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Monitoring Extracellular pH in Cross-Kingdom Biofilms using Confocal Microscopy --Manuscript Draft--

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Cover Letter

Dear Dr. Upponi

Attached please find the manuscript "A confocal microscopy based method to monitor extracellular pH in cross-kingdom biofilms" that we would like to submit for publication in JOVE. The manuscript describes the cultivation of cross-kingdom biofilms consisting of *Candida albicans* and *Streptococcus mutans* and reports a methodology for confocal microscopy based assessment of pH inside the biofilms. We provide a detailed description of all steps of the protocol and present typical results. Moreover, we discuss critical steps of the protocol, as well as advantages and limitations of the method and potential future applications.

Thank you very much for your time and consideration.

Sincerely,

Sebastian Schlafer.

TITLE:

Monitoring Extracellular pH in Cross-Kingdom Biofilms using Confocal Microscopy

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KEYWORDS:

bacteria, biofilm, *Candida albicans*, confocal laser scanning microscopy, cross-kingdom, early childhood caries, fungi, pH ratiometry, *Streptococcus mutans*, yeast

SUMMARY:

The protocol describes the cultivation of cross-kingdom biofilms consisting of *Candida albicans* and *Streptococcus mutans* and presents a confocal microscopy-based method for the monitoring of extracellular pH inside these biofilms.

ABSTRACT:

Cross-kingdom biofilms consisting of both fungal and bacterial cells are involved in a variety of oral diseases, such as endodontic infections, periodontitis, mucosal infections and, most notably, early childhood caries. In all of these conditions, the pH in the biofilm matrix impacts microbehost interactions and thus the disease progression. The present protocol describes a confocal microscopy-based method to monitor pH dynamics inside cross-kingdom biofilms comprising *Candida albicans* and *Streptococcus mutans*. The pH-dependent dual-emission spectrum and the staining properties of the ratiometric probe C-SNARF-4 are exploited to determine drops in pH in extracellular areas of the biofilms. Use of pH ratiometry with the probe requires a meticulous choice of imaging parameters, a thorough calibration of the dye, and careful, threshold-based post-processing of the image data. When used correctly, the technique allows for the rapid assessment of extracellular pH in different areas of a biofilm and thus the monitoring of both horizontal and vertical pH gradients over time. While the use of confocal microscopy limits Z-profiling to thin biofilms of 75 µm or less, the use of pH ratiometry is ideally suited for the noninvasive study of an important virulence factor in cross-kingdom biofilms.

INTRODUCTION:

Cross-kingdom biofilms comprising both fungal and bacterial species are involved in several pathologic conditions in the oral cavity. *Candida* spp. have frequently been isolated from endodontic infections¹ and from periodontal lesions^{2,3}. In mucosal infections, streptococcal species from the mitis group have been shown to enhance fungal biofilm formation, tissue

invasion, and dissemination in both in vitro and murine models^{4–7}. Most interestingly, oral carriage of *Candida* spp. has been proven to be associated with the prevalence of caries in children⁸. As shown in rodent models, a symbiotic relationship between *Streptococcus mutans* and *Candidas albicans* increases the production of extracellular polysaccharides and leads to the formation of thicker and more cariogenic biofilms^{9,10}.

In all of the above-mentioned conditions, early childhood caries in particular, the biofilm pH is of importance for disease progression, and the eminent role of the biofilm matrix for the development of acidogenic microenvironments¹¹ calls for methodologies that allow studying pH changes inside cross-kingdom biofilms. Simple and accurate confocal microscopy-based approaches to monitor pH inside bacterial¹² and fungal¹³ biofilms have been developed. With the ratiometric dye C-SNARF-4 and threshold-based image post-processing, extracellular pH can be determined in real-time in all three dimensions of a biofilm¹⁴. Compared to other published techniques for microscopy-based pH-monitoring in biofilms, pH ratiometry with C-SNARF-4 is simple and cheap, because it does not require the synthesis of particles or compounds that include a reference dye¹⁵ or the use of two-photon excitation¹⁶. The use of just one dye prevents problems with probe compartmentalization, fluorescent bleed-through, and selective bleaching^{16–18} while still allowing for a reliable differentiation between intra- and extracellular pH. Finally, incubation with the dye is performed after biofilm growth, which allows studying both laboratory and in situ-grown biofilms.

The aim of the present work is to extend the use of pH ratiometry and provide a method to study pH changes in cross-kingdom biofilms. As proof of concept, the method is used to monitor pH in dual species biofilms consisting of *S. mutans* and *C. albicans* exposed to glucose.

PROTOCOL:

The protocol for saliva collection was reviewed and approved by the Ethics Committee of Aarhus County (M-20100032).

1. Cultivation of cross-kingdom biofilms

1.1 Grow *S. mutans* DSM 20523 and *C. albicans* NCPF 3179 on blood agar plates at 37 °C under aerobic conditions.

1.2 Transfer single colonies of each organism to test tubes filled with 5 mL of brain heart infusion (BHI). Grow for 18 h under aerobic conditions at 37 °C.

1.3 Centrifuge the overnight cultures at 1,200 x g for 5 min. Discard the supernatant, resuspend the cells in physiological saline and adjust the OD_{550 nm} to 0.5 for *C. albicans* (~10⁷ cells/mL) and *S. mutans* (~10⁸ cells/mL). Dilute the *S. mutans* suspension 1:10 with sterile physiological saline (~10⁷ cells/mL).

1.4 Pipette 50 μL of sterile salivary solution, prepared according to the method of de Jong et al. 19,

into the wells of an optical bottom 96-well plate for microscopy. Incubate for 30 min at 37 °C.
 Wash the wells 3x with 100 μL of sterile physiological saline. Empty the wells.

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1.5 Add 100 µL of *C. albicans* suspension to each well. Incubate at 37 °C for 90 min. Wash 3x with sterile physiological saline.

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NOTE: Empty the wells, leaving a reservoir of 20 μ L. Do not empty the wells completely during washing. Leave a reservoir of 20 μ L to avoid excessive shear forces.

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1.6 Add 100 μ L of heat-inactivated fetal bovine serum (inactivated at 56 °C for 30 min) to each well. Incubate at 37 °C for 2 h. Wash 3x with sterile physiological saline. Empty the wells, leaving a reservoir of 20 μ L.

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1.7 Add 100 μ L of *S. mutans* suspension (prepared in step 1.3) to each well. Add 150 μ L of BHI containing 5% sucrose. Incubate at 37 °C for 24 h or longer. When cultivating older biofilms, change the medium daily to fresh BHI. At the end of the cross-kingdom biofilm growth phase, wash 5x with sterile physiological saline.

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2. Ratiometric pH imaging

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NOTE: Ratiometric pH imaging needs to be performed immediately after biofilm growth is complete.

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2.1 For ratiometric pH imaging, use an inverted confocal laser scanning microscope with a 63x oil or water immersion lens, a 543 nm laser line, and a spectral imaging system (i.e., META detector) to allow for the imaging of overlapping fluorescent signals. Use an incubator to warm the microscope stage to 35 °C.

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2.2 Set the detector to ensure the detection of green fluorescence from 576–608 nm and simultaneous detection of red fluorescence from 629–661 nm. Choose an appropriate laser power and gain to avoid over- and underexposure.

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NOTE: Exposure of the images is best seen in palette images with false coloring.

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2.3 Set the pinhole size to 1 Airy Unit or an optical slice of ~0.8 μm. Set the image size to 512 x
 512 pixels and the scan speed to 2. Choose a line average of 2, using the mean option.

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NOTE: At the beginning of a series of experiments, check that the chosen microscope setting provides a clear contrast between bacterial cells, fungal cell walls, biofilm matrix, and fungal cytoplasm.

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- 2.4. Prepare 100 μL of sterile physiological saline containing 0.4% (w/v) of glucose, titrated to pH
- 7. Prepare a stock solution of C-SNARF-4 (1 mM in dimethyl sulfoxide). Add the dye to a final
- 132 concentration of 30 μM.

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2.5 Empty one of the wells with a cross-kingdom biofilm, leaving a reservoir of 20 μL. Add the sterile saline containing glucose and the ratiometric dye. Place the 96-well plate on the microscope stage and start imaging the biofilm.

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2.6 Acquire single images or Z-stacks in different locations of the biofilm. Mark the X-Y position in the microscope software to follow pH changes in particular fields of view over time. At regular intervals, take images with the laser turned off to correct for detector offset.

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2.7 Repeat steps 2.4–2.6 for the analysis of each biofilm grown in a different well.

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3. Calibration of the ratiometric dye

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NOTE: Calibration of the dye and the fitting of a calibration curve can be performed on a different day than ratiometric pH imaging.

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3.1 Prepare a series of 50 mM 2-morpholinoethanesulfonic acid (MES) buffer titrated to pH 4.0-7.8 in steps of 0.2 pH units at 35 °C. Pipette 150 μ L of each buffer solution into the wells of an optical-bottom 96-well plate for microscopy.

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3.2. Add the dye to the buffer-filled wells of the 96-well plate to a final concentration of 30 μ M. Let equilibrate for 5 min.

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3.3 Warm the microscope stage to 35 °C. Choose the same microscope settings as for ratiometric pH imaging. Place the 96-well plate on the microscope stage. Focus on the bottom of the wells. Acquire two images (green and red channel) for all buffer solutions, 5 μ m above the bottom of the well. At regular intervals, take images with the laser turned off to correct for detector offset.

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3.4 Perform the calibration experiment in triplicate.

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3.5 Export all images as TIF files. Import them into dedicated image analysis software (i.e., ImageJ²⁰). Subtract the images taken with the laser turned off from the respective images of the buffer solutions by clicking **Process** | **Image Calculator** | **Subtract** |.

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NOTE: If necessary, crop the images to eliminate artifacts at the image borders by performing rectangular selection using **Image** | **Crop**.

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3.6. Divide the green channel images through the red channel images and calculate the average
 fluorescence intensities in the resulting images by clicking Analyze | Histogram.

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3.7 From the triplicate experiments, plot the average green/red ratios against the pH. Use dedicated software to fit a function to the calibration data (i.e., MyCurveFit).

4. Digital image analysis

NOTE: Digital image analysis can be performed at any time point after calibration of the dye and ratiometric pH imaging.

4.1 Store the green and red channel biofilm images in separate folders and rename both series of files with sequential numbers (e.g., GREEN_0001). Import the images into dedicated image analysis software (i.e., ImageJ). Click **Analyze | Histogram** to determine the average fluorescence intensity in the images taken with the laser off and subtract the value from the biofilm images by clicking **Process | Math | Subtract**.

4.2 Import the two-image series into dedicated image analysis software (i.e., daime²¹). Perform a threshold-based segmentation of the red channel images (**Segment | Automatic segmentation | Custom threshold**). Set the 'low' threshold above the fluorescence intensity of the fungal cytoplasm and the 'high' threshold below the intensity of the fungal cell walls and the bacteria.

NOTE: When appropriate thresholds have been chosen, only extracellular areas are recognized as objects.

4.3 Transfer the object layer of the segmented image series to the green channel image series. To do so, click **Segment | Transfer object layer**.

NOTE: If the contrast between extracellular areas and fungal cytoplasm is too weak in the individual color channels, add the green channel image series to the red channel series prior to segmentation by clicking **Edit** | **Image calculator** | **Addition**. Perform the segmentation as described under 4.2 and transfer the object layer of the segmented image series to both the green and the red channel image series.

4.4 Employ the object editor to delete non-object pixels in the red and green channel image series (Visualizer | Object editor | In all images | Delete non-object pixels). Now the biofilm images are cleared from both bacterial and fungal cells. Export the processed image series as TIF files.

4.5 Import both image series to ImageJ. ImageJ assigns an intensity of 0 to all non-object pixels. Remove those pixels by dividing the red image series (R1) by itself (Process | Image calculator | Image 1: R1; Operation: Divide; Image 2: R1) and multiplying the resulting image series (R2) with the original red image series (Process | Image calculator | Image 1: R1; Operation: Multiply; Image 2: R2). A third image series (R3) is created, identical to R1, except for the fact that NaN is assigned to all pixels with an intensity of 0 in R1. Proceed in the same way with the green image series.

4.6 Use the 'Mean' filter (**Process** | **Filters** | **Mean** | **Radius** | **1 pixel**) on the red and green channel image series to compensate for detector noise. Divide the green channel image series by the red channel image series (**Process** | **Image calculator** | **Image 1** : **G3**; **Operation**: **Divide**;

4.7 Calculate the average ratio for each image (**Analyze** | **Histogram**). Apply false coloring for better visual representation of the ratios in the images (**Image** | **Lookup Tables**). Convert the green/red ratios to pH values employing the function fitted under 3.6.

REPRESENTATIVE RESULTS:

After 24 h and 48 h, robust cross-kingdom biofilms developed in the well plates. *C. albicans* showed varying degrees of filamentous growth, and *S. mutans* formed dense clusters of up to 35 µm in height. Single cells and chains of *S. mutans* grouped around fungal hyphae, and large intercellular spaces indicated the presence of a voluminous matrix (**Figure S1**).

Calibration of the ratiometric dye yields an asymmetrical sigmoidal curve^{13,14}. The extracellular pH in the biofilms dropped quickly in the first 5 min after exposure to glucose at different rates in different microscopic fields of view. Thereafter, acidification slowed down, typically reaching values of 5.5–5.8 after 15 min (**Figure 1**). Replicate biofilms showed a similar behavior, with only slight variations in average pH (**Figure S2**).

The accuracy of the pH calculations is strongly dependent on a careful choice of image settings during acquisition. For the biofilms in question, an optical slice of 0.8 µm proved to be ideal to obtain the best possible contrast between extracellular areas and fungal cytoplasm (Figure S3). Moreover, a pixel size of 0.28 µm (Figure S4), a pixel dwell time of 102.4 µs (Figure S5), and a line average of 2 (Figure S6) provided a good contrast along with an acceptable image acquisition time of ~1 min. During image post-processing, all bacterial and fungal cells were removed from the images, while most of the surrounding extracellular areas were included in the subsequent pH analysis. Depending on the brightness of the images, different upper and lower thresholds had to be chosen (Figure 2).

The pH data shown in **Figure 1** and **Figure S2** were recorded 5 μ m from the biofilm-substratum interface. With increasing distance from the substrate, and depending on the cell density in the biofilms, the fluorescence intensity decreases, resulting in a lower contrast between cells and matrix. However, in the thin biofilms (~35 μ m) grown in the present study, contrast at the top of the biofilm allowed reliable image analysis (**Figure S7A**).

FIGURE AND TABLE LEGENDS:

Figure 1. Use of pH ratiometry in cross-kingdom biofilms exposed to glucose. (A) A field of view in a stained biofilm was imaged with confocal microscopy and the area covered by bacterial and fungal cells was removed prior to ratiometric analysis. (B) The pH in the extracellular space was calculated and visualized using a lookup table (16 colors). Bars = $20 \mu m$. (C) The pH in the 24 h cross-kingdom biofilms dropped rapidly upon exposure to glucose at slightly different rates in different fields of view. Error bars = standard deviation (SD).

Figure 2. Threshold-based image segmentation of stained biofilms. Due to the local pH changes, fluorescence intensity in the biofilms changes over time. (A) and (B) show the same microscopic

field of view after 1 min and 15 min of exposure to glucose, respectively. During image segmentation, high and low thresholds need to be chosen adequately to eliminate all areas covered by bacterial and fungal cells. (**C**) Blue areas were eliminated by the low threshold (40), red areas by the high threshold (115). (**D**) Only extracellular areas were recognized as objects (surrounded by orange lines). (**E**) After elimination of the area covered by microbial cells, the pH calculation could be performed in the biofilm matrix. Bars = $20 \mu m$.

Figure S1. Typical 24 h cross-kingdom biofilm stained with the ratiometric dye. Many *C. albicans* cells displayed filamentous growth, while *S. mutans* (yellow) cells formed dense clusters or localized around fungal hyphae as single cells or chains. Large intercellular spaces indicated the presence of a voluminous biofilm matrix. Bar = $20 \mu m$.

Figure S2. The pH development in 24 h cross-kingdom biofilms exposed to glucose. (A), (B), and (C) The pH was determined ratiometrically in three replicate biofilms for 15 min after exposure to glucose. After a rapid pH drop in the first 5 min, acidification slowed down, and levels of 5.5–5.8 were reached after 15 min. Slightly different rates were observed in different microscopic fields of view (each represented by a line). Calibration of the ratiometric probe was only performed once. Error bars = SD.

Figure S3. Impact of optical slice thickness on image contrast. Images of stained biofilms were acquired with (A) a pinhole size of 2 Airy Units and an optical slice thickness of 1.6 μ m, and (B) 1 Airy Unit and an optical slice of 0.8 μ m. At 1 Airy Unit, a higher laser power/gain was needed for image acquisition, but the contrast between fungal cytoplasm and extracellular areas improved. Bar = 20 μ m.

Figure S4. Impact of resolution on image contrast. Images of stained biofilms were acquired with pixel sizes of (A) 1.12 μ m, (B) 0.56 μ m, (C) 0.28 μ m, (D) 0.14 μ m, and (E) 0.11 μ m. Reduction of the pixel size below 0.28 μ m did not improve the contrast between extracellular areas and fungal cytoplasm. Bars = 20 μ m.

Figure S5. Impact of scan speed on image contrast. Images of stained biofilms were acquired with pixel dwell times of (A) 12.8 μ s, (B) 25.6 μ s, (C) 51.2 μ s, (D) 102 μ s, and (E) 164 μ s. Increasing the pixel dwell time beyond 102 μ s did not improve the contrast between extracellular and intracellular areas. Bars = 20 μ m.

Figure S6. Impact of averaging on image contrast. Images of stained biofilms were acquired with line averages (mean) of (A) 1, (B) 2, and (C) 4. A line average of 2 provided the best compromise between contrast and acquisition time. Bars = $20 \mu m$.

Figure S7. Image contrast in purely fungal biofilms and cross-kingdom biofilms. (A) The top of a stained cross-kingdom biofilm was imaged 30 μm from the interface. The contrast between extracellular and intracellular areas was lower than at the biofilm base but still sufficient for ratiometric pH analysis. (B) A *C. albicans* monospecies biofilm was imaged for pH ratiometry. Laser power/gain could be increased to optimize the contrast between fungal cell walls and

cytoplasm. (C) In cross-kingdom biofilms, the brightly fluorescent bacteria precluded further increase of laser power/gain. Hence, the contrast between fungal cell walls and cytoplasm is less pronounced than in purely fungal biofilms. Bars = $20 \mu m$.

DISCUSSION:

Different protocols for the cultivation of cross-kingdom biofilms involving *C. albicans* and *Streptococcus* spp. have been described previously^{9,22-25}. However, the present setup focuses on simple growth conditions, a time schedule compatible with regular working days, a balanced species composition, and the development of a voluminous biofilm matrix. Moreover, 96-well plates were coated with salivary solution to mimic oral conditions of adhesion to some extent.

The use of pH ratiometry with C-SNARF-4 is an inexpensive method for rapid measurements of biofilm pH, the advantages and disadvantages of which have been discussed in detail elsewhere^{12,26}. In brief, the technique allows for the assessment of pH in different locations inside biofilms and thus the monitoring of horizontal pH gradients and pH developments over time²⁷. Vertical pH profiles can be recorded, too, but the penetration depth is limited by the use of confocal microscopy. Hence, pH ratiometry represents a faster, more versatile, and less invasive alternative to pH microelectrodes, which are currently the method of choice for Z-profiling of thick biofilms²⁸. Compared to other microscopy-based methods for pH recordings in biofilms, pH ratiometry with C-SNARF-4 has the advantage of only employing one stain. Therefore, several problems that may arise from the differential fluorescent behavior and the interaction between different dyes are circumvented²⁶.

In cross-kingdom biofilms, threshold-based differentiation between microbial biomass and extracellular areas poses some problems when compared to biofilms that only comprise bacterial or fungal cells. Bacterial cells internalize the ratiometric dye and display a bright fluorescent signal compared to the biofilm matrix, but also compared to fungal cell walls. Fungal cell walls appear somewhat brighter than the biofilm matrix, which in turn shows higher fluorescence than the fungal cytoplasm. In biofilms that only consist of fungal cells, images can be acquired with a high laser power/gain, which results in sufficient contrast between the biofilm matrix and the fungal cytoplasm (**Figure S7B**). When bacteria are present, laser power/gain must be reduced to avoid overexposure of bacterial cells (**Figure S7C**). Hence, the contrast between fungal cytoplasm and extracellular areas is not as pronounced, and image settings such as pinhole size, pixel size, and pixel-dwell time must be chosen with great care for ratiometric analysis.

As for all microscopy-based techniques, image quality and acquisition time are inversely correlated. In the present cross-kingdom biofilms, an imaging time of ~30 s, ideally 1 min, was necessary to obtain an appropriate quality for subsequent pH analysis. In comparison, biofilms that only harbor bacteria can be imaged with sufficient quality in 10 s or less²⁹. For microscopy analyses of biofilm growth, structure, or composition, acquisition times of 1 min do not pose a problem, and in many instances, this may also apply to pH recordings. However, extreme pH changes, such as the rapid acidification observed immediately after exposure to glucose ($\Delta pH > 1$ unit/min), are difficult to monitor in cross-kingdom biofilms.

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353 The present work demonstrates that ratiometric pH recordings with C-SNARF-4 are feasible in 354 cross-kingdom biofilms composed of S. mutans and C. albicans. Future studies may employ the 355 method to analyze pH changes in cross-kingdom biofilms with greater species diversity, especially in biofilms grown in situ (i.e. in children with early childhood caries)³⁰. In this study, the pH in the 356 357 biofilms was monitored under static conditions and for a limited observation time of 15 min, right 358 after a glucose pulse. Further advancements may include the application of a fluid flow to mimic 359 in situ conditions more closely^{14,31,32}. Moreover, pH ratiometry may be used to study the impact of various nutritional conditions on long-term pH changes in cross-kingdom biofilms and 360 361 contribute to elucidate the effect of biofilm pH on the underlying host tissues. Again, the 362 demineralization of dental enamel in early childhood caries may serve as a prominent example.

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DISCLOSURES:

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The authors have nothing to disclose.

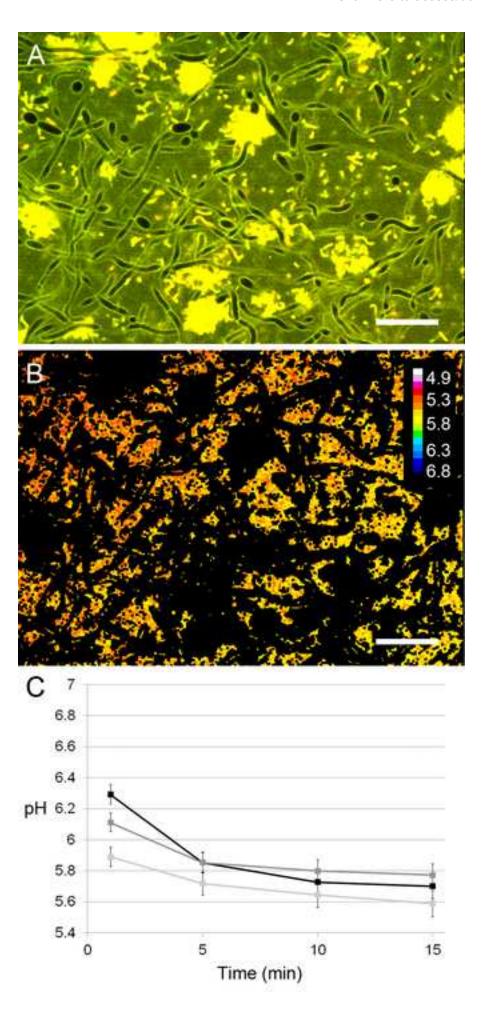
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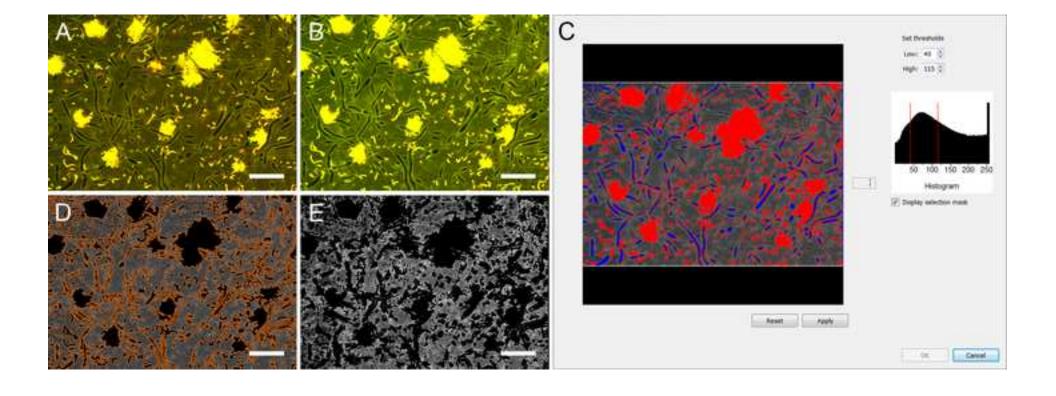
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Name of Material/Equipment	Company	Catalog Nu	umber
Blood agar plates	Statens Serum Institut		677
Brain heart infusion	Oxoid	CM1135	
Brain heart infusion + 5 % sucrose	BDH laboratory supplies		10274
Candida albicans	National Collection of Pathogenic	NCPF 3179	
	Fungi		
D-(+)-Glucose	Sigma-Aldrich	G8270	
daime: digital image analysis in microbial ecology	Universität Wien	N/A	
Dimethyl sulfoxide	Life Technologies	D12345	
Fetal bovine serum	Gibco Life technologies		10270
GS-6R refrigerated centrifuge	Beckman	N/A	
ImageJ	National Institutes of Health	N/A	
Java	Oracle	N/A	
μ-Plate 96 Well Black	Ibidi		89626
MyCurveFit	MyAssays Ltd.	N/A	
2-(N-Morpholino)ethanesulfonic acid (MES) buffer	Bioworld		700728
PHM210 pH-meter	Radiometer Analytical		
Plan-Apochromat 63x oil immersion objective	Zeiss	N/A	
SNARF®-4F 5-(and-6)-Carboxylic Acid	Life Technologies	S23920	
Sterile physiological saline	VWR		6404
Streptococcus mutans	Deutsche Sammlung von	DSM 20523	
	Mikroorganismen und Zellkulturen		
Vis-spectrophotometer V-3000PC	VWR	N/A	
XL Incubator	PeCON	N/A	
Zeiss LSM 510 META	Zeiss	N/A	

Comments/Description

Freeware; V2.1; https://dome.csb.univie.ac.at/daime

Freeware; V1.46r; https://imagej.nih.gov/ij

Freeware necessary to run ImageJ; V8.0; https://java.com/en/download

NA=1.4



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Reply to reviewers:

Reviewer #1:

Manuscript Summary:

The authors describe an innovative method for monitoring extracellular pH in mixed bacterial/fungal biofilms. The methods are sufficiently described and I recommend to publish this work. I have no further comments.

Major Concerns:

none

Minor Concerns:

none

Reviewer #2:

Manuscript Summary:

This manuscript provides a detailed and informative explanation of a method for assessing extracellular pH in interkingdom biofilms relevant to the oral cavity. The Introduction sets the context well, and the methods are clearly explained with helpful notes where needed.

Major Concerns:

There are no major concerns.

Minor Concerns:

1. Since the method relies heavily on software, it would be very helpful to give details of the software versions that were used along with relevant websites to obtain the software. I believe that ImageJ runs on Java, so this should also be included. Undoubtedly, there will be options for different types/versions of software, but it is useful to give an example of a setup that is known to work.

Answer: We now provided the software versions for daime, ImageJ and Java, as well as links to the websites, in the Table of Materials.

2. Protocol 1.3: Is C. albicans predominantly in yeast form at this point, or are hyphae present?

Answer: After planktonic growth C. albicans is predominantly in yeast form.

3. Protocol 1.6: Is it important to use heat inactivated fetal bovine serum?

Answer: We used heat inactivated fetal bovine serum to eliminate any potential antimicrobial effect of the complement system. We added this detail to the protocol.

4. Prot. 2.1: What is a 'META' detector?

Answer: The META detector employs a spectral imaging system to simultaneously detect fluorescence emissions from overlapping fluorophores. We added a short explanation to the protocol.

5. Prot. 3.1: It seems strange to use MES buffer for such a wide range of pH since MES only buffers effectively between pH 5.5-6.7. Why not use a broader buffer system such as citric acid/Na phosphate?

Answer: Thank you for the comment. We have employed MES buffer for calibration, as its range fits nicely with the pH values typically encountered in dental biofilms (pH 5-7). Adjusting the solutions to low pH takes a bit of time, but since the solutions contain only MES buffer and C-SNARF-4, the pH is stable across the entire range. In previous reports, HEPES buffer solutions (pK_a=7.31) have been employed for calibration (Hunter & Beveridge; Appl Environ Microbiol. 2005 May;71(5):2501-10). We will consider using citric acid/Na phosphate in future studies, provided that it does not interfere with the fluorescence emission of C-SNARF-4.

6. Figure S5: What is 'xx'?

Answer: Thank you for spotting the mistake. We now inserted 102 µs.

7. References: Full details of references are needed (e.g. refs 13 and 14).

Answer: We have now added missing details in the reference list.

