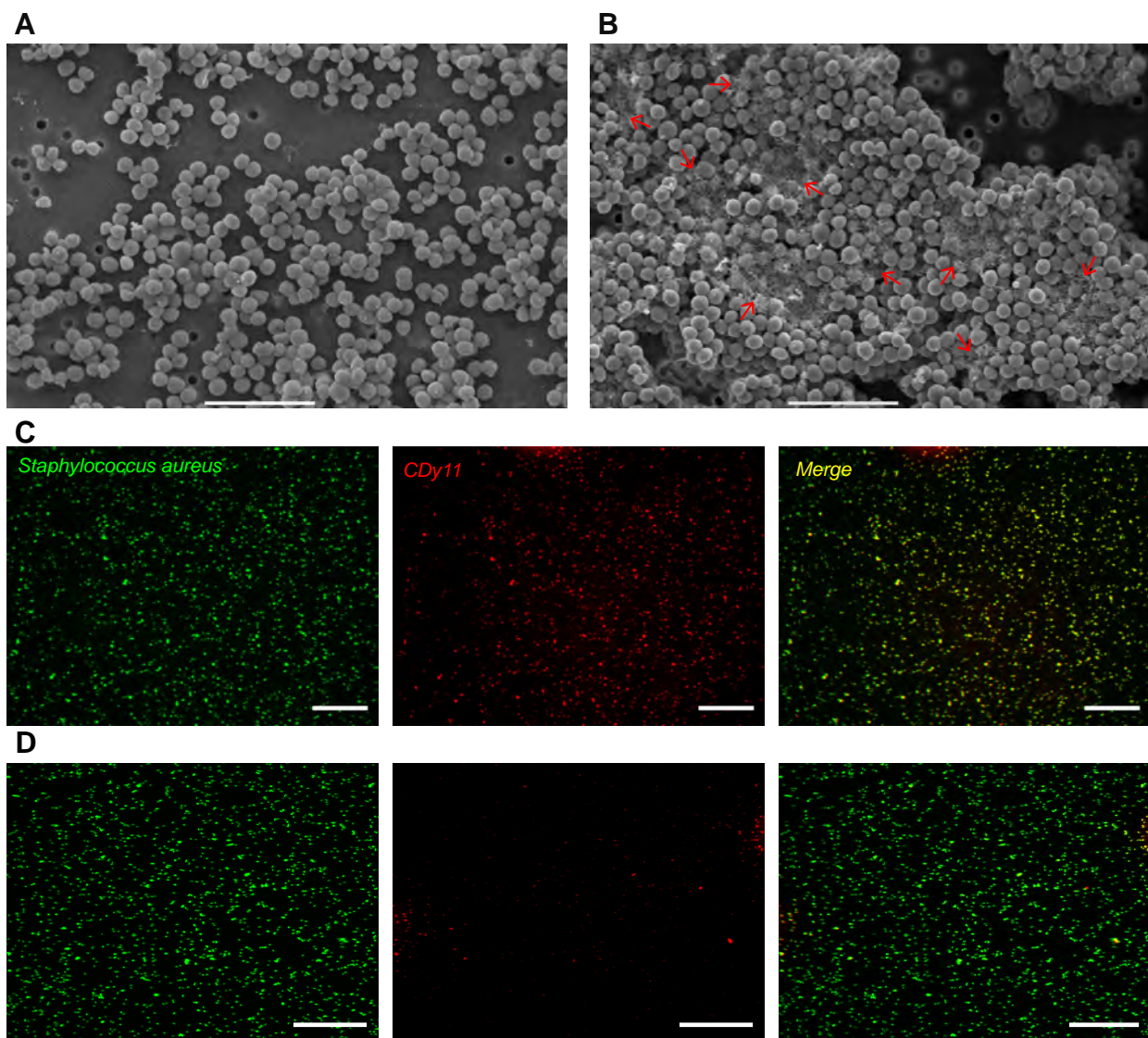
688 Table presenting the average speed of cells in contact with bacteria or not in infected mice at
689 late time points. Raw data extracted from Imaris software.

690 **Additional file 13: Table S8**

691 **Raw data used for Figure 4H (xls 199 Kb)**

692 Table presenting the straightness of cells in contact with bacteria or not in infected mice at
693 late time points. Raw data extracted from Imaris software.

Figure 1. Abdul Hamid et al.



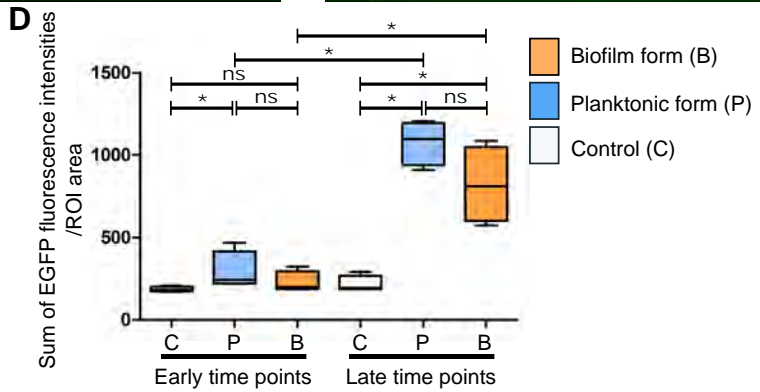
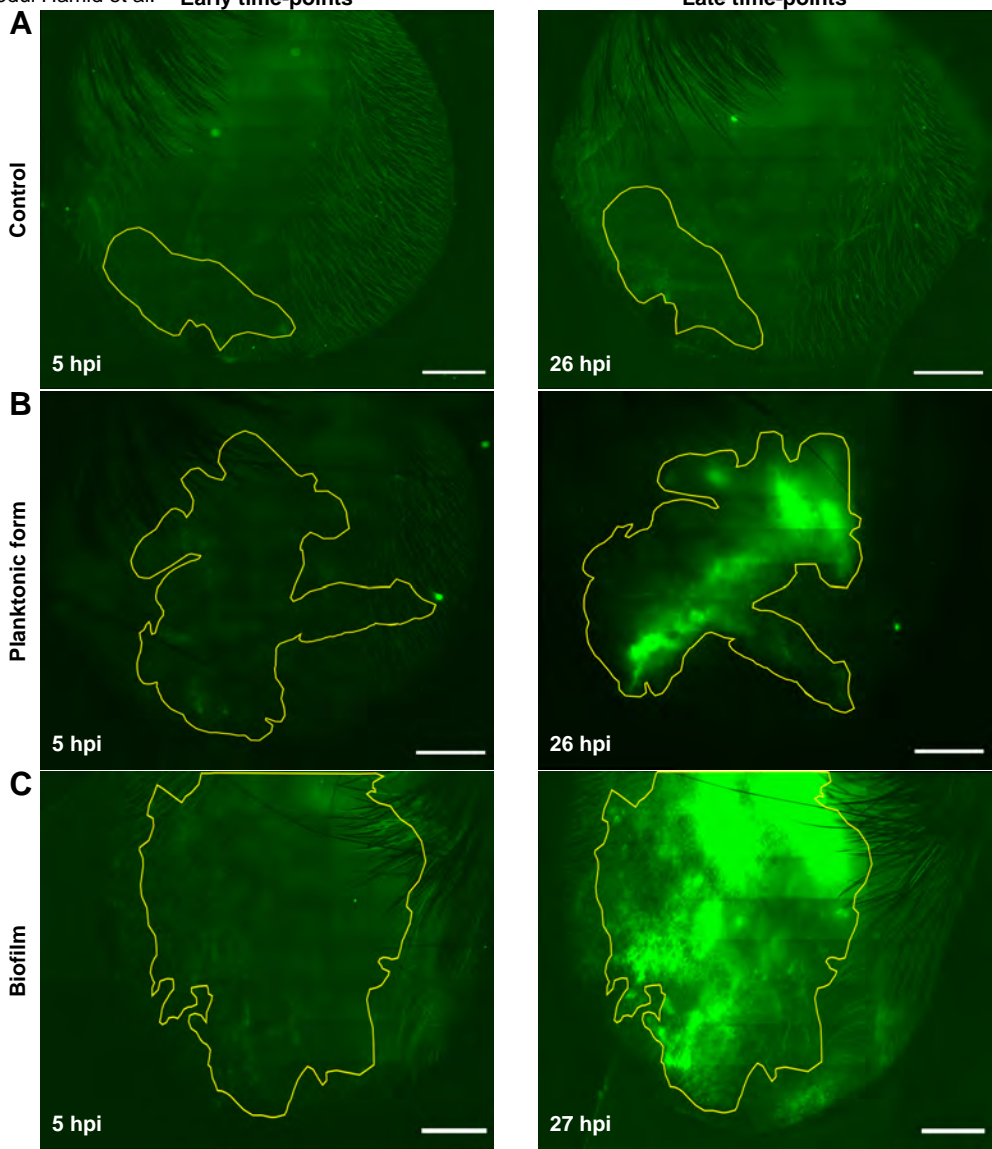


Figure 3. Abdul Hamid et al.

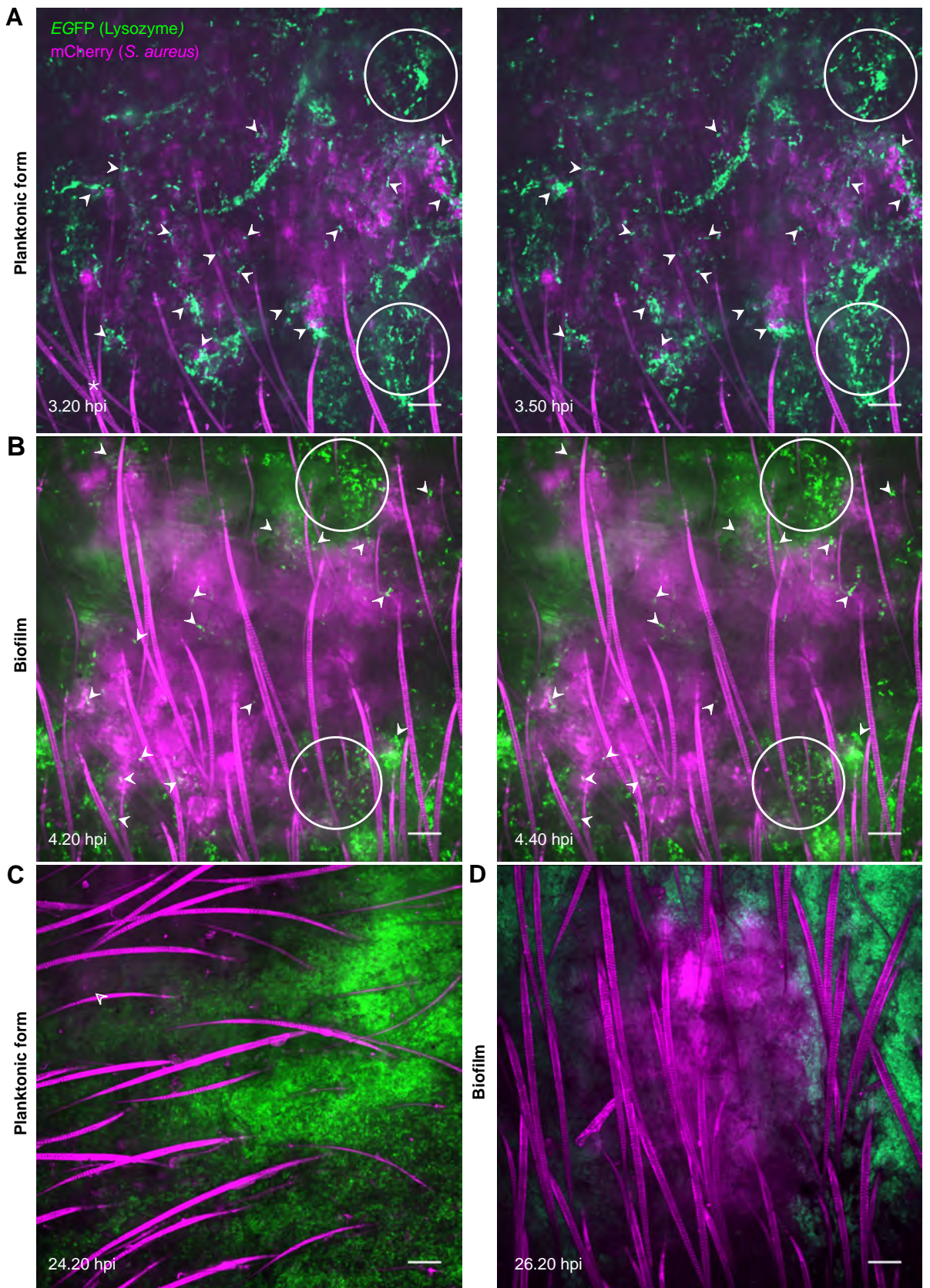


Figure 4. Abdul Hamid et al.

