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# Biofouling of filtration membranes in wastewater reuse: in situ visualization with confocal laser scanning microscopy

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## Abstract

The filtration membranes utilized in advanced treatment of municipal wastewater are prone to biological fouling. Permeability loss and water recovery limitations due to fouling necessitate additional pretreatment and membrane cleaning, which raise the cost of water reuse. Better fouling mitigation calls for a deeper understanding of biofilm behavior on water reuse membranes. This study uses confocal microscopy to monitor biofouling under realistic conditions. To simulate water reuse, a miniature flow cell with microfiltration and ultrafiltration membranes was operated under typical filtration conditions using secondary effluent from California's East Bay Municipal Utility District. The growing biofilms were stained for extracellular polymeric substances and live and dead cells, and they were monitored in situ under filtration conditions with a confocal laser scanning microscope to produce 2-D and 3-D images. In situ image series revealed biofilm growth and removal processes including internal and external fouling, extracellular polymeric substance production, wrinkling, delamination, and film re-deposition. Additionally, changes in biofilm morphology between in situ and ex situ images highlighted limitations of ex situ imaging. These results provide insight into the physical and biological mechanisms of biofouling in hopes of informing the development of improved techniques for biofouling mitigation.

*Keywords:* water reuse, membrane fouling, confocal laser scanning microscopy, ultrafiltration, microfiltration

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## 1. Introduction

While membrane-based potable reuse of municipal wastewater is rapidly expanding as a low-energy method of mitigating water scarcity [1], membrane fouling remains a common

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4 concern. In water reuse facilities, membrane fouling raises water cost by necessitating ex-  
5 tensive pretreatment and frequent membrane cleaning as well as limiting water recovery  
6 [2]. Membrane fouling occurs due to the buildup of various constituents including microbes,  
7 dissolved organic and inorganic species, and suspended solids on the membranes through  
8 which water permeates [3, 4]. Biological fouling (biofouling) is particularly problematic in  
9 water reuse due to the relatively high microbial count and organic content of municipal  
10 wastewater [5]. Fouling can be mitigated to some extent through feed water pretreatment  
11 and specialized membrane coatings, and can also be partially or fully reversed by cleaning  
12 the membrane with detergents, acidic or alkaline solutions, chelating agents, or biocides  
13 [6], but the limited efficacy of current techniques drives further research in fouling preven-  
14 tion. Better understanding of the interplay between the biology and transport phenomena  
15 that dictate biofilm formation [7] and the mechanics that influence biofilm removal has the  
16 potential to inform the development of improved fouling mitigation techniques.

17 Visualization of fouling behaviors under relevant filtration conditions has the potential to  
18 inform the design of innovative fouling mitigation strategies. For example, a novel method  
19 of chemical-free, deformation-induced cleaning [8] of organic-fouled reverse osmosis (RO)  
20 membranes derived inspiration from in situ observation of alginate fouling delamination [9].  
21 Optical coherence tomography (OCT) has been used to create in situ videos and time-lapse  
22 image series of the fouling layer in plane view and in cross section to better understand  
23 biofouling of UF membranes [10, 11, 12]. Epifluorescence microscopy was used with a high-  
24 pressure RO membrane flow cell to visualize the spatial distribution of bacterial deposition  
25 and understand the influence of feed spacers on RO membrane biofouling [13]. Other nonde-  
26 structive optical techniques for biofouling characterization include two-photon femtosecond  
27 imaging [14] and optical microscopy [15], both of which have been used with yeast. Such  
28 visualization studies complement quantitative studies using metrics such as permeate flux  
29 decline and longitudinal pressure drop, which have enabled comparisons of biofouling rates

30 between different operating conditions (e.g., temperature [16]) and processes (e.g., RO vs.  
31 forward osmosis (FO) [17]). Together, quantitative measurements and visualization tech-  
32 niques enable researchers to better understand, predict, and mitigate biofouling.

33 Confocal laser scanning microscopy (CLSM) enables visualization of the composition  
34 and morphology of biofilms [18, 19], and it has occasionally been used in situ to observe  
35 biofilm growth as it occurs. CLSM provides a 3-D view of biofilm morphology, which can  
36 be complex and hard to predict or measure [20], and the various labeling methods avail-  
37 able allow for visualization of biofilm composition and morphology together, especially when  
38 CLSM is combined with other microscopy techniques [21, 22]. For example, activated sludge  
39 supernatant biofilms formed on a glass slide were examined with both CLSM and Raman  
40 microscopy [23] to determine the change in EPS composition over a period of three months.  
41 Fluorescent labeling of extracellular polymeric substances (EPS), which serve a host of pur-  
42 poses in biofilms [24], and other biofilm constituents, such as proteins and cells, enables  
43 CLSM to reveal biofilm structure and behavior. Klausen et al. [25] used in situ CLSM to  
44 create high-resolution, time-lapse 3-D images of the mushroom-shaped structures formed  
45 by *Pseudomonas aeruginosa* on glass and determine the effect of cell motility on biofilm  
46 conformation. Their study tagged mutated and unmutated cells with different fluorescent  
47 proteins to avoid photobleaching and stain cytotoxicity, which pose challenges to long-term  
48 CLSM imaging of live cells. Due to its ability to differentiate species and strains, the fluo-  
49 rescent in situ hybridization (FISH) method of fluorescence-tagging particular DNA or RNA  
50 sequences has been particularly successful in enabling CLSM to elucidate the structure and  
51 dynamics of multispecies biofilms [26, 27]. In addition to providing qualitative visualization  
52 of biofilm behavior, analysis of 3-D CLSM images allows for quantification of biofilm volume  
53 and spatial distribution of biofilm constituents [28].

54 CLSM has been used ex situ to examine the structure and composition of biofilms formed  
55 on water treatment membranes, specifically, to complement macroscopic fouling metrics such

56 as flux decline. For example, Farias et al. [29] collected ex situ CLSM images of biofoul-  
57 ing of RO membranes treating membrane bioreactor effluent to reveal changes in biofilm  
58 morphology and composition over time. Kwan et al. [17] visualized biofilm conformation ex  
59 situ on RO and forward osmosis (FO) membranes and found that the harvested FO biofilms  
60 were thicker and more heterogeneous in shape than those grown in RO, which enabled  
61 interpretation of the mechanisms for the slower flux decline typically observed in FO.

62 Although CLSM has previously been used in situ for non-membrane biofilms (e.g., in Ref.  
63 [25]), permeation through a membrane has significant effects on biofilm thickness and shape  
64 [30], and thus it is important to study membrane biofouling under permeation conditions.  
65 Bar Zeev et al. [31] examined the effect of a subtle change in sample preparation for ex situ  
66 biofilm imaging—the addition of a glass coverslip—and found significant changes in biofilm  
67 conformation occurred. Ex situ visualization of biofilm morphology, while capable of high  
68 resolution, is limiting because removal from the environment in which the biofilm was grown  
69 has the potential to cause conformational changes. However, to our knowledge, biofilms  
70 grown under permeation conditions have only been imaged with CLSM ex situ due to the  
71 challenge of imaging pressurized liquids with optical microscopy.

72 In this study, CLSM was used to visualize biofouling of water treatment membranes  
73 in situ and *operando*, i.e., under conditions of permeation and cross-flow. Biofouling in  
74 microfiltration (MF) and ultrafiltration (UF)—key filtration processes in water reuse that  
75 typically precede RO [2]—were simulated using a transparent, pressurizable, miniature flow  
76 cell outfitted with an MF or UF membrane. Secondary effluent from a municipal wastewater  
77 treatment plant was filtered through the flow cell under cross-flow and flux conditions typi-  
78 cal of water reuse facilities, including frequent membrane backwashes. This novel method of  
79 in situ CLSM imaging of membrane biofouling under permeation conditions allows for visu-  
80 alization of hydrodynamic and biological mechanisms of biofouling. We hope these images  
81 will enhance understanding of biofouling behavior and inform the development of future

82 biofouling mitigation strategies for water reuse.

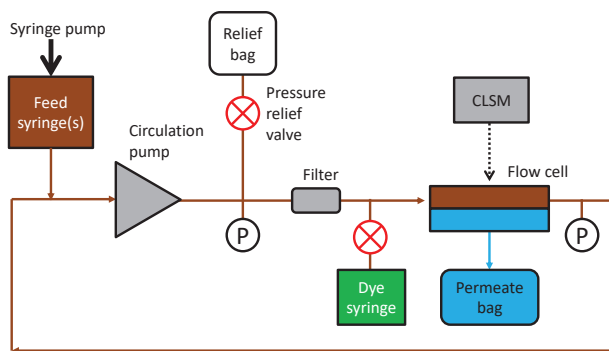
## 83 **2. Methods**

84 In this study, the MF/UF step in municipal wastewater reuse was simulated using a  
85 custom-designed flow cell. Secondary municipal wastewater effluent was circulated through  
86 a transparent, pressurizable cross-flow filtration cell designed for in situ optical probing with  
87 CLSM.

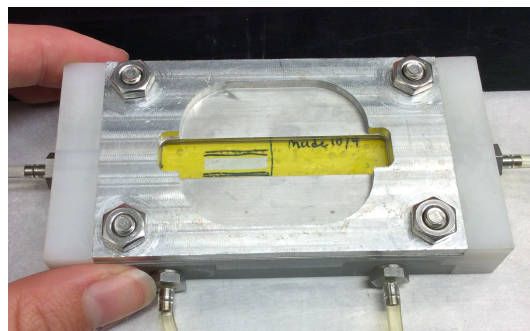
### 88 *2.1. Apparatus*

89 The experimental apparatus, which is depicted in Fig. 1, allowed for pressurized filtration  
90 of municipal wastewater effluent through MF and UF membranes under conditions typical  
91 of wastewater reuse while allowing for in situ visualization of membrane fouling. A custom  
92 flow cell (Fig. 1b) was designed to hold a membrane between a feed channel containing a feed  
93 spacer (Sterlitech 31 mil diamond), which enabled backwashing of the flat sheet membrane,  
94 and a permeate channel containing a permeate carrier cut from a commercial spiral-wound  
95 RO membrane (Membrane Solutions).

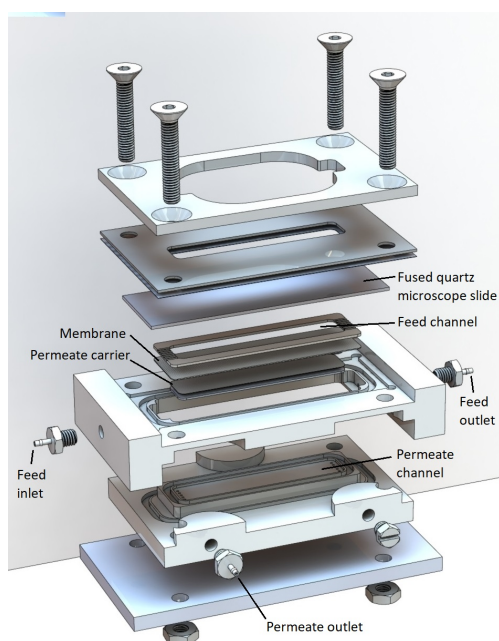
96 The cell was designed to incorporate a standard 1 mm-thick, 1"×3" (approximately  
97 25×75 mm) glass or quartz microscope slide; quartz was used in this study for its supe-  
98 rior shatter resistance. In preliminary tests, quartz slides in the experimental apparatus  
99 withstood 3–6 barg of pressure before shattering, which was sufficient to operate filtration  
100 through MF and UF membranes, and the 1 mm slide was thin enough to allow biofilm  
101 visualization with long working distance objectives. The small opening in the metal plate  
102 above the slide, the plastic sheet between the slide and the plate, and the O-ring grooves  
103 were designed to minimize stress concentration on the slide. However, despite careful design  
104 and the use of a pressure relief valve in the flow loop, the slide occasionally cracked during  
105 long experiments, requiring cleaning the apparatus and starting the trial over from the be-



(a)



(b)



(c)

Figure 1: Experimental apparatus: (a) schematic diagram of flow system, where P represents a pressure gauge and a crossed circles represents a valve; (b) photograph of the flow cell; (c) exploded view of flow cell design (sealing O-rings not pictured).

106 ginning. Researchers wishing to replicate or adapt the flow cell used in this work may do so  
 107 by downloading the SolidWorks files provided in the Supplementary Information.

108 Permeate flux was controlled by a syringe pump (New Era Pump), which injected liquid  
 109 into a fluidic system of approximately constant volume (with some variation due to tubing  
 110 flexibility and pressure changes), so that the flow rate through the membrane was approxi-

111 mately equal to the syringe pump flow rate. Backwashing was conducted by programming  
112 the syringe pump to periodically withdraw liquid from the flow loop, thus pulling permeate  
113 back through the membrane into the feed.

114 Cross-flow velocity in the flow cell was set by a magnetic-drive circulation pump (Ismatec  
115 drive with Micropump head) that was calibrated with a rotameter before fouling tests. A 25  
116  $\mu\text{m}$  filter was installed in the flow loop to catch pieces of biofilm removed from the membrane  
117 during the backwash step, since these would be swept away during the backwash step in a  
118 real treatment facility.

119 The flow system was cleaned and disinfected between trials by flushing effluent out with  
120 water; disassembling the flow cell and scrubbing parts with soap and water; flushing the  
121 reassembled system with ethanol; flushing the ethanol out with tap water; and, finally,  
122 flushing the tap water out with deionized water.

## 123 *2.2. Membranes*

124 Commercial membranes were used for both filtration processes. The UF membrane  
125 was a polyethersulfone (PES) 10 kDa molecular weight cutoff (MWCO) Microdyn Nadir  
126 UF membrane. The MF membrane was a polyvinylidene fluoride (PVDF) membrane from  
127 Synder (V0.1) with a nominal pore size of 0.1  $\mu\text{m}$ . SEM images of both membranes are  
128 provided in Appendix C. Although hollow fiber MF and UF membranes are more common  
129 in practice than flat sheet membranes [2], flat sheet membranes were used in this study to  
130 enable plane views of fouling accumulation and removal processes.

131 To define the active area and prepare membranes for use in the miniature flow cell,  
132 membrane coupons were first laid between sheets of 0.001" (0.025 mm) adhesive-backed  
133 polyimide film, from which rectangles had been removed, in order to provide a smooth surface  
134 against which the flow cell's O-ring could form a seal. Nail polish, an adhesive commonly  
135 used to seal microscope slides, was carefully applied to the edges of the rectangular cutout on  
136 the active surface of the membranes to prevent permeation through any defects that might

137 have been introduced at the contact line between the membrane and the adhesive film. The  
138 region of the active layer not covered with adhesive film or nail polish was considered the  
139 active area for calculating permeate flux. All membranes were prepared with active areas of  
140 approximately 0.25 cm<sup>2</sup> to enable visualization of several regions between spacer filaments;  
141 active areas for calculating flux were calculated from microscope images.

142 Before installing in the flow cell, UF membranes were dipped in 1:1 ethanol:water solution  
143 for approximately 10 s to wet the support layer and enhance permeability before rinsing  
144 with deionized water and installing in the flow cell; MF membranes were only rinsed with  
145 deionized water.

### 146 *2.3. Wastewater effluent*

147 Several samples of secondary wastewater effluent prior to disinfection were obtained from  
148 clarifier outflow at the East Bay Municipal Utility District (EBMUD) wastewater treatment  
149 plant in Oakland, California, USA. Samples were stored at 4±1 °C and used within five weeks  
150 of collection. Wastewater composition was characterized by CalTest Labs (USA; Table 1).  
151 Cell counts on Luria broth agar plates after 48 hours at room temperature found the cell  
152 count to range from about 5,000–30,000 CFU/ml, a fluctuation which is similar in magnitude  
153 to the seasonal fluctuations seen in plate counts of other treatment plants [32]. However,  
154 Ref. [32] found that the typical cell count by flow cytometry was 3–4 orders of magnitude  
155 higher than plate counts due to the difficulty of growing wastewater effluent bacteria on  
156 plates, so it is likely that the density of cells in the wastewater effluent was higher than  
157 determined by plate counts. It is likely that the wastewater microbial composition changed  
158 both between samples taken in different seasons and between collection and testing, which  
159 we acknowledge is a limitation of this study.

Table 1: Wastewater effluent composition at approximately one week of sample age

Characteristic	Value
pH	7.2
Total dissolved solids (TDS) (mg/L)	970
Total suspended solids (TSS) (mg/L)	<2
Total nitrogen (mg/L)	72
Nitrate and nitrite (mg/L)	<0.07
Dissolved organic carbon (DOC) (mg/L)	20
Biological oxygen demand (BOD) (5-day) (mg/L)	7
Phosphorus (mg/L)	6.0

160 *2.4. Fouling and cleaning*

161 Operating conditions were chosen to mimic realistic MF/UF operation to the extent  
 162 possible in the miniature flow cell. MF and UF were operated at average fluxes (during  
 163 permeate production) of 300 and 80 LMH, respectively, except where noted, and with a feed  
 164 flow rate chosen to create a cross-flow velocity of 25 cm/s.

165 When backwashes were employed, both membrane types were operated with backwash  
 166 cycles every 45 min for 2.25 min at equal and opposite flux as the permeate production step  
 167 to simulate an overall backwash-limited recovery ratio of 90%, which is within the typical  
 168 range of 90–95% [33]. Each trial with backwash continued for long enough for a clear and  
 169 consistent fouling pattern to develop (2–4 days); new room-temperature wastewater was  
 170 added each day. Additional trials were conducted for shorter times to elucidate fouling  
 171 behavior in the absence of backwashing.

172 *2.5. Staining*

173 Stains were selected to differentiate live cells from dead, highlight EPS, and be distin-  
 174 guishable by CLSM. DAPI (4',6-diamidino-2-phenylindole) or Hoechst 33342 (referred to as  
 175 Hoechst in the following) are cell-permeable live cell stains that were used (individually)  
 176 to dye all cells; propidium iodide (PI), which is not cell-permeable, was used to stain dead



177 cells only. Wheat germ agglutinin–Alexa Fluor 555 conjugate (referred to as WGA in the  
178 following) was chosen as a stain to highlight EPS because it adheres to *N*-acetylglucosamine  
179 and sialic acid [34] and was found to highlight EPS in a wide range of bacterial biofilms  
180 by Neu and Kuhlicke [35]. However, because EPS includes a wide range of polysaccharides  
181 and proteins that are not all bound by WGA, WGA cannot highlight all EPS. Due to the  
182 limitation of selecting a few stains that are distinguishable by CLSM and the common de-  
183 sire to additionally stain cell DNA, fluorescence-based biofilm imaging studies often choose  
184 a single lectin to stain glycoconjugates of EPS and stand in for an EPS stain (e.g., *Aleuria*  
185 *aurantia* in Ref. [28]). In the present study, WGA was chosen as a sole lectin to capture  
186 the general shape of EPS accumulation at scales larger than the optical resolution of the  
187 present imaging method (several  $\mu\text{m}$ ). Although EPS will be used as shorthand for the parts  
188 of the biofilm highlighted by fluorescence-tagged WGA in the remainder of this paper, it is  
189 possible for there to be additional EPS that is not bound by WGA and thus not visualized.

190       Approximately 1 h before imaging, concentrated dyes (1 mg/mL for Hoechst, WGA, and  
191 PI and 1-14 mM for DAPI) were added to approximately 0.5 mL of deionized water and  
192 injected slowly into the flow loop upstream of the flow cell. Live cell dyes were sometimes  
193 used throughout the duration of experiments and sometimes applied only at the end in order  
194 to determine whether dye toxicity affected fouling behavior; as discussed in Appendix A,  
195 no marked differences were seen. Stain concentrations stated in the results section (on the  
196 order of  $\mu\text{g}/\text{mL}$ ) are based on an estimated feed loop volume of 17 mL; concentration of  
197 stains in the biofilm may be greater due to permeation of stains through the biofilm as water  
198 flows through the membrane. Stain concentrations and contact times were chosen through  
199 preliminary experimentation to find the lowest concentrations that would allow for clear  
200 imaging and the lowest contact time (about 30 min) that would allow for equilibration of  
201 image brightness. Final (post-dilution) concentrations were within the ranges recommended  
202 for individual stains in their respective manuals [36, 37, 38, 34]; the chosen wait time of 30

203 min met or exceeded recommended minimum incubation times for all stains. The 30-min  
204 wait time also allowed for adequate mixing of injected stains with the wastewater effluent  
205 by circulation more than 200 times through the flow loop.

206 Because injection occurred upstream of the flow cell while filtration was occurring, the  
207 stained biofilm formed through two mechanisms: The existing biofilm was stained as con-  
208 vection and diffusion brought unattached stain molecules to the biofilm, and also water  
209 constituents (unattached microbes, carbohydrates, etc.) may have been stained and then  
210 assembled into a biofilm. Any water constituents that were stained and that did not form  
211 the biofilm either passed through the membrane (and thus were not imaged) or stayed in  
212 the bulk solution, where their concentration was low enough compared to the biofilm to  
213 not meaningfully affect the images. The concentration of dissolved organic carbon (a lower  
214 bound on the concentration of organic water constituents) in the wastewater effluent was 20  
215 mg/L (Table 1), whereas the stains were an order of magnitude lower in final concentration  
216 (typically 1-3 mg/L), so stains should not have significantly interfered with the buildup of  
217 water constituents into a biofilm. That said, the potential for interference of stains with  
218 biofilm formation is a possible limitation of staining-based in situ imaging, as discussed in  
219 Sec. 3.5.3.

220 The membranes used in this study were largely permeable to the stains, with one ex-  
221 ception. Although the MF membrane's nominal pore size of 0.1  $\mu\text{m}$  far exceeds the size of  
222 all four stains, allowing unattached stain molecules to pass through so as not to affect the  
223 biofilm images, the stains adhered to the clean MF membranes somewhat, as seen in CLSM  
224 images (e.g., in Fig. A.20) at the beginning of fouling trials. The 10 kDa UF membrane's  
225 pores are large enough to pass PI (at 668 Da), DAPI (at 277 Da), and Hoechst 33342 (at 453  
226 Da), but not WGA (at 38 kDa), so it is possible for WGA to build up on the UF membrane  
227 independently of the biofilm, as will be discussed in Sec. 3.4. However, 3-D views of UF  
228 fouling do not show disproportionate accumulation of WGA on the membrane surface (see,

229 e.g., Fig. 10).

## 230 2.6. Confocal imaging

231 Confocal imaging was conducted using a Zeiss LSM710 confocal laser scanning micro-  
232 scope. 405 nm and 561 nm lasers were used to excite DAPI/Hoechst and PI/Alexa Fluor  
233 555, respectively. Emission detector ranges were selected using ZEN 2011's dye spectrum  
234 database and built-in optimization<sup>1</sup>. The thickness of the fluid channel and enclosing glass  
235 required the use of long-working-distance objective lenses. A 10x objective (10x Plan-  
236 Neofluar 0.30 NA, Carl Zeiss) was used with a pinhole corresponding to one Airy unit for  
237 the lowest wavelength imaged in all trials except where noted. A 50x long working distance  
238 objective (50x Epiplan Neofluar 0.55 NA/DIC, Carl Zeiss) was also used in some trials, but  
239 clearer images were typically obtained with the 10x objective, digital magnification of 5x,  
240 and a slow imaging speed, the combination of which allowed for visualization of some indi-  
241 vidual cells. All confocal images included in this study were captured in situ and *operando*  
242 (i.e., with the membrane in place in the flow cell and with permeation and cross-flow) unless  
243 otherwise noted. Coordinates of imaged regions were recorded relative to the corner of the  
244 membrane active area to enable repeated imaging of the same region.

245 Image collection and initial processing were conducted with ZEN 2011 (Zeiss). Channel  
246 intensities were normalized consistently within each series of images. When estimating  
247 biofilm volume from 3-D stacks, ImageJ was used with a thresholding procedure similar to  
248 that of Staudt et al. [28]: a single threshold was set manually for what minimum intensity  
249 in the WGA channel indicated the presence of biofilm in a time series of 3-D images, and  
250 product of the number of voxels above the threshold intensity and the voxel volume gave

---

<sup>1</sup>Detector ranges varied depending on the set of dyes used as follows: 410–509 nm for Hoechst or DAPI (regardless of other dyes used); 550–579 nm for Alexa Fluor 555 when used with Hoechst and PI; 540–579 nm for Alexa Fluor 555 when used with DAPI and PI; 545–578 nm for Alexa Fluor 555 when used with PI only; 623–719 nm for PI when used with Alexa Fluor 555 and Hoechst or DAPI; and 623–685 nm for PI when used with Alexa Fluor 555 only.

251 an estimate of the total biofilm volume on the region of membrane imaged. Dividing by the  
252 imaged area allowed for calculation of average thickness. However, significant staining of the  
253 MF membrane prevented the calculation of biofilm volume in MF because the membrane  
254 and biofilm could not be distinguished by intensity alone.

255 In each imaging session, a tiled image was collected of the entire fouled region of each  
256 membrane, and higher-resolution images (some in 3-D) were collected at several locations  
257 that were consistent throughout a given fouling trial (see Fig. 3 for an example). With this  
258 combination of images at each time step, we hoped to capture a range of fouling behaviors  
259 at different scales. More images were collected than are presented in this report; the im-  
260 ages shared here were selected with the intention to represent and illustrate features seen  
261 throughout the larger set of images.

### 262 3. Results and discussion

263 The images collected in this study show the progression of biofouling on MF and UF  
264 membranes through in situ visualization of simulated water reuse with real wastewater efflu-  
265 ent. Although periodic backwashing of porous membranes is integral to fouling mitigation in  
266 real water reuse facilities, fouling observation is conducted both with and without backwash  
267 (as described in Sec. 2.4) to elucidate phenomena that occur between and in the absence of  
268 backwash cycles.

#### 269 3.1. Microfiltration without backwash

270 MF of wastewater effluent was conducted for 8 h at 300 LMH flux without backwash  
271 to determine baseline MF fouling behavior. Stains were added with the effluent to reach  
272 concentrations of 1.8  $\mu\text{g}/\text{mL}$  each of Hoechst and PI and 2.9  $\mu\text{g}/\text{mL}$  WGA. Half the initial  
273 amount of dye was added before collecting images at 4 h and 8 h.

274 Figure 2 shows the progression of the biofilm in one region, including magnified images  
275 of the region's center, over the 8 h period. Live and dead cells (in blue and red, respectively)

276 are visible in the magnified images (Fig. 2 d, and f). In Figs. 2a and b, the membrane is  
277 shown before significant biofilm deposition, highlighting the membrane's large (50  $\mu\text{m}$ -scale)  
278 internal pores just below the membrane surface, which appear in the CLSM images and are  
279 also visible in SEM images of the clean MF membrane (Fig. C.25b). Due to convection of  
280 material to the membrane, a flat biofilm containing live cells, dead cells, and EPS develops  
281 over much of the membrane surface. Tiled images of the entire active area (collected but  
282 not included here) show similar behavior across the membrane.

283 The magnified images (Fig. 2 d, and f) show an accumulation of live cells around the  
284 edges of larger internal pores. Cell accumulation in large pores was confirmed in SEM images  
285 of fouled MF membranes (Fig. C.24d). The enhanced image brightness near the pore edge  
286 may be due to the steepness of the pore there and thus the large number of live cells seen by  
287 the long working distance objective, which has poor vertical resolution. The surface biofilm  
288 is less bright above the fouled internal pores, perhaps indicating a difference in external  
289 biofouling behavior where internal biofouling occurs.

### 290 *3.2. Microfiltration with backwash*

291 MF was conducted with wastewater effluent for 4 d at 300 LMH with backwashes every  
292 45 min (as described in Section 2.4) to determine fouling behavior under the influence of  
293 periodic backwashing. Before imaging, dye was added to reach concentrations of 1.8  $\mu\text{g}/\text{mL}$   
294 each of Hoechst and PI and 2.9  $\mu\text{g}/\text{mL}$  WGA. As explained in Appendix A, a similar test was  
295 conducted separately without Hoechst for comparison because of the effects of Hoechst on  
296 cell reproduction [39]; qualitatively, results were similar with and without daily application  
297 of Hoechst.

298 Figure 3a shows the entire fouled membrane after 1 d as well as several close-up images.  
299 Initially, the MF biofilm builds up in a relatively flat layer with some internal fouling, as  
300 observed in MF without backwashing, but two new features emerge: a striped pattern occurs  
301 parallel to the expected direction of flow and a few small cracks appear in the biofilm. Later,

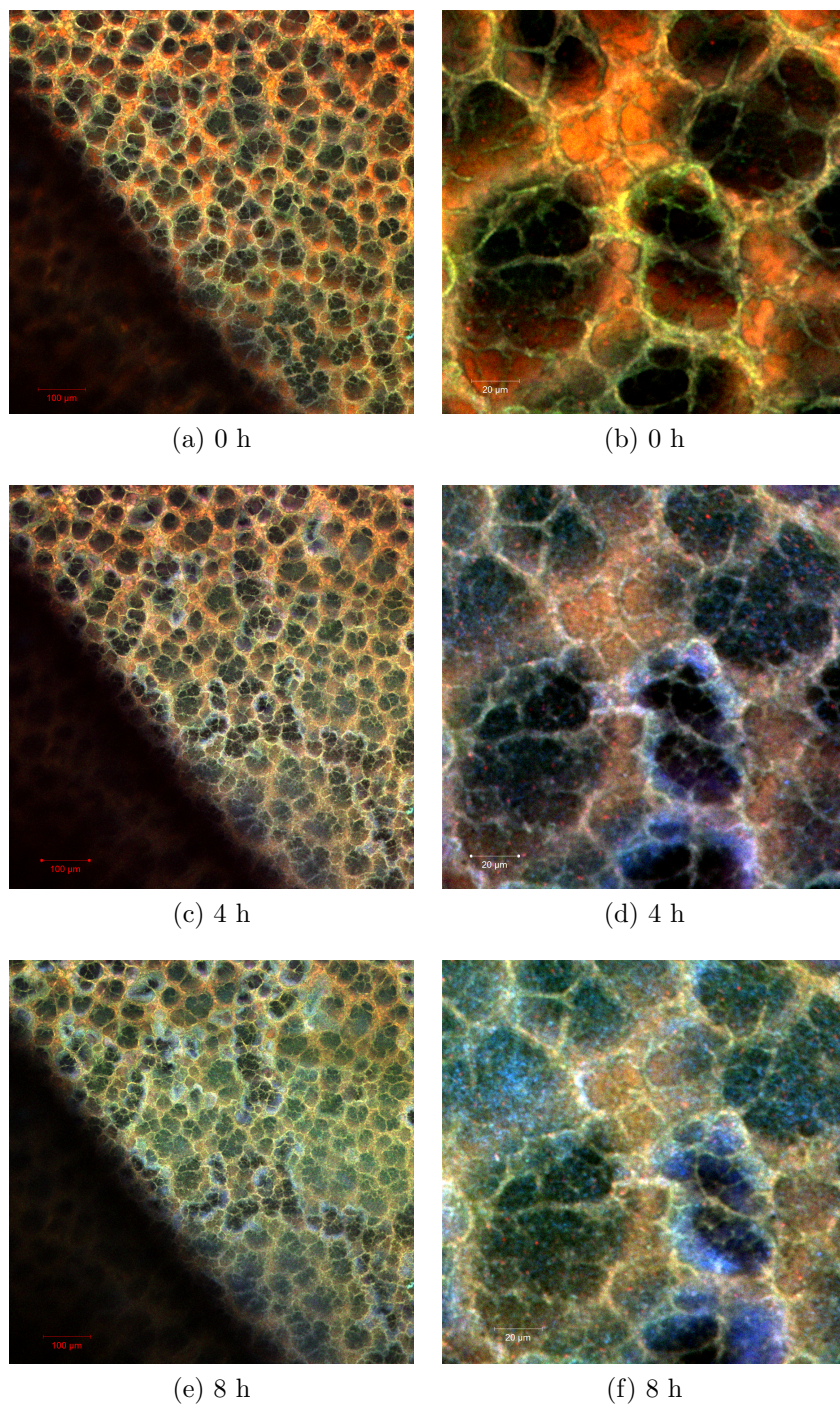


Figure 2: Confocal images of a region of MF membrane after different durations of fouling without backwash. (b), (d), and (f) are magnified images collected at the center of (a), (c), and (e), respectively. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI).

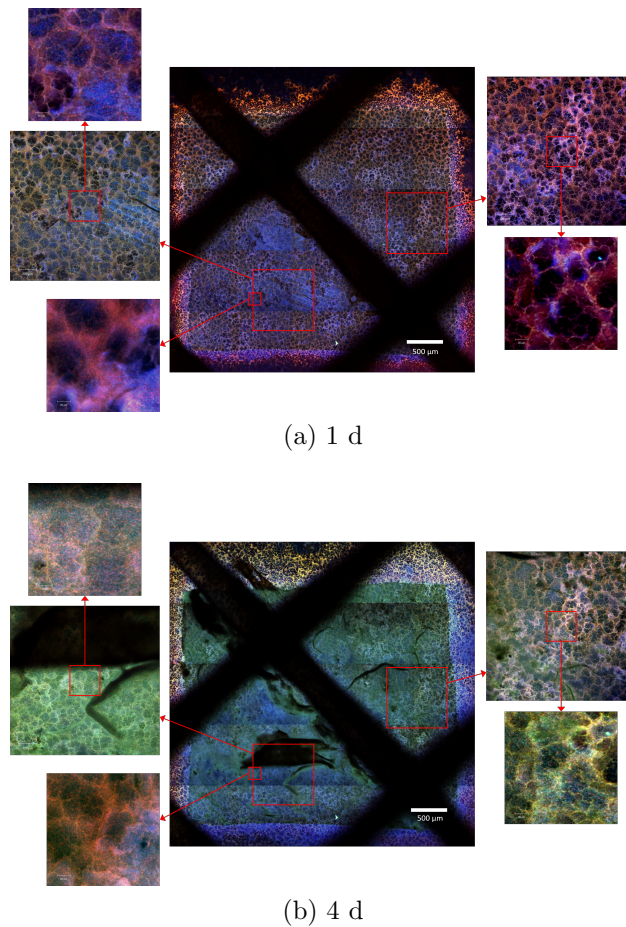


Figure 3: Tiled confocal images of fouled MF membrane after (a) 1 d and (b) 4 d of fouling with higher-resolution images taken at several locations. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI). Cross-flow was from right to left; dark diagonal lines are spacer filaments. Apparent horizontal stripes in the tiled images are an artefact of tiling.

302 two 3-D structures appear: large wrinkles, which appear as dark, curved lines; and flaps,  
 303 which appear as large dark patches. These structures are visible in Fig. 3b, which shows the  
 304 membrane after 4 d of fouling. These structures may appear dark due to extending above  
 305 the confocal plane. The flaps are concentrated downstream of the spacer filament that lays  
 306 flat on the membrane (the diagonal line from upper left to lower right) in arcs, which may  
 307 indicate peeling in the direction of flow from the line of contact between the spacer and the  
 308 membrane. Both flaps and wrinkles grow and move over time. The remainder of the biofilm  
 309 remains relatively flat; refer to Fig. 17a for a 3-D image of a section of this biofilm after 4 d.



310 Dark spots—both holes in the biofilm and small domes where the film appears to be locally  
311 detached from the membrane—also occur, and are more visible in cross-section (Fig. 17a).  
312 Holes and wrinkles in the biofilm are also visible in SEM images of fouled MF membranes  
313 (Fig. C.24a,c).

314 Magnified images of the MF biofilm (Fig. 4) reveal stages of biofilm growth above and  
315 below the active layer. Individual dead cells (red specks) are visible, appearing primarily  
316 on the upper surface of the membrane, particularly in regions between large internal pores.  
317 Live cells (in blue) congregate in the vicinity of several internal pores, densely populating the  
318 rightmost pore over time and leaving a dark hole in the middle. EPS (green) is less obvious  
319 in Fig. 4, but the color shift over time from red to orange on the membrane surface and  
320 blue to turquoise in the open pore indicates EPS buildup over time both on the membrane  
321 surface and in the internal pore.

322 Figure 5 separates the image of the fouled MF membrane at 4 d (Fig. 4d) into individual  
323 detector channels, corresponding to individual dyes, to more clearly show the distribution  
324 of biofilm constituents on the membrane. Live cells (i.e., specks of light that appear in Fig.  
325 5a but not Fig. 5b) can be seen congregating in the lower right, while dead cells are spread  
326 more evenly over the membrane. Dead cells appear brighter in Fig. 5b (the PI channel)  
327 than Fig. 5a (the Hoechst channel) due to the use of minimal levels of the cell-permeable  
328 stain Hoechst. EPS appears cloudy relative to the cells, and is brighter where there are  
329 more live cells. The ratio between total cells and dead cells in the internal pore, which  
330 appears light blue in the lower-right portion of Fig. 4, is shown to be higher than that on  
331 the membrane surface by the intensity profiles shown in Fig. 5d, suggesting the occurrence  
332 of cell proliferation within the pore.

333 The apparent internal fouling of the MF membrane shown in Figs. 2, 4, and 5 is confirmed  
334 by CLSM images of the edges of MF membranes, 3-D CLSM scans, and ex situ SEM images.  
335 At the edges of the active membrane area, which were sealed as described in Section 2.2,



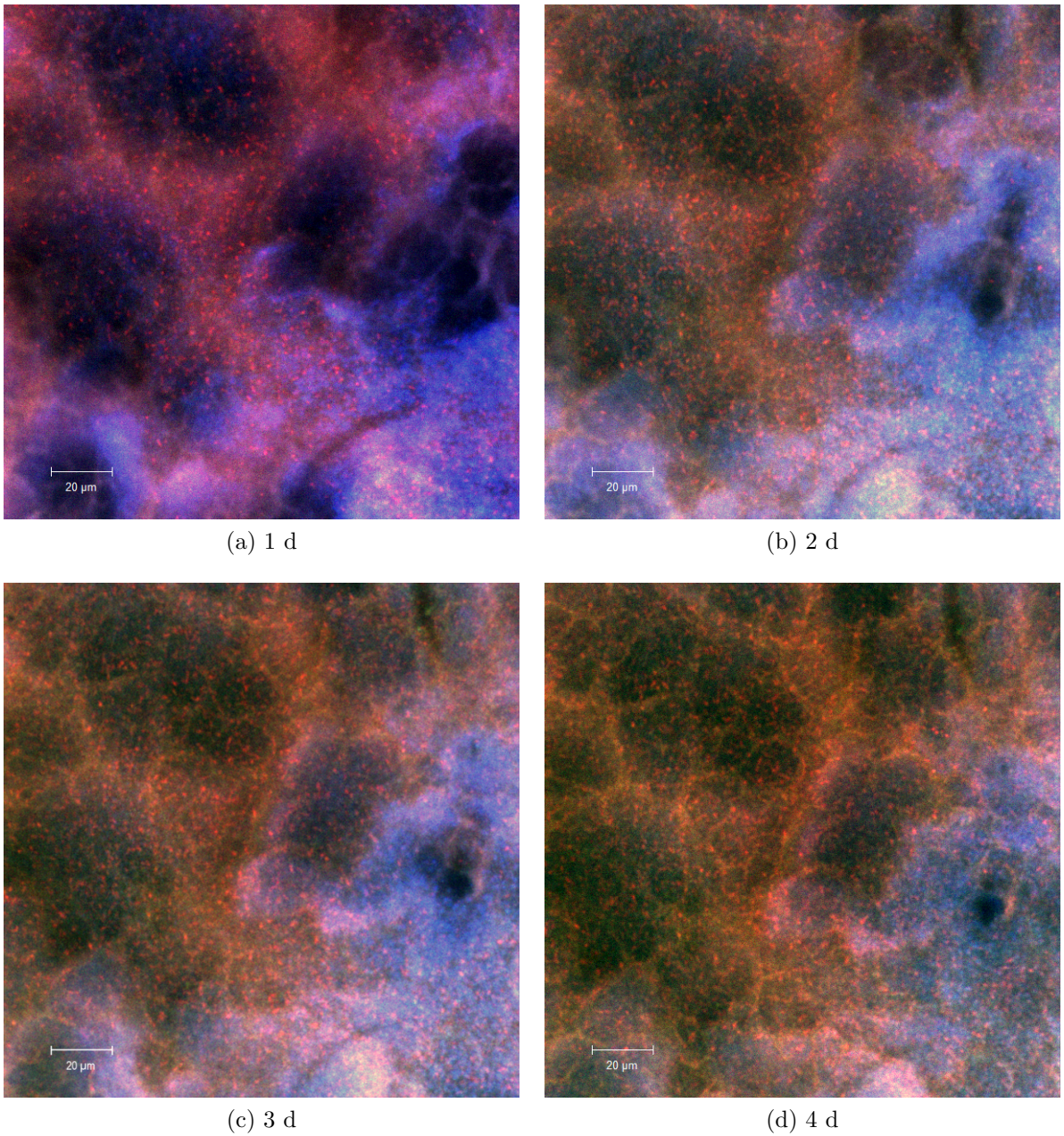


Figure 4: High-resolution confocal images of fouled MF membrane shown in Fig. 3, collected after 1–4 d of fouling with 50x objective. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI).

336 EPS can be seen accumulating beneath the sealed surface along the edges of networks of  
 337 large pores beneath the active layer in Fig. 6. The biofilm appears to grow easily within a  
 338 given network of connected internal pores, perhaps beginning with a large-enough hole in the



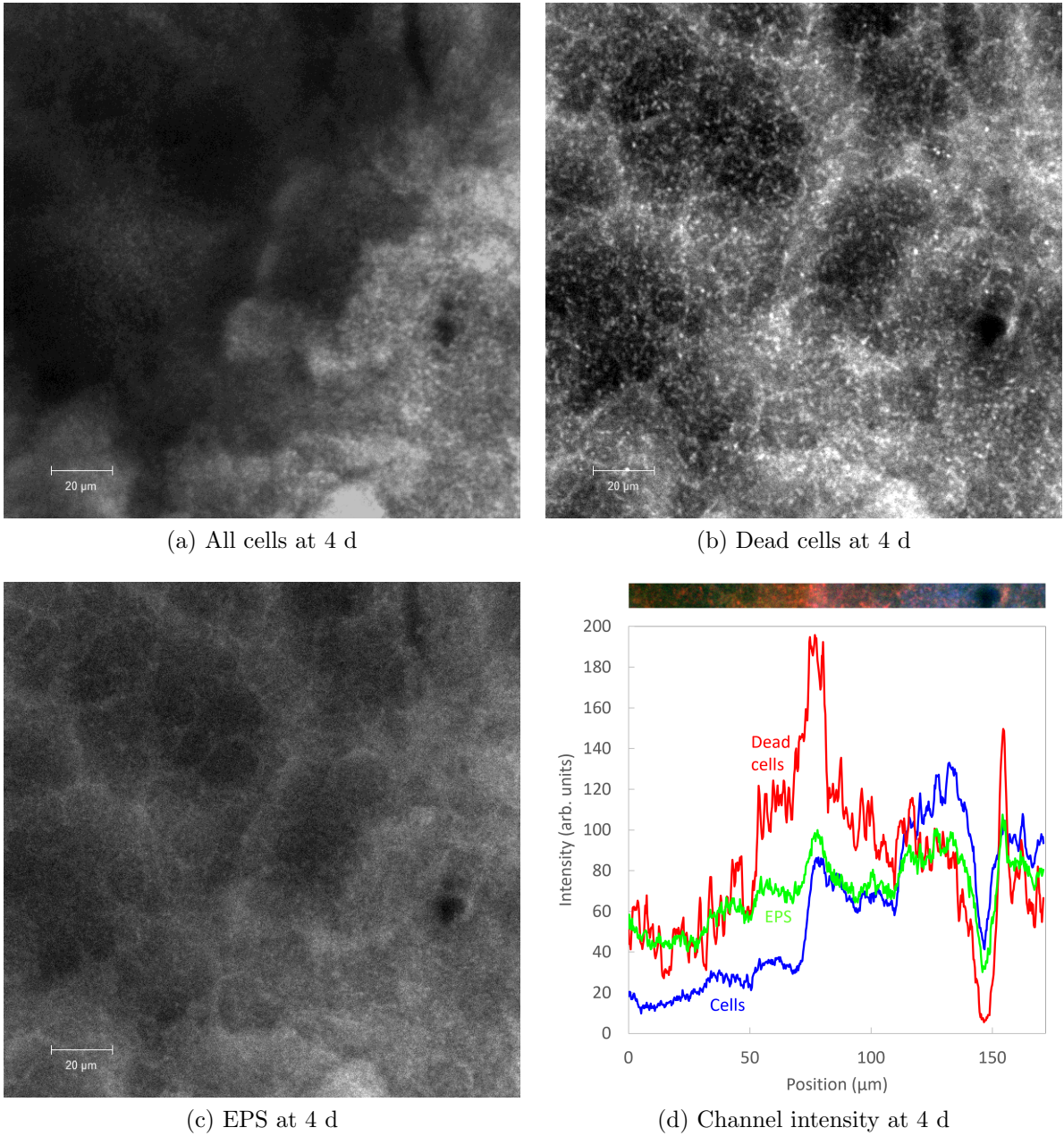


Figure 5: (a–c) The fouled MF membrane of Fig. 4d (at 4 d of age) visualized using separate detector channels for each dye to show the distribution of biofilm constituents. (d) Plot of intensity at 4 d along the horizontal region of Fig. 4d shown above the plot, which intersects the dark spot seen in the micrographs.

339 unsealed membrane surface for bacteria to pass through; such holes are shown in an SEM  
 340 image of an unfouled MF membrane (Fig. C.25a). Bacteria are also seen lining these large  
 341 internal pores in ex situ SEM images (Fig. C.24d) and a 3-D CLSM image (Fig. 7, below) of

342 the same fouled membrane imaged by SEM, in which cells and EPS can be seen extending  
343 below the surface of the membrane within some relatively large internal pores. These images  
344 demonstrate that internal biofouling, which is difficult to remove through backwashing, can  
345 be significant in MF.

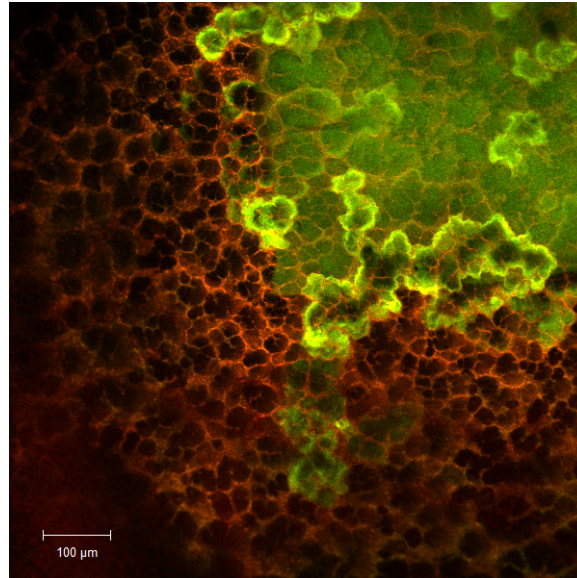


Figure 6: Internal fouling under top-sealed edges of active MF membrane region. This membrane was fouled for 3 days without Hoechst as described in Appendix A. Green = EPS (WGA); red = dead cells (PI).

346 The efficacy of backwashing in cleaning this MF membrane was limited. A separate  
347 experiment was conducted to visualize the membrane before and after backwashing, with  
348 results shown in Fig. 8. Comparing the before and after images, this particular backwash  
349 cycle was ineffective at removing both internal and external fouling; one flap of biofilm in  
350 the middle of the frame moved slightly, but the film largely stayed attached, with several  
351 flaps peeling away from the membrane in the direction of flow. This result highlights the  
352 cohesiveness of biofilms and the difficulty of removing a biofilm adhered to a flat sheet mem-  
353 brane. This result also highlights a limitation of the experimental backwashing procedure:  
354 it did not include air scouring, which would typically accompany backwash in water reuse  
355 facilities [33].



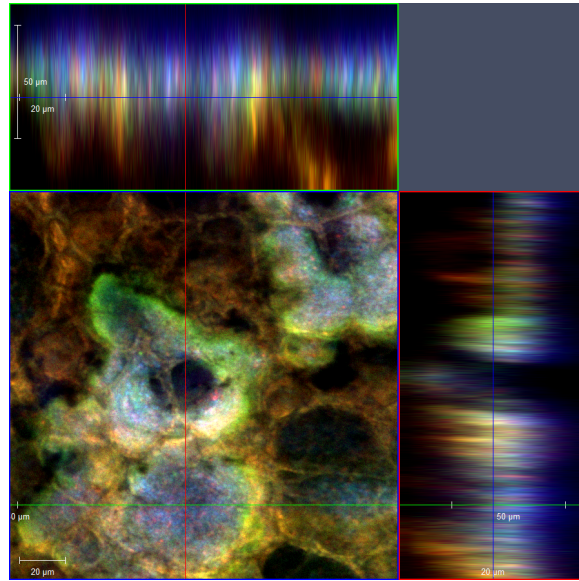
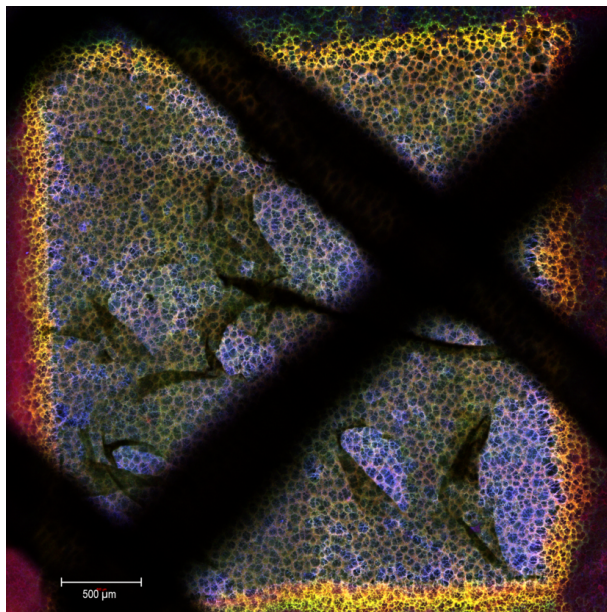
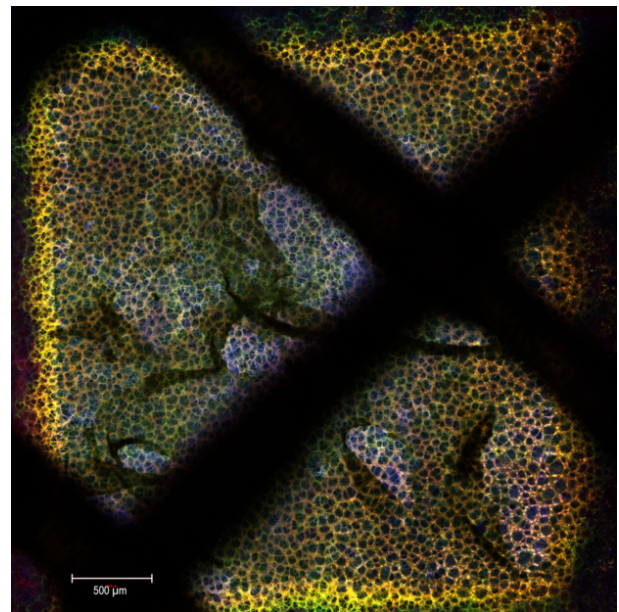


Figure 7: CLSM image showing internal and external fouling in MF. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI).



(a) Before backwash



(b) After backwash

Figure 8: Confocal images of 2-day fouled MF membrane before and after a backwash. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI).

356 *3.3. Ultrafiltration without backwash*

357 To visualize baseline biofouling behavior in UF, wastewater effluent was stained with 2.4  
358  $\mu\text{g}/\text{mL}$  WGA, 2.4  $\mu\text{M}$  DAPI, and 4.7  $\mu\text{g}/\text{mL}$  PI, and was filtered through a UF membrane  
359 at a flux of 48 LMH with a cross-flow velocity of 20 cm/s. During one hour of filtration,  
360 pressure increased from 1.1 to 2.1 bar, indicating significant fouling. CLSM images of the  
361 membrane were collected in situ during filtration.

362 Figure 9 shows a tiled scan of the entire active area using a 10x objective. The fouled  
363 membrane exhibited curved stripes of both cells and EPS as well as patches of black dots,  
364 likely representing holes or channels through the biofilm. Figure 10 shows 3-D cross-sections  
365 of a primarily striped region of the biofilm. Stripes and holes appear in a regular pattern  
366 between consecutive cells of the spacer, with stripes appearing approximately tangential  
367 to the expected direction of flow. The holes congregate near spacer filaments and on the  
368 concave sides of the striped arcs.

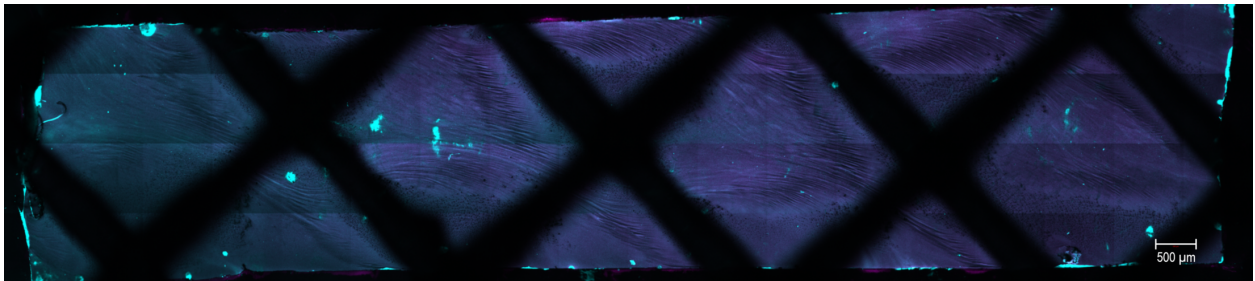


Figure 9: UF membrane fouled for approximately one hour during cross-flow from right to left. Cyan = DAPI; magenta = EPS (WGA); the combination appears purple. Readers may zoom in to see details of fouling pattern.

369 Other in situ membrane biofouling visualization studies with comparable feed spacers  
370 have revealed somewhat similar macroscopic fouling patterns despite differences in imaging  
371 methods, hydrodynamics, and membranes. In an in situ optical coherence tomographic  
372 study of biofouling on unpressurized NF membranes with the same type of spacer, Fortunato  
373 et al. [12] saw membrane fouling concentrated upstream of each spacer filament that was

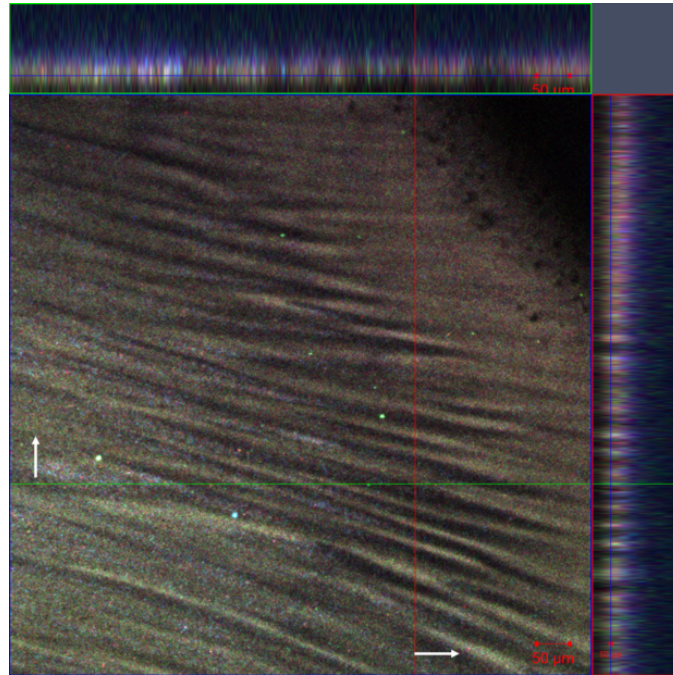


Figure 10: Depth section of stripe and spot patterns from the fouled membrane of Fig. 9. Cyan = cells (DAPI); green = EPS (WGA); red = dead cells (PI). Depth section planes are indicated with arrows.

374 farther from the membrane. Perhaps due to the significant flux through the UF membrane  
 375 in the present study and the association between higher permeate flux and increased spatial  
 376 uniformity [30], the biofilm is distributed more uniformly over the membrane; however,  
 377 there is heterogeneity in patterning within each cell of the spacer. The region upstream of  
 378 each elevated spacer filament that showed the most membrane fouling in Fortunato et al.  
 379 is the region of the UF membrane in Figs. 9 and 10 that has the least patterning (neither  
 380 pronounced stripes nor pits). Similarly, the downstream corner of each spacer cell (which  
 381 is the left corner of each cell in Fig. 9) had the highest bacterial load in forward osmosis  
 382 biofouling in a study by Kastl et al. [30]. In contrast, however, Huang et al. [13] found the  
 383 fastest accumulation of bacteria on an RO membrane to be in the upstream corner of each  
 384 feed spacer cell, although the spacer did not fill the entire feed channel in that study. Stripes  
 385 and holes were not visible in any past in situ visualization studies of filtration membrane  
 386 biofouling known to the authors. Given that the pattern of stripes and spots observed in Fig.

387 9 is repeated in the spaces between each set of spacer filaments, the lines and spots likely  
388 represent self-organization of the microbes and other constituents that make up the biofilm,  
389 rather than a pattern related to membrane structure or surface heterogeneity. A similar  
390 pattern of stripes was recorded after 1 d of MF fouling (Fig. 3a). Patterned morphologies  
391 such as these may help the biofilm survive in local hydrodynamic conditions, as discussed  
392 below.

393 Patterns of holes in biofilms, as seen in Figs. 9 and 10, have been noted when biofilms  
394 disperse. McDougald et al. [40] review mechanisms of biofilm dispersal, which often involves  
395 EPS degradation, cell lysis, and dispersal of a select few bacteria, leaving a hollow micro-  
396 colony. Increase or rapid decrease in nutrient concentration or depletion of oxygen can lead  
397 to dispersal. The biofilm forming *P. aeruginosa* is known to form hollow microcolonies prior  
398 to dispersal of cells [41]. Similar round channels through the biofilm have been observed  
399 with CLSM on *P. aeruginosa*-fouled RO membranes after cleaning by osmotic backwashing  
400 [42].

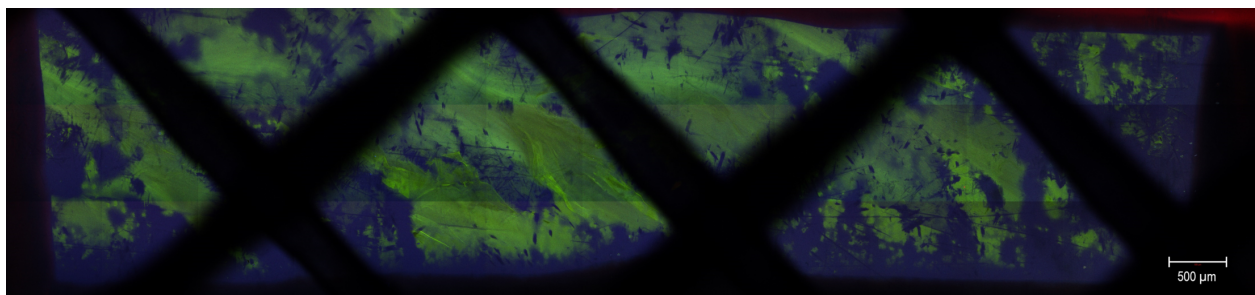
401 Various rough patterns occur in biofilms when nutrient scarcity leads to unstable growth.  
402 Stewart [43] reviewed effects of convection on biofilm formation, emphasizing how increased  
403 mass transfer at peaks in a rough biofilm structure leads to increased local growth rates,  
404 creating an instability resulting in increasingly rough biofilms. For example, *P. aerugi-*  
405 *nosa* biofilms grown under oxygen-depleted conditions [44] developed stripes. The stripes  
406 and possibly also the holes observed in the present study's biofilms may results from the  
407 improvement in nutrient capture enabled by biofilm roughness under convection conditions.

#### 408 3.4. Ultrafiltration with backwash

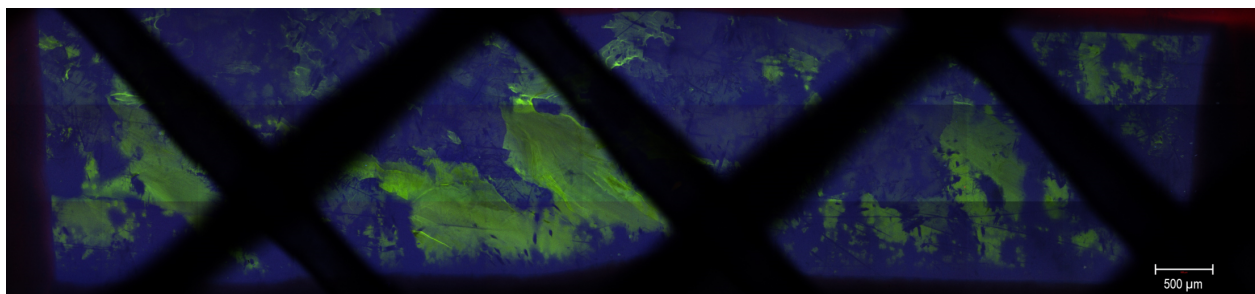
409 The in situ CLSM images in this section show how fouling progressed on the UF mem-  
410 brane over four days of operation with periodic backwashing. Before imaging, dyes were  
411 added to each a final dilution of 0.6  $\mu\text{g}/\text{mL}$  of WGA and PI and 0.6  $\mu\text{M}$  DAPI. After the  
412 first day of fouling at 54 LMH, limited fouling was observed, and flux was increased to 80



413 LMH. Over the 4 d fouling period, operating pressure increased by a factor of two, indicating  
414 significant fouling. Foulant accumulation was heterogeneous on a range of scales. By the  
415 third day, Fig. 11 shows that an overall pattern develops: large flakes of EPS-rich material  
416 stick to the membrane, while some areas are nearly clean, particularly after backwashing.  
417 Some of the EPS-rich regions appear to be composed of multiple layers of folded or wrinkled  
418 flakes. A very large flake, which could be seen flapping in the flow through the microscope  
419 eyepiece, appears as a dark, diagonal blur in the left-hand side of Figs. 12a and 12b. Some  
420 small regions of the membrane appear to perpetually remain clean.



(a) Before backwash

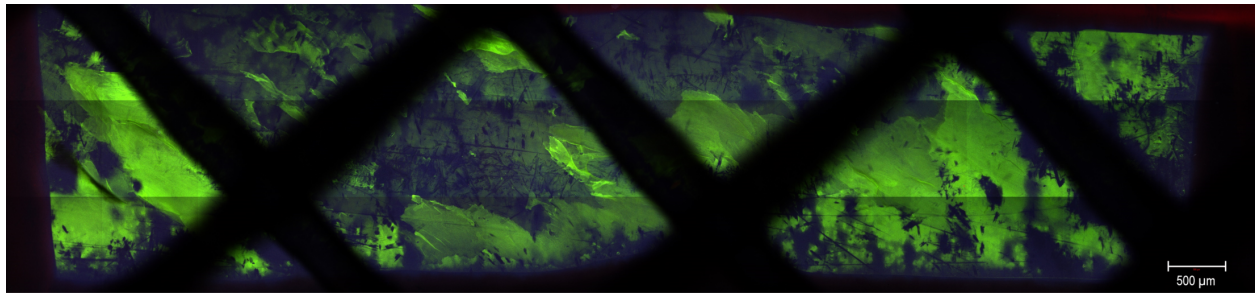


(b) After backwash

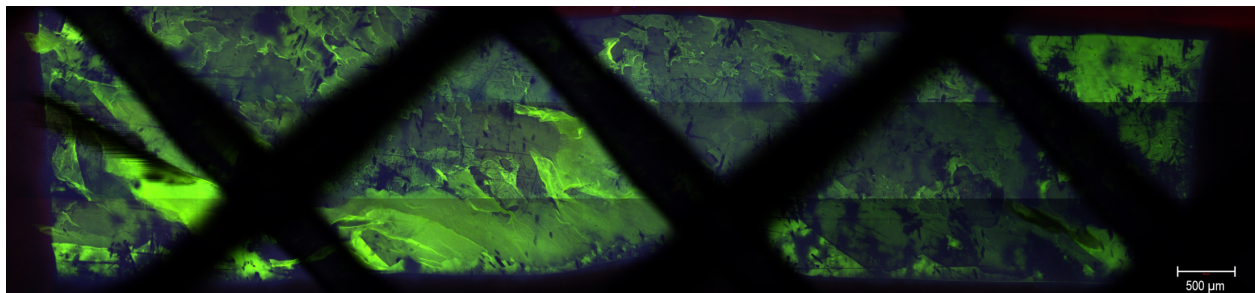
Figure 11: Tiled confocal images of 2 day-old UF biofilm captured in situ before and after a backwash. Flow was right to left. Blue = DAPI; green = EPS (WGA); red = dead cells (PI)..

421 The accumulation process between backwash cycles is illustrated in Fig. 13, showing three  
422 accumulation behaviors: (1) small regions (order 10 μm in width) remain dark throughout  
423 the permeate production step, indicating low local membrane permeability; (2) most of  
424 the membrane is relatively clean of EPS right after a backwash, but gradually accumulates





(a) 3 d



(b) 4 d

Figure 12: Confocal images of UF biofilm captured in situ after (a) 3 d and (b) 4 d. Flow was right to left. Blue = DAPI; green = EPS (WGA); red = dead cells (PI).

425 foulant (possibly including fluorescent but unbound WGA lectins) over the course of perme-  
 426 ate production; and (3) regions of membrane that were not cleaned by the backwash step  
 427 are already so thickly fouled that local permeation is lower and further foulant accumulation  
 428 during permeation is limited. In regions exhibiting the second behavior (periodic fouling  
 429 and cleaning), a lacy pattern of dark spots in the WGA-rich material develops beginning on  
 430 the third day (see Figs. 14c,e and the upper-right corner of Figs. 14a,c) that is most evident  
 431 later in each permeate production cycle (see Fig. 13). These spots (likely holes in the biofilm,  
 432 as discussed in Section 3.3) are similar in size to those seen on the UF membrane without  
 433 backwash (Figs. 9 and 10). Stripes and wrinkles in the direction of flow, as discussed in  
 434 Section 3.3, persist, but appear more subtle with periodic backwash than without.

435 The progression of fouling on the UF membrane is substantially shaped by the effects of  
 436 backwashing. Comparison of Figs. 11a and 11b shows that backwashing removes biofouling

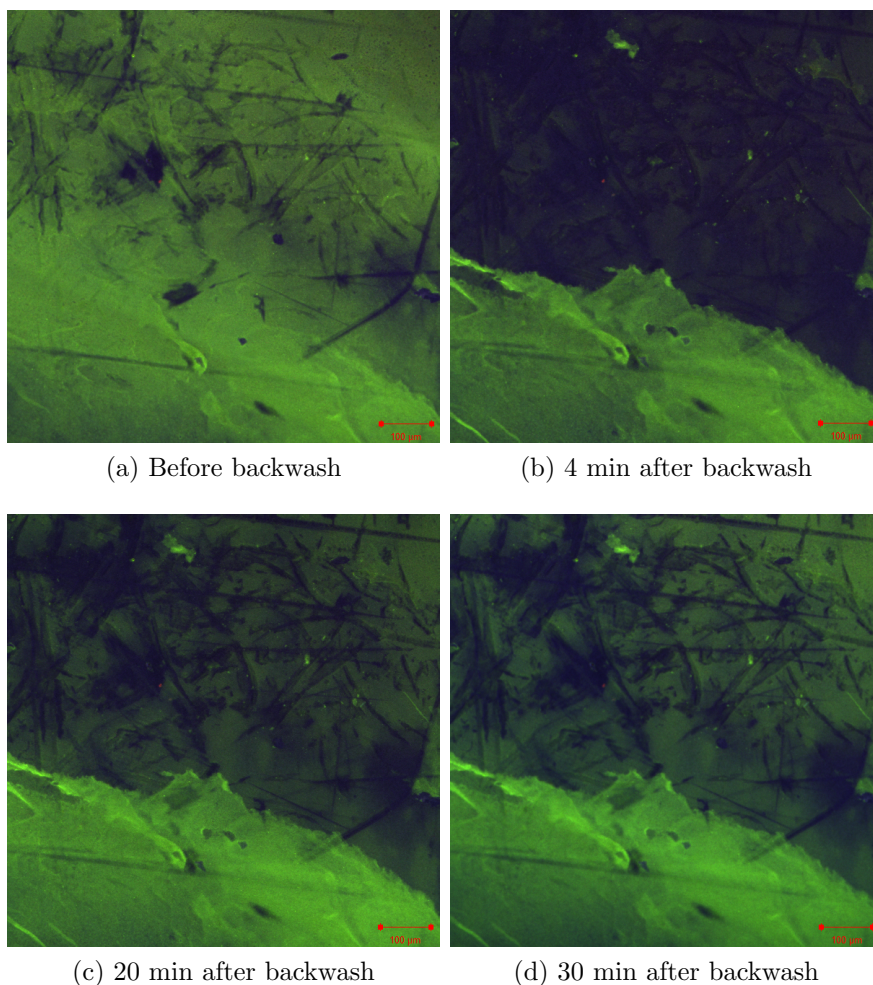


Figure 13: Confocal images of the progression of UF membrane fouling after a backwash. All images are collected after 3 d of fouling and the time since the last backwash is specified below each image. Region A is also depicted in Fig. 14 and Region B is also depicted in Fig. 15. Blue = DAPI; green = EPS (WGA); red = dead cells (PI).

437 in large ( $>100\ \mu\text{m}$ ) flakes with well-defined edges and leaves large areas of membrane nearly  
 438 clean, but has little effect on other membrane regions. The effect of backwashing is captured  
 439 in more detail in Figs. 14 and 15, which highlight smaller regions of the membrane at several  
 440 different times. While large regions of the membrane are left almost clean (see, e.g., Fig.  
 441 15), other areas in each image show no change and seem to be unaffected by backwashing.  
 442 Some sections of foulant move in the direction of flow but stay attached to the membrane  
 443 (compare Figs. 15c and d). Some regions even accumulate more foulant as sheets that peeled

444 off elsewhere pile up on the membrane (see, e.g., Fig. 14d). The thicker regions of layered or  
445 wrinkled foulant are evident in a 3-D rendering of the biofilm (Fig. 15c). Repeated disruption  
446 by backwashing also makes the biofilm accumulate unevenly over time, as demonstrated by  
447 the nonlinear progression of average biofilm thickness (see captions of Fig. 14) over time  
448 (computed as described in Sec. 2.6 from 3-D images taken prior backwashing at the same  
449 location shown).

450 In a separate experiment to observe the backwashing process itself, a UF membrane  
451 was fouled with periodic backwashes for 2 d. A cohesive film formed, as shown in Fig.  
452 16a. During the backwash process, images showed the film detaching from the membrane in  
453 several places and ballooning into a bubble wrap-like structure (Fig. 16b); in the imaging  
454 plane of the 3-D structure, the cross-section of the biofilm appears as a set of closed forms.  
455 (For reference, both 2-D and 3-D images of a similar structure are shown in Fig. 17.) As  
456 seen in Fig. 16c, this backwash was ultimately not successful at removing the film, and the  
457 film appears largely unchanged after backwashing but for a slight change in the shape of  
458 large the wrinkle near the middle of the image.

459 The ineffectiveness of the backwash shown in Fig. 16 and the removal of foulant in  
460 large flakes in the more successful backwashes shown earlier (e.g., Fig. 14) demonstrates  
461 the cohesiveness of the biofilms formed on the UF membrane and provide some insight into  
462 biofouling mitigation strategies. Removal of organic foulant from RO and FO membranes in  
463 large flakes and sheets was previously demonstrated [9, 8]. Furthermore, the dark lines on  
464 the membrane that never seem to accumulate foulant often coincide with the edges of patches  
465 where flakes are removed during backwashing. A similar pattern is also seen in SEM image  
466 of the unfouled UF membrane (Fig. C.26), suggesting the membrane texture defines the  
467 placement of these low-fouling areas. We hypothesize that the edges of the thinner regions  
468 of biofilm that form over less-permeable membrane regions are more likely to tear and allow  
469 for the removal of flakes. Intentionally patterning membranes with low-permeability regions

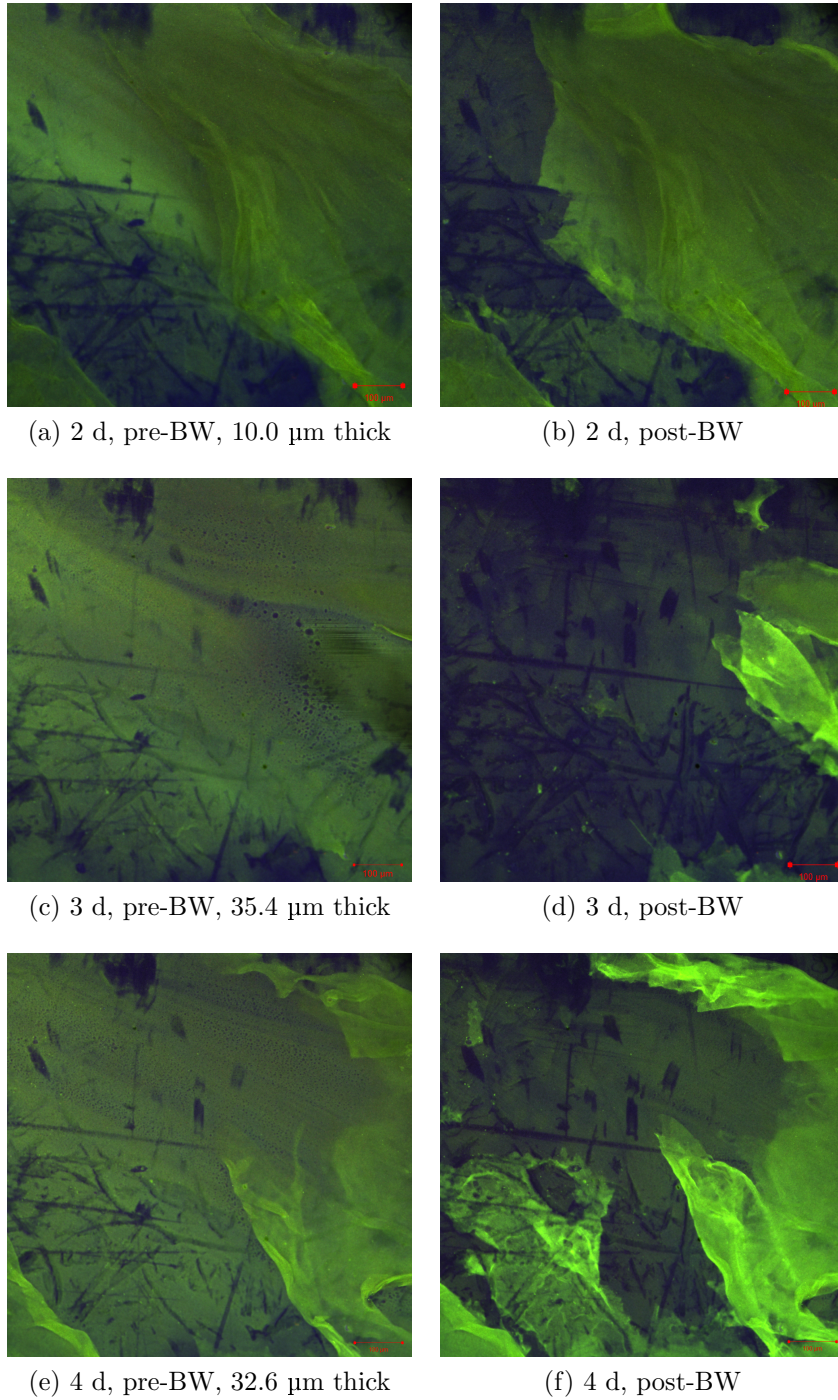


Figure 14: Confocal images of a single region of fouled UF membrane captured in situ with a 10x objective before and after a backwash (BW). Blue = DAPI; green = EPS (WGA); red = dead cells (PI). Average biofilm thicknesses before backwash, calculated from 3-D images, are included in sub-figure captions to show nonlinear biofilm accumulation.



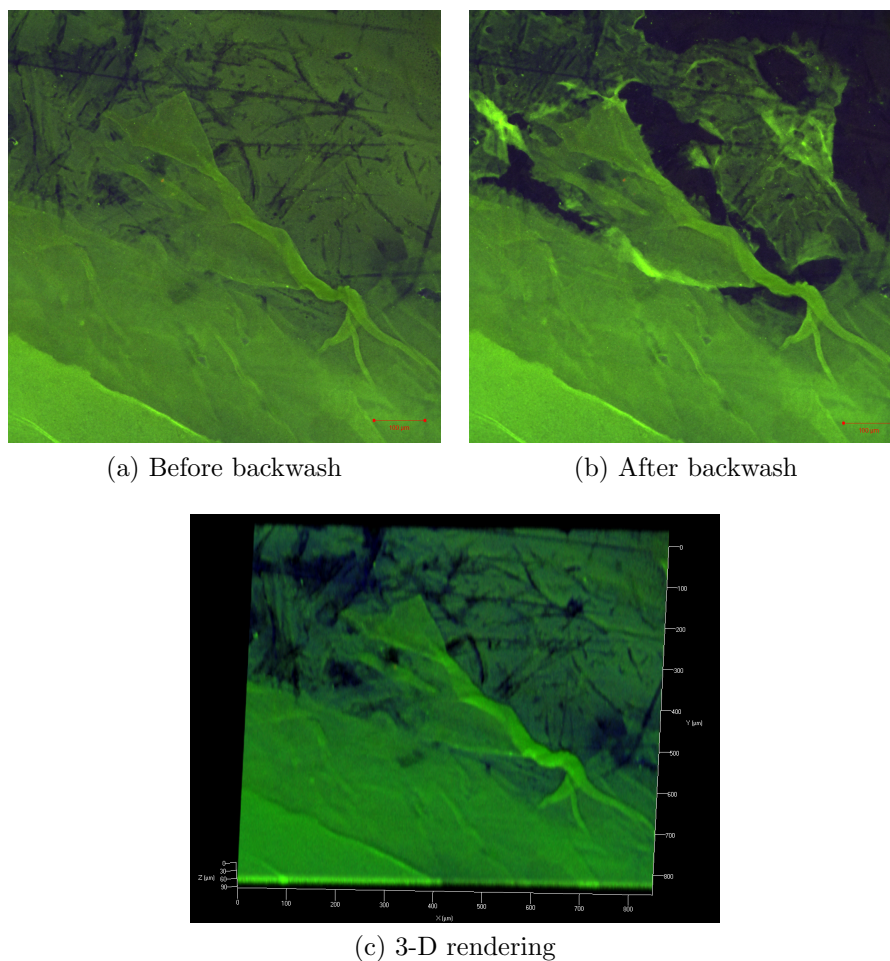


Figure 15: Confocal images of a single region of fouled UF membrane after 4 d, captured in situ with a 10x objective. Blue = cells (DAPI); green = EPS (WGA); red = dead cells (PI).

470 could be a method of raising the effectiveness of backwashing with the potential outcome  
 471 of raising the flux at which the membrane can be operated long-term or extending the time  
 472 between chemical cleans.

### 473 3.5. Discussion

#### 474 3.5.1. In situ vs. ex situ imaging

475 This study was carried out to better understand fouling progression as well as to separate  
 476 realistic membrane biofouling behaviors from artefacts of ex situ imaging. As shown in Fig.  
 477 17, sample removal and simple preparation for imaging (placing on a slide in a drop of feed

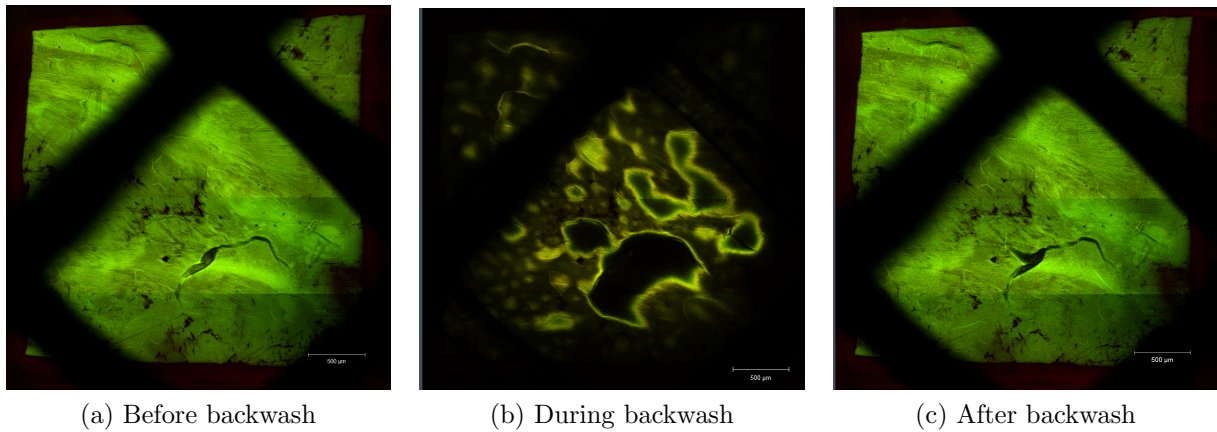


Figure 16: Confocal images of a single region of 2-day fouled UF membrane captured in situ with a 10x objective (a) before, (b) during, and (c) after an ineffective backwash. Green = EPS (WGA); red = dead cells (PI). Scale bars (lower right of each image) span 500  $\mu\text{m}$ .

478 liquid under a coverslip) caused a rapid ( $<30$  min) change in foulant conformation from  
 479 flat to buckled; in this case, a 2-D slice and 3-D rendering of the biofilm imaged ex situ  
 480 show delamination of the biofilm from the MF membrane in a pattern reminiscent of bubble  
 481 wrap. The in situ and ex situ images in Fig. 17 do not correspond to the same location on  
 482 the membrane, but the in situ image of the entire membrane (Fig. 3) shows that bubble  
 483 wrap-like delamination did not occur anywhere on the membrane during filtration.

484 Similar buckled conformations have been observed in ex situ CLSM images of membrane-  
 485 bound biofilms previously [17, 31]. In the present study, cleaning processes involving relax-  
 486 ation (see Appendix B.1) and salinity change (see Appendix B.2) created similar confor-  
 487 mations in the biofilm in situ. Bar-Zeev et al. [31] noticed a similar buckled conformation  
 488 in ex situ imaging of biofilms, including changes in biofilm conformation when a glass cov-  
 489 erslip was placed on a hydrated biofilm for ex situ imaging. They attributed the flatter  
 490 conformation under the coverslip to a change (flattening) induced by the coverslip; however,  
 491 without in situ imaging, the original biofilm conformation before removal from the operating  
 492 conditions of the flow cell cannot be known.

493 Biofilm delamination from the membrane during preparation for ex situ imaging may

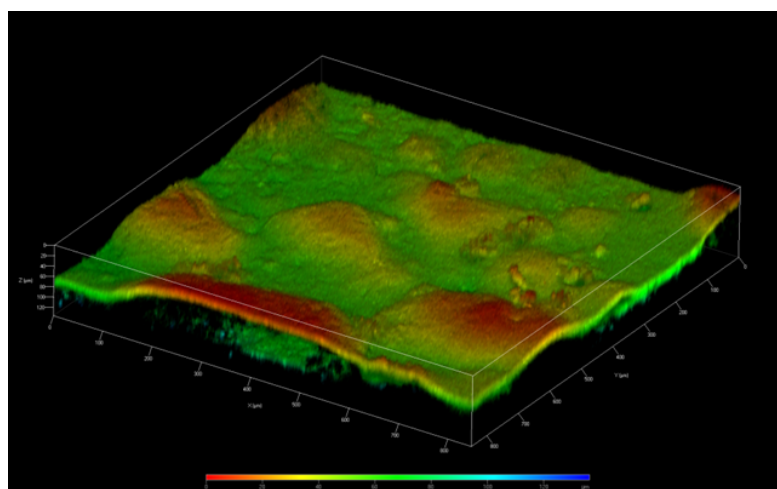
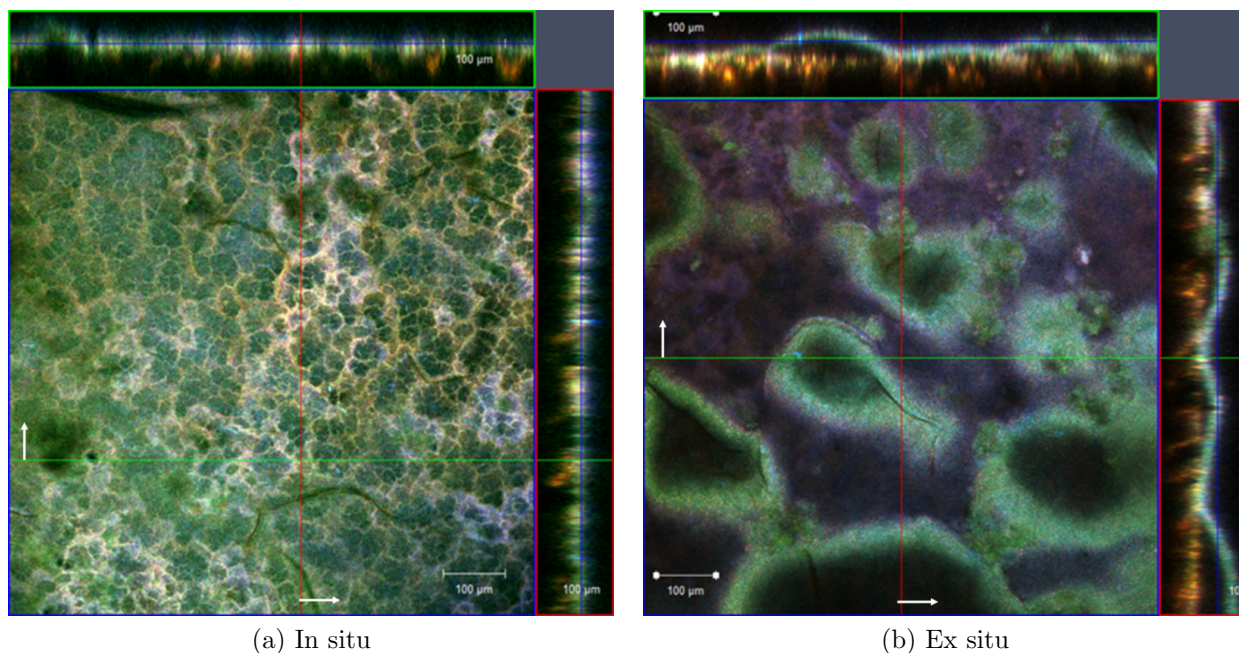


Figure 17: In situ and ex situ images of a 4 day-old MF biofilm. (a) shows the biofilm is initially flat, but image (b) (and its rendering in (c)) of a region of the biofilm shortly after removal from the flow cell demonstrates buckling into a characteristic bubble wrap-like conformation. Cyan = cells (Hoechst); green = EPS (WGA); red = dead cells (PI). Depth section planes are indicated with arrows.

494 occur due to changes in nutrient availability when cross-flow and transmembrane flux no  
 495 longer bring sufficient nutrients to the biofilm. Hunt et al. [45] observed that *P. aeruginosa*  
 496 biofilms detached from a stainless steel substrate both under stagnation conditions (when

497 flow was stopped) or when the flowing solution was replaced with a nutrient-scarce solution.  
498 A biofilm grown on a membrane under particular conditions may begin to detach when  
499 conditions change in preparation for ex situ imaging. As shown in Fig. 17, the remarkable  
500 change in the membrane-grown biofilm upon removal from filtration conditions demonstrates  
501 the limitations of ex situ imaging and the importance of not only in situ but *operando*  
502 imaging for understanding the conformations and behaviors of membrane biofouling.

### 503 3.5.2. Fouling behaviors and mechanisms

504 Several accumulation and removal behaviors were revealed through in situ imaging of  
505 wastewater effluent filtration with periodic backwashing. UF membrane images in Fig.  
506 18 highlight regions where foulant advection, sloughing, and accumulation are significant.  
507 In parts of the membrane that were successfully cleaned by a recent backwash, high flux  
508 through the clean membrane brings new biofilm components to the surface. In parts of  
509 the membrane where local permeation is low due to membrane damage or prior biofouling,  
510 additional foulant accumulation occurs slowly or not at all. During a backwash, part of  
511 the membrane is cleaned, while other sections remain covered in biofilm; some regions even  
512 accumulate additional fouling during the backwashing process if the delaminated biofilm  
513 folds or re-deposits elsewhere. MF displayed similar accumulation and removal behaviors  
514 and additionally exhibited internal fouling with significant EPS accumulation inside large  
515 internal pores, as shown in Fig. 6.

516 Biofouling behaviors on membranes with permeation, cross-flow, and periodic backwash-  
517 ing are shaped by a variety of mechanisms. Observed behaviors are qualitatively arrayed,  
518 using micrographs from this report, from biologically- to physically-mediated in Fig. 19 to  
519 illustrate the range of filtration membrane biofouling behaviors observed in this study. Bet-  
520 ter understanding of membrane biofouling through in situ visualization enables insight into  
521 potential improvements to membranes and processes in water reuse applications, such as  
522 patterning membranes to encourage biofilm flake sloughing, as discussed in Sec. 3.4.



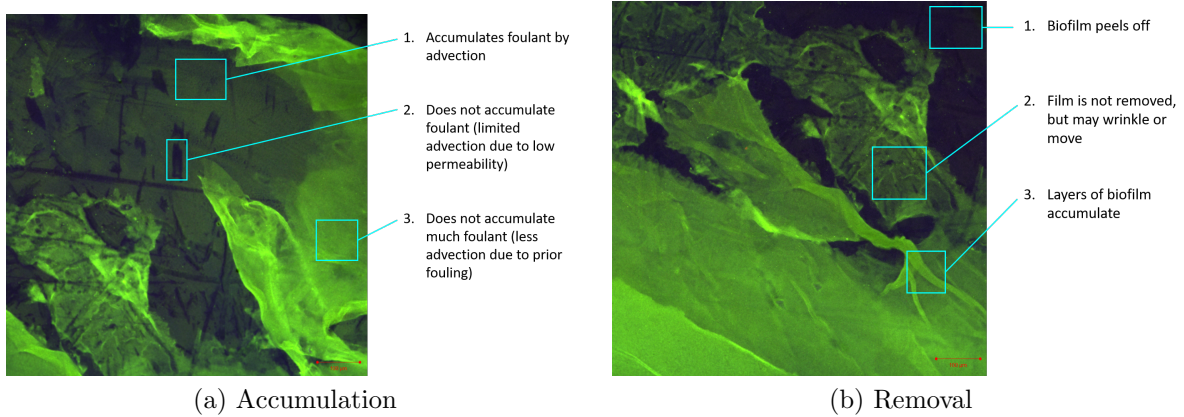


Figure 18: Summary of accumulation and removal behaviors in ultrafiltration with periodic backwash

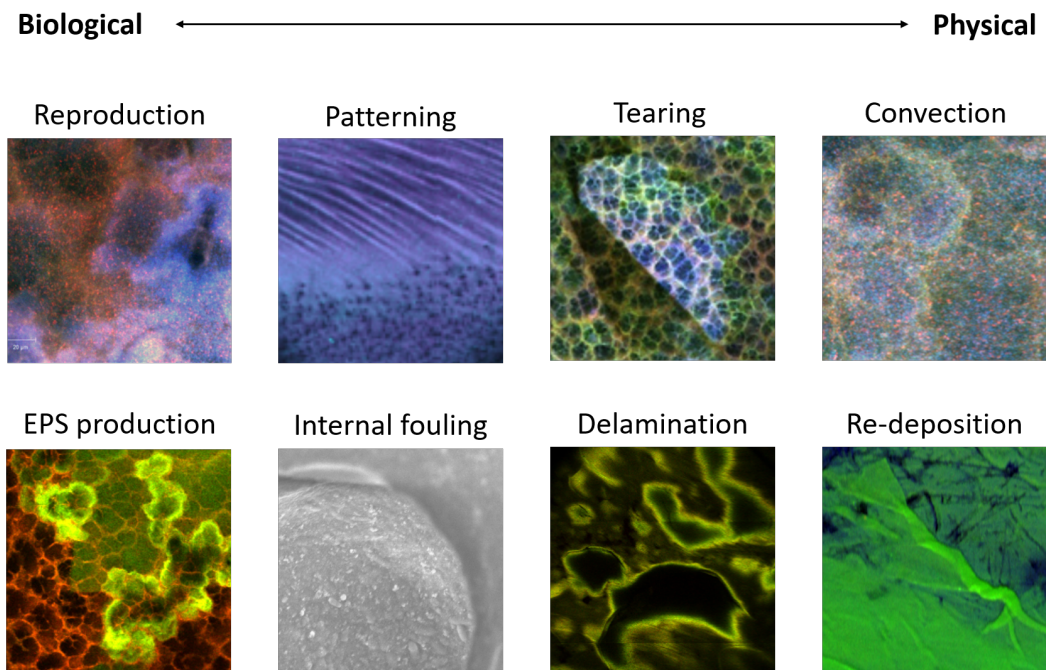


Figure 19: Graphical summary of biofouling behaviors observed throughout this study

523 *3.5.3. Limitations*

524 We acknowledge several limitations of this study's method of visualizing biofouling mech-  
 525 anisms in water reuse. The longest experiments, at 4 d in duration, were still significantly  
 526 shorter than the multi-year lifespan of an MF or UF membrane, so long-term fouling be-

527 haviors were not captured by this study. The experimental apparatus also used a relatively  
528 low flux for backwashing and did not incorporate air scouring, which is an aspect of back-  
529 wash procedures in reuse facilities [33]. The persistent use of stains, especially live cell  
530 stains (see Appendix A), had an unknown effect on the behavior of the biofilm-forming  
531 microbes; similarly, the unrealistically long feedwater storage times prior to filtration (days  
532 to weeks) had an unknown effect on fouling behavior. While the small imaging area was  
533 large enough to observe heterogeneity in fouling behavior on the scale of a feed spacer cell,  
534 it was far too small to capture variations that might exist between distant parts of a com-  
535 mercial membrane element. Finally, the thin glass slide used to seal the flow cell and allow  
536 imaging was both too thin to allow application of the high pressures used in RO, another  
537 biofouling-prone membrane process common in potable water reuse, and too thick to allow  
538 for CLSM imaging at the resolution required to visualize the biofilm conformation at the  
539 level of individual cells due to the short working distances associated with high-resolution  
540 objectives. While these limitations prevented visualization of the exact behavior of real  
541 wastewater reuse biofilms, the use of realistic hydrodynamic conditions and real wastewater  
542 effluent provide some insight into biofouling behaviors that can only be observed in situ.

543 Future work might mitigate several of the aforementioned limitations by improving the  
544 flow cell design. Using hollow fiber membranes could allow for both plane views and cross-  
545 sectional views of the biofilm, despite the poor vertical resolution of the long working distance  
546 objectives; hollow fiber membranes are also more realistic in MF and UF for water reuse.  
547 Designing a flow cell into which a water immersion objective could be integrated would  
548 also improve resolution and enable quantitative analysis of biofilm characteristics such as  
549 porosity. Using new ultra-permeable RO membranes could enable in situ visualization of  
550 RO membrane fouling with UF/MF permeate at relatively low pressures, although dye  
551 rejection by the membranes would be a concern that would need to be mitigated, perhaps  
552 by replacement of the wastewater effluent shortly after introducing stains.

#### 553 4. Conclusion

554 In situ confocal laser scanning microscopy of membrane biofilms under filtration con-  
555 ditions was used to elucidate biofouling behaviors in MF and UF. Biofilms formed within  
556 minutes due to convection of water constituents to the membranes at high flux, causing ex-  
557 ternal fouling of both MF and UF membranes. Additionally, in situ CLSM and ex situ SEM  
558 images showed that MF membranes exhibited internal fouling, with apparent cell prolifera-  
559 tion and EPS production within networks of large pores beneath the membrane surface.

560 Biofouling behavior in conditions with periodic backwashing was influenced by the co-  
561 hesiveness that is a hallmark of biofilms. Behaviors including wrinkling, delamination, tear-  
562 ing, and re-deposition after backwashes were influenced by the mechanics of the biofilm,  
563 including its cohesion and its adhesion to the membrane. Buckling of the biofilm in a char-  
564 acteristic bubble wrap-like conformation was observed during backwashing, relaxation, and  
565 high-salinity soaking as well as after removal from the flow cell for ex situ imaging. Pat-  
566 terning of the biofilm at the scale of 10–50 microns was also observed, rendering visible the  
567 complex interplay of microbiology, local hydrodynamics, mechanics, and mass transfer that  
568 dictates biofouling behavior.

569 Observed changes in biofilm conformation between in situ and ex situ CLSM images high-  
570 light the importance of in situ and *operando* imaging for understanding biofouling behavior,  
571 particularly when biofilm conformation is of interest. Biofilm delamination was observed  
572 even during in situ imaging under relaxation conditions (i.e., no flux or cross-flow), indicat-  
573 ing the responsiveness of the biofilm to subtle changes in its environment. While promising  
574 in situ imaging methods continue to be developed, we invite fouling researchers to recognize  
575 the possibility of rapid conformational change when biofilms are removed from filtration  
576 conditions for ex situ imaging.

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## 587 **CRedit authorship statement**

588 EWT: Conceptualization, methodology, investigation, data curation, formal analysis,  
589 resources, writing—original draft, writing—review and editing, visualization, funding ac-  
590 quisition. BR: Methodology, formal analysis, resources, writing—review and editing. RK:  
591 supervision, writing—review and editing, funding acquisition.

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## 700 **Appendix A. Effect of live cell stain**

701 Due to the potential cytotoxicity and mutagenicity of Hoechst 33342 [39], an analogous  
702 experiment to that reported in Section 3.2 was conducted in which Hoechst was only added  
703 prior to the last imaging session to qualitatively determine the effect of frequent Hoechst  
704 application on the fouling behavior in the experiments reported on throughout this study.  
705 2.9  $\mu\text{g}/\text{mL}$  of WGA and 1.2  $\mu\text{g}/\text{mL}$  of PI were added daily before imaging, and 1.8  $\mu\text{g}/\text{mL}$   
706 Hoechst was added just before the final imaging session after 4 d of fouling. MF membrane  
707 images with and without the daily application of Hoechst are compared in Fig. A.20. Both  
708 with and without Hoechst, after 4 d of fouling, a flat layer of live and dead cells and EPS  
709 forms on the active layer and internal fouling with live cells and EPS is apparent within  
710 large internal pores. The addition of Hoescht seems to dull the appearance of EPS whether  
711 it is added daily or just before the final imaging session. Without Hoechst, it is particularly  
712 apparent how concentrated EPS are at the edges of internal pores in which fouling occurs.  
713 The similarity of the biofilm at 4 d with and without daily application of Hoechst suggests  
714 that the daily use of Hoechst as reported on in Sec. 3 did not significantly affect qualitative  
715 fouling behavior.

716 The membrane of Fig. A.20 that was only treated with Hoechst once at 4 d was depth-  
717 scanned in Fig. A.21, confirming that what appears to be internal fouling does occur below  
718 the membrane active layer.



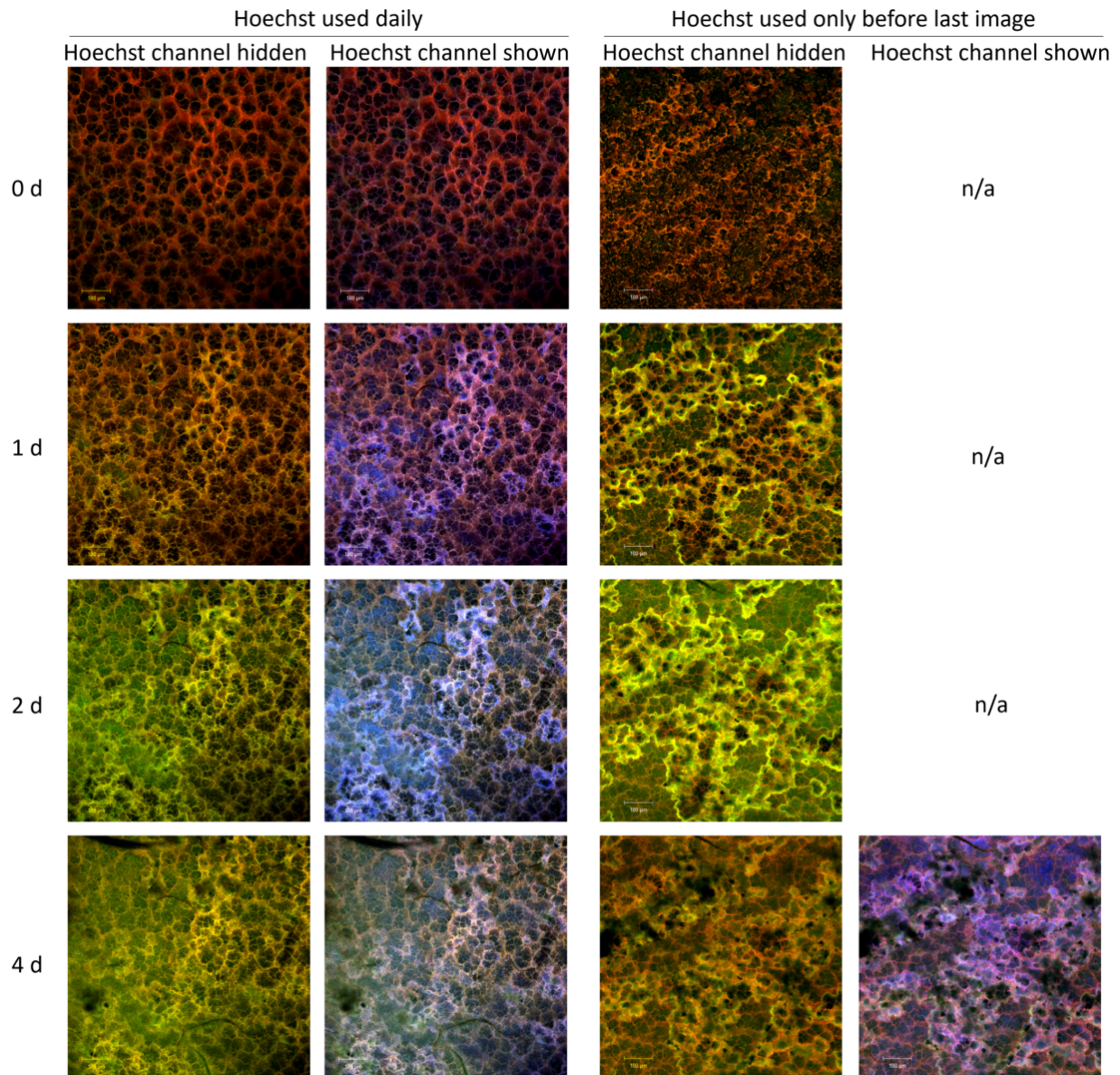


Figure A.20: Confocal images of fouled MF membrane with application of Hoechst daily and only just before the last imaging session, shown with and without the Hoechst channel illuminated. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI).

## 719 Appendix B. Additional cleaning procedures

720 Due to the cohesive external fouling observed on UF membranes, two cleaning pro-  
 721 cedures beyond backwashing were investigated. Anecdotal reports of relaxation-induced  
 722 biofilm sloughing inspired the use of relaxation as a potential cleaning strategy, and past

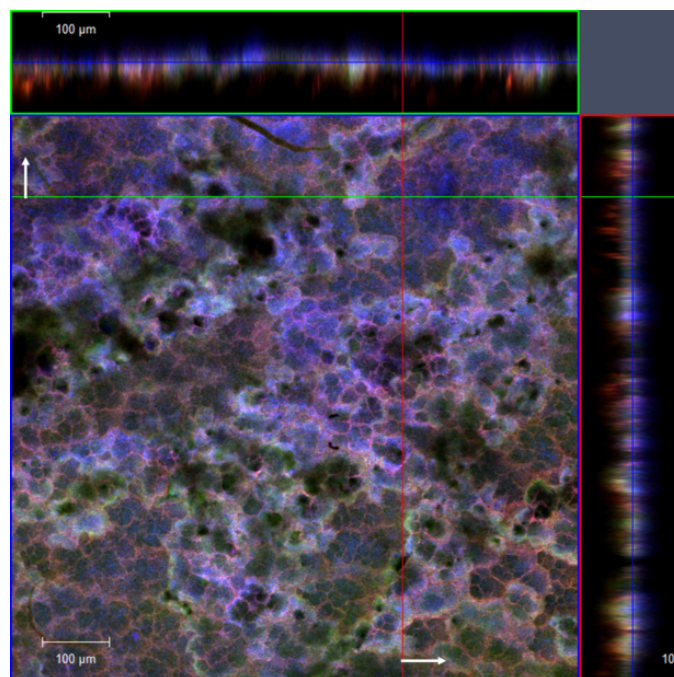


Figure A.21: 3-D scan of biofilm on MF membrane after 4 d with Hoechst added only before this final imaging session. Blue = cells (Hoechst); green = EPS (WGA); red = dead cells (PI). Depth section planes are indicated with arrows.

723 observation of effective organic fouling removal by salt cleaning [46] inspired the used of a  
 724 saline soak. Neither strategy was effective as removing the biofilm, but both revealed biofilm  
 725 delamination behavior, as discussed further below. Visualization of membrane cleaning with  
 726 bleach was also attempted, but bleaching of the fluorescent probes occurred and the resulting  
 727 images were not interpretable.

### 728 *Appendix B.1. Relaxation*

729 The biofilm shown in Fig. 9 was allowed to relax in order to observe biofilm behavior  
 730 during stoppage of flux and cross-flow. After approximately 3 h of fouling at 20 cm/s cross-  
 731 flow and approximately 48 LMH flux, both circulation and syringe pumps were stopped.  
 732 Flux declined as pressure wound down to atmospheric. After about five hours with both  
 733 pumps off, the biofilm was imaged again (Fig. B.22), revealing slight delamination of the  
 734 biofilm from the membrane in a pattern of mounds, approximately 100  $\mu\text{m}$  wide and 20  $\mu\text{m}$



735 tall, and oriented similarly to the stripe pattern shown in Figs. 9 and 10. These wrinkles  
736 show some similarity to the bubble wrap pattern seen in a biofilm imaged ex situ (Fig.  
737 17b,c). Relaxation, as performed here, was not effective in cleaning the membrane, but did  
738 cause delamination, which points to potential for biofilm removal.

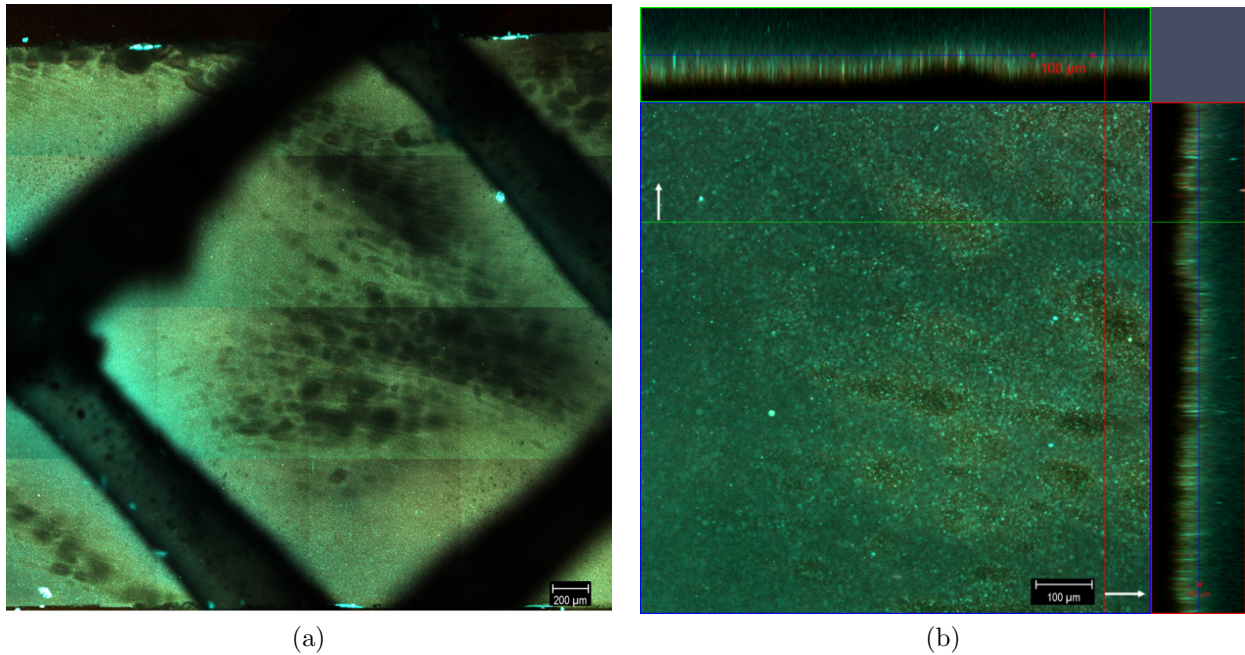


Figure B.22: 3-D wrinkling of UF biofilm after relaxation: (a) plane view and (b) close-up with cross-sectional views showing delamination. Cyan = cells (DAPI); green = EPS (WGA); red = dead cells (PI). Depth section planes are indicated with arrows.

### 739 *Appendix B.2. Saline soak*

740 A saline soak was used in an attempt to clean a UF membrane that had been fouled for  
741 2 d with periodic backwashing. Shortly after a backwash that cleared part of the membrane,  
742 a 10% NaCl solution was introduced to the flow loop and the system was left to sit without  
743 cross-flow or applied pressure for approximately 30 min, during which time the membrane  
744 was imaged (Fig. B.23c), revealing delamination of sections of the remaining biofilm from  
745 the membrane. When cross-flow was restarted, the biofilm flattened again and was not  
746 removed. Although ineffective as a cleaning procedure, the finding of delamination during

747 a short saline soak (similar to behavior during a backwash, Fig. 16b, or a longer relaxation,  
748 Fig. B.22) is noteworthy.



Figure B.23: Backwashing and a saline soak were performed sequentially on a UF membrane fouled for 2 d. (a) and (b) show the effect of a backwash, while (c) and (d) show the effect of a saline soak.

## 749 **Appendix C. SEM of clean and fouled membranes**

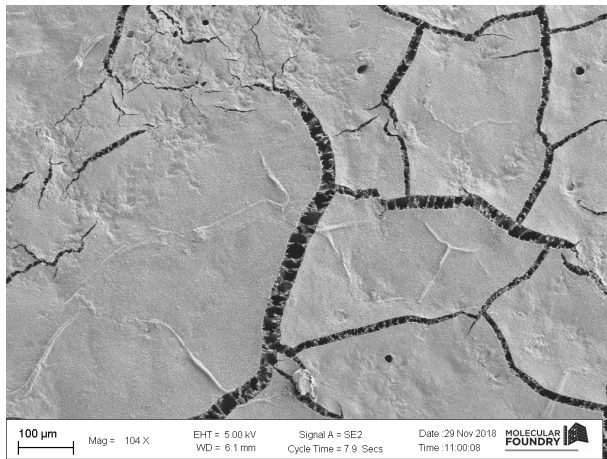
750 Fouled MF membranes were examined with SEM (Ultra 55 FESEM, Carl Zeiss) to  
751 confirm suspected internal fouling.

752 To preserve the biofilm conformation, the fouled membrane was prepared for SEM as  
753 follows: Glutaraldehyde was added to the flow loop at a final concentration of 2.5% and left  
754 to fix cells for 30 min. The fouled membrane was removed from the cell and dipped for 10-15  
755 min each in 10%, 25%, 75%, and 90% ethanol in deionized water and then twice in 100%  
756 ethanol to dehydrate the sample. The sample was finally vacuum dried and sputtered with  
757 gold using a SC7620 Mini Sputter Coater/Glow Discharge System (Electron Microscopy  
758 Sciences).

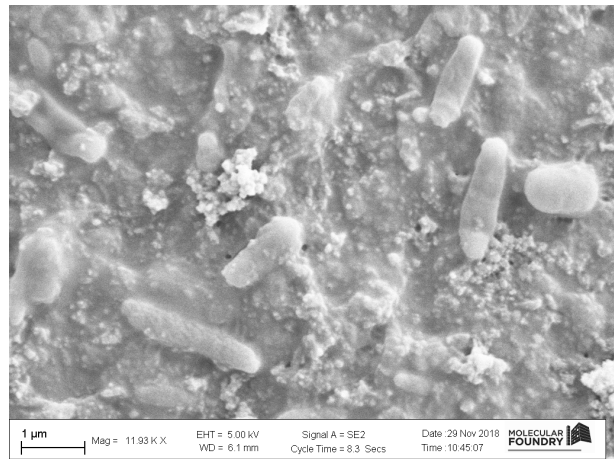
759 In SEM images of a fouled MF membrane (Fig. C.24), rod-shaped bacteria (bacilli) of  
760 approximately 2  $\mu\text{m}$  length are seen lining a large internal pore that is visible through a hole  
761 in the surface; CLSM images in Section 3.2 suggest that such colonization of internal pores  
762 occurred throughout the membrane. Cracks in the fouled MF membrane are also visible,  
763 likely as a result of the drying process.

764 Unfouled MF and UF membranes were gold-sputtered and imaged with SEM to reveal  
765 membrane texture and aid interpretation of CLSM images. The MF membrane (Fig. C.25)  
766 displays large pores beneath the active layer, some of which connect to the active layer  
767 through holes that are large enough to allow bacteria to pass through. The UF membrane  
768 (Fig. C.26) has a smooth surface without visible pores at 168x magnification, but displays  
769 patterns that may have occurred in storage or handling. These membrane surface patterns  
770 appeared to affect fouling patterns, as shown in Section 3.4.

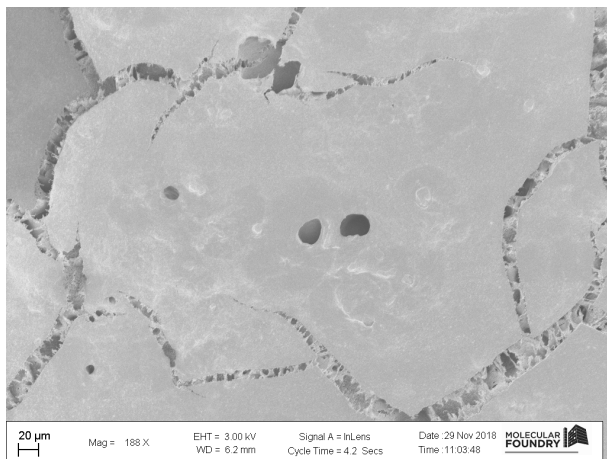




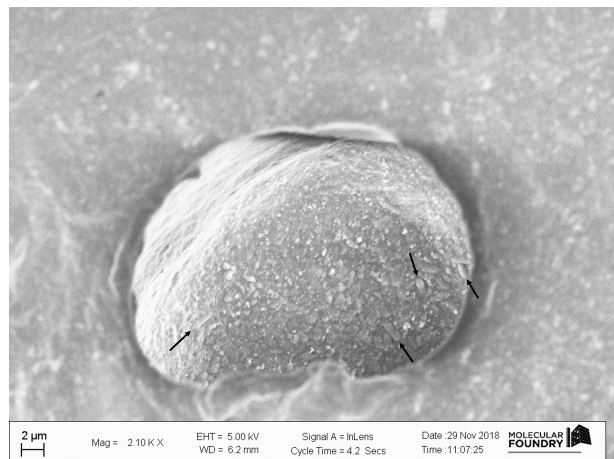
(a)



(b)

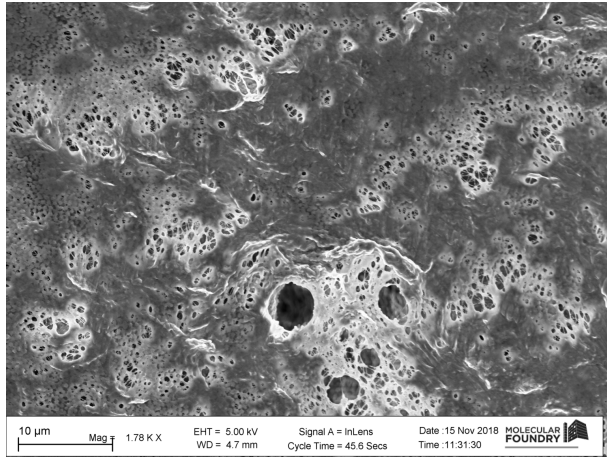


(c)

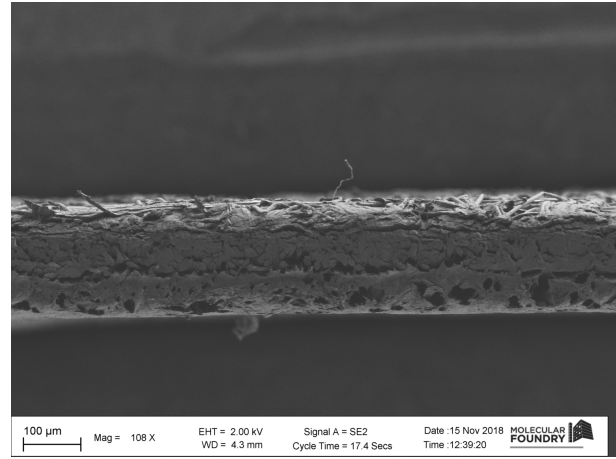


(d)

Figure C.24: SEM images of fouled MF membrane. (a) Wrinkles in the biofilm and cracks in the membrane surface. (b) Rod-shaped bacteria on the outer surface of the active layer. (c) Holes in the membrane and biofilm. (d) Rod-shaped bacteria inside one of the holes shown in (c) (several indicated with arrows; the reader may wish to zoom in; image contrast uniformly enhanced for visibility).

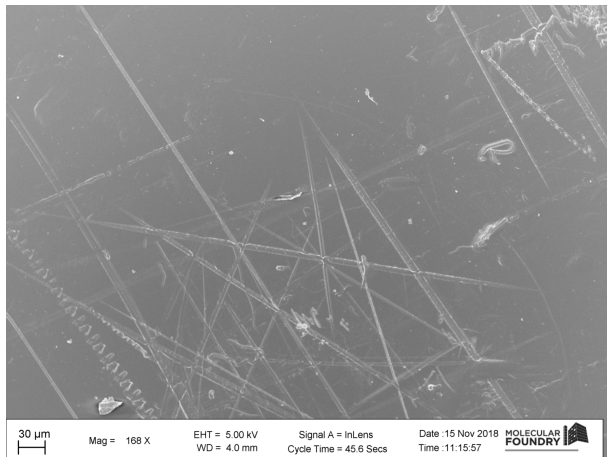


(a) Active layer

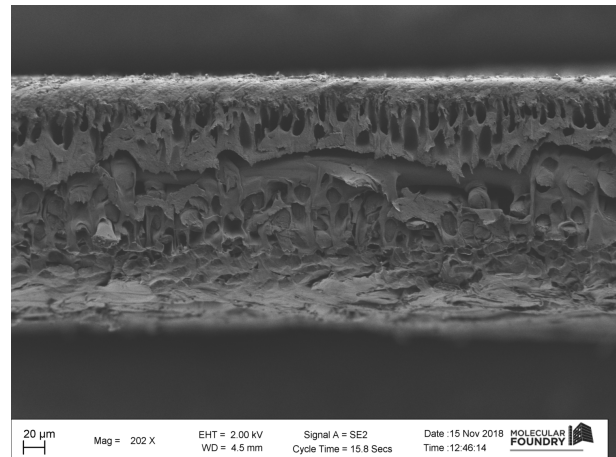


(b) Edge view (active layer facing down)

Figure C.25: SEM images of unfouled MF membrane (PVDF, Synder V0.1).



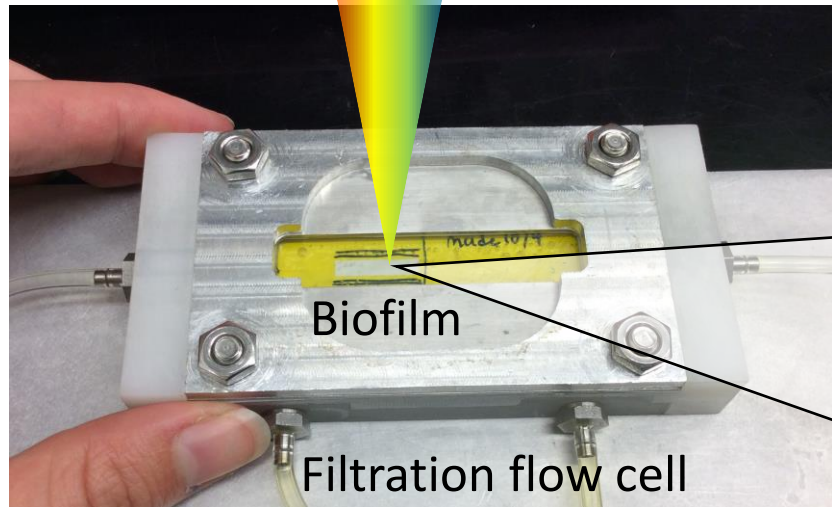
(a) Active layer



(b) Edge view (active layer facing up)

Figure C.26: SEM images of unfouled UF membrane (PES, Microdyn Nadir UP010).

Confocal laser scanning microscopy

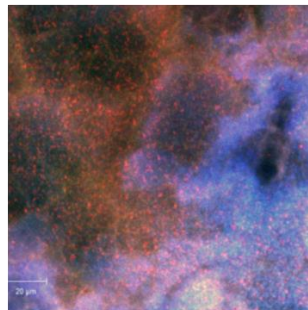


Biofilm

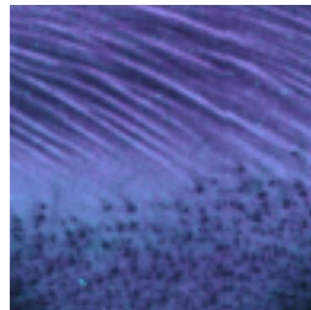
Filtration flow cell

## Biofouling mechanisms visualized in situ

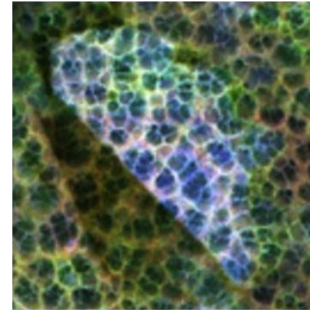
Reproduction



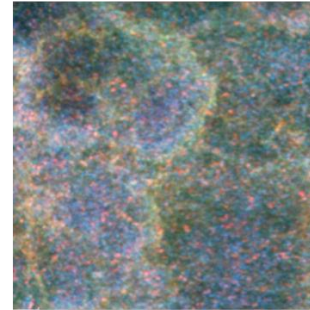
Patterning



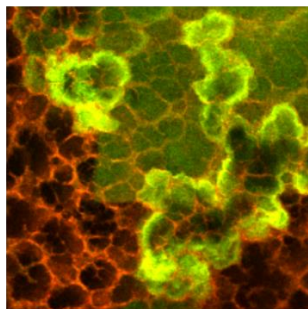
Tearing



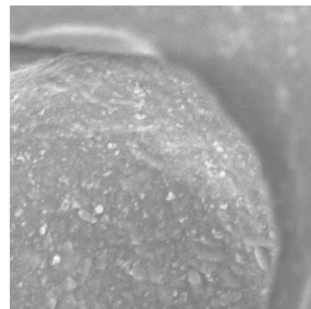
Convection



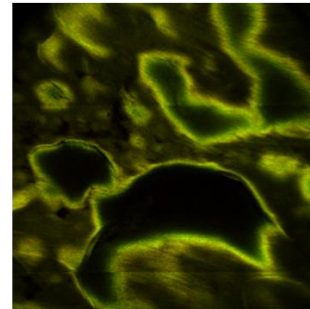
EPS production



Internal fouling



Delamination



Re-deposition

