

Staphylococcus aureus Induces Hypoxia and Cellular Damage in Porcine Dermal Explants

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We developed a porcine dermal explant model to determine the extent to which *Staphylococcus aureus* biofilm communities deplete oxygen, change pH, and produce damage in underlying tissue. Microelectrode measurements demonstrated that dissolved oxygen (DO) in biofilm-free dermal tissue was 4.45 ± 1.17 mg/liter, while DO levels for biofilm-infected tissue declined sharply from the surface, with no measurable oxygen detectable in the underlying dermal tissue. Magnetic resonance imaging demonstrated that biofilm-free dermal tissue had a significantly lower relative effective diffusion coefficient (0.26 ± 0.09 to 0.30 ± 0.12) than biofilm-infected dermal tissue (0.40 ± 0.12 to 0.48 ± 0.12 ; $P < 0.0001$). Thus, the difference in DO level was attributable to biofilm-induced oxygen demand rather than changes in oxygen diffusivity. Microelectrode measures showed that pH within biofilm-infected explants was more alkaline than in biofilm-free explants (8.0 ± 0.17 versus 7.5 ± 0.15 , respectively; $P < 0.002$). Cellular and nuclear details were lost in the infected explants, consistent with cell death. Quantitative label-free shotgun proteomics demonstrated that both proapoptotic programmed cell death protein 5 and antiapoptotic macrophage migration inhibitory factor accumulated in the infected-explant spent medium, compared with uninfected-explant spent media (1,351-fold and 58-fold, respectively), consistent with the cooccurrence of apoptosis and necrosis in the explants. Biofilm-origin proteins reflected an extracellular matrix-adapted lifestyle of *S. aureus*. *S. aureus* biofilms deplete oxygen, increase pH, and induce cell death, all factors that contribute to impede wound healing.

Staphylococcus aureus causes a diversity of infections in both humans and animals, ranging from minor skin infections to life-threatening toxic shock syndrome, endocarditis, and septicemia in humans (1). It is one of the most common bacterial pathogens found in both acute and chronic wounds, and its treatment is complicated by the emergence of drug-resistant strains commonly known as methicillin-resistant *S. aureus* (MRSA) (2–4). Formation of biofilms in wounds is a major cause of nonhealing wounds (5–7) and treatment failure (8). Oxygen and pH are two critical factors in wound healing (9–12), but at present we have limited knowledge about how the presence of *S. aureus* biofilms affects oxygen and pH levels in wounds.

Oxygen availability is essential for wound healing (9, 10) and is required for optimal function of leukocytes and fibroblasts (13, 14). Oxygen enhances leukocyte chemotaxis, phagocytosis, and antibacterial activities. Killing of bacteria by leukocytes is compromised when oxygen pressure drops below 40 mm Hg (~1.81 mg/liter) (15, 16). A minimum oxygen level is required for replication and proper functioning of cells, including fibroblasts, which play a critical role in wound repair. For example, fibroblasts can replicate at oxygen levels below 25 mm Hg (~1.13 mg/liter), but their capacity for hydroxylation of proline and lysine, which is required for collagen synthesis, is impaired. Cell replication is arrested at oxygen levels below 0.5% (approximately 0.16 mg/liter) (17). Furthermore, angiogenesis in chronic wounds also requires oxygen. Consequently, hypoxia can impair or prevent wound healing through several mechanisms (18). Because biofilm communities consume oxygen, we surmise that *S. aureus* biofilms negatively impact oxygen content in tissues underlying the wound surface.

Similarly, mass transport of essential nutrients in the tissue will influence the healing process. Mass transfer of O₂ and nutrients can be characterized by a parameter known as the effective diffusion coefficient. To our knowledge, this parameter has never been measured in tissues underlying biofilms.

Normal wound healing proceeds by tissue degradation and reassembly and therefore requires various enzymes and proteins. Enzymes function optimally in a defined pH range. In chronic wounds, for example, pH increases from 7.5 to 8.9, and this enhances the activity of proteolytic enzymes. As healing ensues, wound pH declines to around 6, where the activity of enzyme inhibitors is optimal (11, 19, 20). Nothing is known about the specific contribution to the pH of chronic wounds by various types of bacterial biofilms in general and *S. aureus* biofilms in

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particular, despite *S. aureus* being an important wound pathogen. In addition to oxygen and pH, successful wound healing requires basic functions from surviving cells. While it has been shown that *S. aureus* can kill isolated blood cells (erythrocytes and monocytes) and various cultured cells, little is known about how *S. aureus* biofilms impact host cells within the context of tissue.

We hypothesized that *S. aureus* biofilms can deplete dissolved oxygen and change the pH of tissue and that these changes together with secretion of bacterial products into the tissue result in cell and underlying tissue damage. To test this hypothesis, we developed a porcine explant dermal model that is conducive to measuring these and related parameters. Explants were 500- μm -thick samples of dermis from pig ears, which we selected in part because porcine skin closely resembles human skin (21) and pigs are generally preferable models for studying human infectious diseases (22). The biofilms were grown on dermal tissue, and microelectrodes were used to measure dissolved oxygen and pH profiles throughout the depth of the biofilms and the tissue. In addition, we estimated relative effective diffusion coefficients using pulsed-field gradient nuclear magnetic resonance imaging (or magnetic resonance microimaging) to understand if *S. aureus* biofilms negatively impact mass transfer processes. Cytotoxic effects of biofilm were examined by histology, and proteins released as a result of biofilm–dermal-tissue interaction were identified by quantitative label-free shotgun proteomics. The explant model allowed us to accurately determine these parameters, which are either difficult or impossible to determine accurately in an *in vivo* animal model.

MATERIALS AND METHODS

Explant preparation and infection. Ears (pinnae) were collected from healthy pigs (*Sus scrofa domestica*) within 1 h of slaughter at a local abattoir and transported to the lab on ice (no study-specific animals were purchased or housed for the work reported herein). The ears were rinsed with water, dried with a paper towel, clipped, and shaved. Ear tissue was then lightly wetted with 70% ethanol and then sectioned longitudinally in 2- to 3-cm-wide strips using a sterile surgical blade. A 500- μm -thick slice of skin was removed from each strip using a Padgett's dermatome (Nouvag, Goldach, Switzerland). After removal of the epidermis, one more layer of 500- μm thickness containing mostly dermis was sliced from each strip. These dermal strips were placed in petri dishes containing ice-cold phosphate-buffered saline (PBS; pH = 7.2). Discs 12 mm in diameter were punched out from each dermal strip using surgical biopsy punches. The discs were placed in insert wells (Greiner Bio-one Inc., Monroe, NC, USA) on a membrane (0.4- μm pore size). Care was taken to ensure that the inner surface (in reference to intact skin) of the discs rested on the membrane for proper nutrient acquisition by the explants (Fig. 1). The insert wells were placed in 6-well plates that were pre-filled with 2 ml of Dulbecco's modified Eagle medium (DMEM) with high glucose (HyClone Laboratories Inc., Logan, UT, USA) containing the antibiotics streptomycin (100 $\mu\text{g}/\text{ml}$), penicillin (100 units/ml), and chloramphenicol (5 $\mu\text{g}/\text{ml}$) and the antifungal amphotericin B (0.4 $\mu\text{g}/\text{ml}$). The spent explant medium was replaced with fresh medium on the 4th and 7th days after initial explant culture. pH and lactate dehydrogenase (LDH) activity was measured with a CytoTox 96 kit (Promega Corporation, Madison, WI, USA), described below, from a fraction of the spent medium, and the remaining medium was stored at -80°C for other analyses. All equipment was either autoclaved or disinfected before use. The work was primarily carried out in a biological safety cabinet to limit the probability of external contamination of the explants.

Explants were infected with a green fluorescent protein (GFP)-expressing *S. aureus* strain (strain ALC7559) that was derived from *S. aureus* LAC (USA300), a well-characterized clinical strain (23). ALC7559 carries

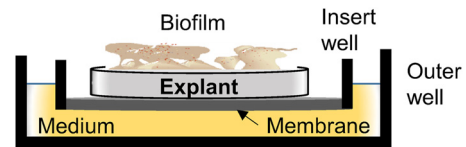


FIG 1 Cross-sectional schematic of a culture well with inner membrane cup, explant tissue, and biofilm.

the plasmid pALC1743 (24), with genes for GFP expression and ampicillin (Amp^r) and chloramphenicol (Cm^r) resistance. Frozen stocks of bacteria were streaked onto Luria-Bertani (LB; Becton Dickinson and Company, Sparks, MD, USA) agar containing chloramphenicol (5 $\mu\text{g}/\text{ml}$), penicillin (100 $\mu\text{g}/\text{ml}$), and streptomycin (100 $\mu\text{g}/\text{ml}$) and incubated overnight at 37°C . A small portion of the bacterial growth on the agar surface was scraped off and suspended in 1 ml of PBS. A 5- μl volume, containing approximately 500 to 800 CFU, of this suspension was then deposited onto the center of each explant to initiate a local infection and development of a biofilm. CFU were confirmed from each bacterial suspension used to infect the explants.

Growth of *S. aureus* in DMEM. To determine if *S. aureus* grows in the explant culture medium, cells were grown overnight on LB agar, and the resultant cells were detached and retrieved by washing with PBS (1 ml). After adjustment of the optical density at 600 nm (OD_{600}) to 0.7 unit, 50 μl of the suspension was added to phenol red-free DMEM with high glucose (950 μl) and LB broth (950 μl). Aliquots (200 μl) from each of the two inoculated media were separately dispensed into 100-well honeycomb plates (Growth Curves USA, Piscataway, NJ, USA). The plates were loaded into a Bioscreen C plate reader (Growth Curves USA) to record growth (OD_{600}) at 37°C every 30 min.

Assessing explant viability. Explant viability was assessed using PrestoBlue cell viability reagent (Invitrogen Corporation, Carlsbad, CA, USA), which is a resazurin-based cell viability reagent. Explants were incubated in 300 μl of 10% PrestoBlue (in DMEM with antibiotics) for 2 h at 37°C in a 5% CO_2 incubator. After incubation, the absorbance of the medium was measured at 570 and 600 nm. Percent reduction of PrestoBlue was calculated from absorbance and molar extinction coefficients of oxidized and reduced PrestoBlue.

Diffusion coefficient in explants. Pulsed-field gradient nuclear magnetic resonance (PFG-NMR) was used to measure diffusion coefficients and generate relative effective diffusion coefficient profiles through the *Staphylococcus aureus* biofilm and explant tissue, similar to the method described by Renslow et al. (25). Specifically, the NMR experiments were conducted at 500.40 MHz for proton (^1H) detection using an 89-mm wide-bore 11.7-T magnet with a Bruker Avance III digital NMR spectrometer (Bruker Instruments, Billerica, MA). Bruker ParaVision 5.1 imaging software was used to collect and process the data for the diffusion tensor imaging method (DtiStandard) (26). This method was run with a 750-ms repetition time, an echo time of 15.5 ms, and 128 averages. The pulse gradient width (δ) was 2 ms, and the diffusion time interval (Δ) was 10 ms. Signal intensity was measured for seven independent b factors: 0, 100, 250, 425, 625, 850, and 1050 s/mm^2 . The field-of-view dimensions were 5 mm in the biofilm/explant depth direction and 5 mm in the perpendicular direction, with a 500- μm -thick slice. A total of 64 complex points were sampled in the depth direction, with 32 phase-encoding steps, for an in-plane resolution of 78 μm by 156 μm . The total experiment time per depth profile was 307 min. Diffusion coefficients were calculated using semilogarithmic regression of the b -factor-dependent intensity value of each measurement pixel in accord with the Block-Torrey differential equation (26, 27). Depth profiles were generated by averaging and performing statistical analysis on the diffusion coefficients in the middle 780 μm of the sample. Therefore, the 78- μm -resolution depth profiles represent the center 780- by 500- μm^2 region of the biofilm/explant. PFG-NMR measures diffusion coefficient of water. The measured diffusion coefficient was normalized by dividing the value at a local point by the diffusion

coefficient in the bulk. This normalized diffusion coefficient is called the relative effective diffusion coefficient (28).

Histological assessment of explants. A subset of explants from days 1, 4, and 7 were fixed in 10% neutral buffered formalin, dehydrated by exposure to increasing concentrations of ethyl alcohol, embedded in paraffin, and sectioned to a thickness of 5 μm . The tissue sections were mounted on glass slides, deparaffinized, hydrated, and stained with hematoxylin and eosin according to standard protocols followed by the Washington Animal Disease Diagnostic Laboratory, Washington State University, Pullman, WA. Resulting light micrographs were interpreted by a board-certified veterinary anatomic pathologist.

Biofilm imaging. Biofilms on 4- and 7-day explants were examined using a Leica SP5 confocal microscope (Leica Microsystems Inc., Deerfield, IL, USA). GFP-labeled bacteria were imaged using 488-nm excitation and 510-nm emission light. Biofilm structure was calculated using Image Structure Analyzer software, developed by our research group (29). We used environmental scanning electron microscopy (ESEM) to image cells on explants at the end of experiments. The samples were first fixed in a solution of glutaraldehyde (2%) and paraformaldehyde (2%) in 0.1 M cacodylate buffer at 4°C. The samples were rinsed with 0.1 M cacodylate buffer twice for 10 min and then treated with osmium tetroxide (2% in 0.1 M cacodylate buffer) overnight at 4°C. Afterwards the samples were rinsed twice in 0.1 M cacodylate buffer for 10 min followed by rinsing with double-distilled water twice for 10 min. The samples were examined using a Quanta 200F (FEI, Hillsboro, OR, USA) scanning electron microscope in environmental mode, in which the samples were frozen to -1°C and the chamber humidity was adjusted to 35% before the electron beam was switched on for scanning the sample surfaces.

Biofilm and explant oxygen and pH depth profiles. We followed previously described protocols to construct microelectrodes (30). We used dissolved-oxygen microelectrodes with a tip diameter $<20\ \mu\text{m}$ in diameter. The dissolved oxygen microelectrodes were polarized to $-800\ \text{mV}_{\text{Ag}/\text{AgCl}}$. Prior to use, the microelectrodes were calibrated using two-point calibration: in the air (oxygen saturation) and in Na_2SO_3 solution (zero oxygen concentration). pH microelectrodes were constructed using a liquid ion-exchange (LIX) membrane tip (28). We used an external Ag/AgCl reference electrode. The pH microelectrodes were calibrated in buffer solutions (pH = 4, 7, and 10).

A Keithley 6517A electrometer/high-resistance meter was used to operate the dissolved-oxygen or pH microelectrode. The surface of the biofilm/explant sample was located using a Zeiss Stemi 2000 stereomicroscope (Carl Zeiss Microscopy, Thornwood, NY, USA). The microelectrode tip was lowered with 10- μm step sizes into the biofilm/explant sample. A stepper motor (PI M-230.10S, part no. M23010SX; Physik Instrumente, Auburn, MA), controlled by custom MicroProfiler software, was used to manipulate the movement of the microelectrodes. Collected data were recorded using an analog/digital converter (ADC; USB-1608FS; Measurement Computing, Norton, MA). The profiles showing the change of oxygen or pH by depth are called “depth profiles.” The pH in spent medium was also measured using a pH electrode (Mettler Toledo, Columbus, OH, USA).

Lactate dehydrogenase activity in spent media. Lactate dehydrogenase activity in the spent medium was assayed using a CytoTox 96 kit (Promega Corporation). Reconstituted substrate mix (50 μl) was added to 50 μl of diluted spent medium, and the solution was incubated at room temperature for 30 min. The reaction was stopped by adding stop solution provided in the kit, and the OD_{490} was recorded. The LDH activity of the test samples was calculated from standard curve that was prepared using an LDH-positive control (1,600 U of LDH/ml).

Soluble proteins in the spent media. Soluble proteins in the spent medium were identified by using ultraperformance liquid chromatography/high-bandwidth mass spectrometry (UPLC/MS^E). The proteins in the spent medium from *S. aureus*-infected and *S. aureus*-free 4-day explants were split and concentrated into two fractions using ultrafiltration devices (Merck Millipore Ltd., Tullagreen, Ireland). One fraction contained proteins of $>50\ \text{kDa}$, and the other contained proteins within a

range of 3 to 50 kDa. The DMEM in each sample was replaced with PBS by diluting the medium using an excess of PBS and then filtering the diluted suspension through an appropriate ultrafiltration device. A minimum of three washes in PBS were given to each sample. Albumin in the $>50\text{-kDa}$ fraction was removed using a Pierce albumin depletion kit (Thermo Scientific, Rockford, IL, USA) according to the instructions of the manufacturer.

The concentrated proteins were quantified using the Qubit protein assay kit (Invitrogen) in accordance with the supplier’s protocol for the Qubit 2.0 fluorometer (Invitrogen). Proteins from each sample (100 ng) were digested with trypsin and then analyzed by using a Synapt G2-S HDMS mass spectrometer (Waters Corp., Milford, MA, USA) coupled with a nanoAcquity UPLC system (Waters Corporation). Eluted proteins were identified using a quantitative label-free shotgun proteomic strategy with data-independent scanning (MS^E) and ion mobility spectrometry. Data were processed and searched using TransOmics informatics for proteomics (Waters Corporation), and all searches were performed against a protein database of *Staphylococcus aureus* and *Sus scrofa* from UniProt 2013_09. Quantities of various proteins in infected- and uninfected-explant media were compared using the TM4 microarray software suite (31).

Statistical analysis. Data obtained from multiple explants from a minimum of three different animals were used in the statistical analysis. Unless otherwise stated, we used a *t* test and one- or two-factor analysis of variance (ANOVA) to test the significance among groups, and all the pairwise multiple comparisons were carried out using the Tukey or Holm-Šidák method or with a Bonferroni correction, as applicable. SigmaPlot v. 12.0 (SYSTAT Software Inc.) and the TM4 software suite (31) were used for statistical analyses.

RESULTS

To validate our model, we first verified that *S. aureus* cannot sustain growth in the culture medium (DMEM) and that therefore any biofilm growth is dependent on essential nutrients from the dermal explants (Fig. 2). In addition, we verified that biofilm-free explants were capable of reducing PrestoBlue, a resazurin-based cell viability testing reagent. Dye reduction was evident for up to 7 days ($\sim 70\%$ reduction by day 1, $\sim 60\%$ by day 4, and $\sim 35\%$ by day 7) (see Fig. S1 and Table S1 in the supplemental material). Viable cells reduce PrestoBlue resazurin, a blue compound, to resorufin, a red fluorescent compound. While this assay does not provide an absolute count of viable cells for these tissues, a greater magnitude of chemical reduction is consistent with a greater magnitude of metabolic activity and, presumably, a larger number of viable cells in the explants.

***S. aureus* biofilms increase relative effective diffusion coefficient in dermal tissue.** In live uninfected dermal explants, the relative effective diffusion coefficient dropped from 1.00 near the superficial surface to approximately 0.29 near the bottom 100 μm , consistent with a sigmoidal distribution (Fig. 3). In *S. aureus*-infected explants, these profiles were significantly greater than those in uninfected explants ($P < 0.0001$), indicating that biofilms actively enhance diffusion in the underlying dermal tissue.

***S. aureus* biofilms kill dermal cells.** Histopathological examination of 1-, 4-, and 7-day uninfected explants showed mild changes in cell and nuclear morphology in the more metabolically active cells. These changes included mild vacuolation of myofibers in the tunica muscularis of arterioles, pyknosis and nuclear fragmentation of endothelial cells, and occasional small clusters of perivascular lymphocytes. In *S. aureus*-infected explants, the dermal architecture was diffusely intact; however, nearly all cells were hypereosinophilic. Nuclei were nearly completely absent, indicat-

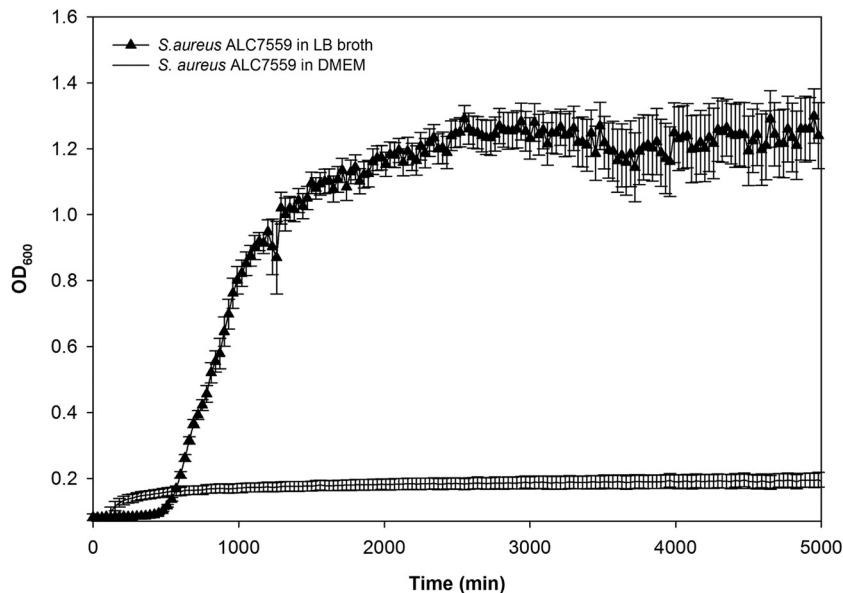


FIG 2 Comparison of growth curves of *S. aureus* (strain ALC7559) grown in DMEM and LB broth (15 independent replicates; vertical bars represent standard deviations [SD]).

ing widespread cell death (Fig. 4). There was no difference in cytotoxicity between 4- and 7-day biofilm-infected explants.

***S. aureus* biofilms deplete oxygen in dermal tissue.** The lowest oxygen concentration (4.45 ± 1.17 mg/liter) was found in high-viability-score, 4-day, biofilm-free explants (Table 1), which corresponds to the approximate partial pressure of oxygen found in most tissues at the capillary level in a resting individual (32). The 7-day biofilm-free explants showed statistically nonsignificant increases in oxygen compared with 4-day explants. The explants with no viable dermal cells had statistically significant ($P < 0.05$) oxygen content (5.87 ± 0.73 mg/liter) compared with 4- and 7-day live dermal tissue (Table 1). With *S. aureus* biofilms growing on the top of explants, oxygen was depleted within the first 50 μm of depth, leaving no measurable oxygen in deeper parts of the biofilms or in the explants; a typical example is shown in Fig. 5 (this result was replicated independently three times). We found similar profiles for day 7 biofilms on explant (results not shown).

Changes in *S. aureus* biofilms with age. Electron microscopy showed that 4-day biofilms were patchy, but these patches coalesced by day 7 (Fig. 6). The bacterial cells in 4-day biofilm were healthy in appearance and rounded, whereas by day 7, the biofilm surface was flat and composed of both lysed and intact bacterial cells that had a flattened appearance (Fig. 6). Confocal laser scanning microscopy (CLSM) examination of dermal biofilms also confirmed bacterial growth interrupted by small spaces in 4-day biofilms, while the growth was dense in 7-day biofilms (see Fig. S2 in the supplemental material). Biofilm structural and volumetric parameters were measured from multiple CLSM images of 4- and 7-day biofilms. The average diffusion distance, defined as the average of all distances between cell clusters and their nearest void area in a biofilm (29), was significantly greater (1.7-fold; $P < 0.05$) in 7-day biofilm, indicating a progressive coalescence of cell clusters with time (Table 2). The biovolume (total volume of the biofilm biomass), and the average biofilm and biomass thicknesses were also greater (about 2-fold) in day 7 biofilm than day 4 biofilm

(Table 2). Biofilm thickness is the volume of the biofilm divided by the total surface area of the substratum, including the areas free of bacterial cell clusters, whereas the average biomass thickness is the volume of the biofilm divided by only that area of the substratum covered by the cell clusters (29). These values nearly doubled in the 7-day biofilms. No significant difference was observed in biovolume-to-surface area ratio between 4- and 7-day biofilms, indicating a uniform bacterial growth in all directions. This is expected in the case of *S. aureus*, which multiplies in all spatial planes. Despite the volumetric and morphological differences between 4- and 7-day biofilms, biofilms completely depleted oxygen from the top to the bottom of the explants for both time points.

In hypoxic tissues, one would expect larger amounts of lactate dehydrogenase and lower pH (33–35). Under hypoxic conditions, LDH converts pyruvate, generated in glycolysis, to lactate, thus decreasing pH (36). The enzyme is also released in larger amounts by damaged cells (37). We measured LDH activity in the spent media from both uninfected and *S. aureus*-infected explants. As expected, significantly greater LDH activity was detected in spent media from infected explants ($1,892 \pm 39$ mU/ml from 4-day and $1,255 \pm 126$ mU/ml from 7-day explants; $P < 0.001$) than in media from uninfected explants (467 ± 68 mU/ml from 4-day and 430 ± 90 mU/ml from 7-day explants) (see Fig. S3 in the supplemental material). Proteomic analysis of the spent media from infected explants revealed the presence of bacterial-origin LDH (see Table S3). This bacterial-origin LDH could contribute to the higher activity of LDH detected in the media of infected explants.

***S. aureus* biofilms increase pH in dermal tissue.** Because pH can influence wound healing by affecting enzymes and cells involved in the healing process (11, 19), we measured pH depth profiles inside the biofilm and dermal explants. The pH was always significantly more alkaline ($P < 0.002$) inside the biofilm and the biofilm-infected dermal tissue than in uninfected tissue (8.08 ± 0.03 versus 7.62 ± 0.04 in 4-day explants, and 7.83 ± 0.16 versus

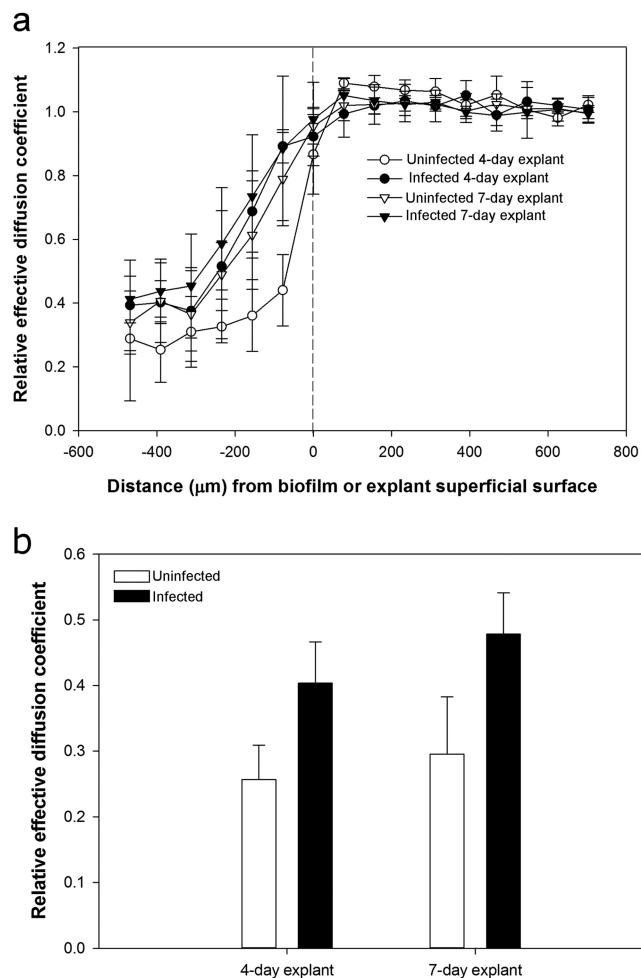


FIG 3 Relative effective diffusion coefficient depth profiles through uninfected and *S. aureus*-infected dermal explants. Relative effective diffusion coefficient is the ratio of the diffusion coefficient of water in biological material (explants) to the diffusion coefficient of water in bulk fluid (DMEM). (a) The biofilm or explant superficial surface begins at 0 μm , and the increasing negative numbers indicate increasing depths within the explants. Error bars represent SD for 3 independent NMR measurements. (b) The relative effective diffusion coefficient was significantly higher in *S. aureus*-infected explants than in uninfected control explants (two-way ANOVA, Holm-Sidak method; $P < 0.0001$; the model included explant age [4 day and 7 day] plus infection [uninfected and infected] plus interaction; the latter was not significant [$P = 0.40$]).

7.39 ± 0.13 in 7-day explants) (Fig. 7). The pH in the spent media was always significantly more alkaline for media from infected dermal explants than biofilm-free dermal explants ($P < 0.001$) (see Fig. S4 in the supplemental material). These findings were unexpected given that the presence of more LDH with infected biofilms should have made the medium more acidic.

***S. aureus*-infected dermal tissue releases pro- and antiapoptotic porcine proteins and extracellular matrix-colonizing *S. aureus* proteins.** *S. aureus*-infected explants released ~ 2.5 -fold more protein in the spent medium than uninfected explants. A comparison of soluble proteins from infected and uninfected explants showed a greater diversity of proteins from explants that were infected with *S. aureus*. A total of 211 porcine-origin and 96 *S. aureus*-origin proteins were detected in the spent media.

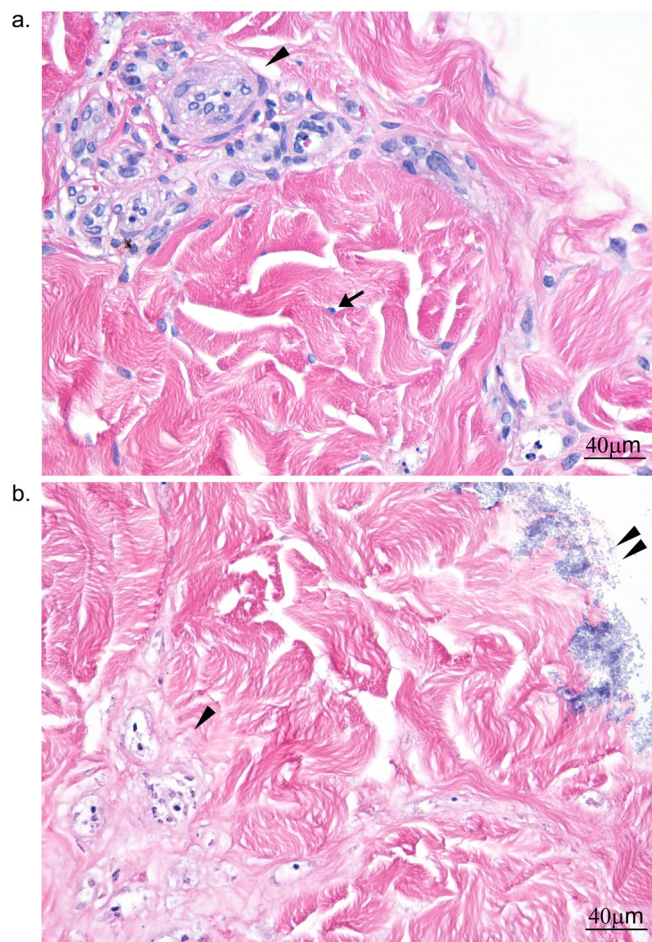


FIG 4 Histopathological changes in 4-day *S. aureus*-infected dermal explants. (a) Uninfected explant; (b) *S. aureus*-infected explant. The arrow in panel a points to a fibroblast nucleus; no equivalent nucleus was present with infected dermal explant. An arrowhead highlights cellular detail in blood vessels in panel a, whereas the basophilic stippling associated with the vascular bundles in panel b appears to be nuclear debris. The double arrowheads in panel b point to biofilm on the explant surface.

Of 211 porcine-origin proteins, 77 accumulated differentially (false discovery rate of $< 1\%$) (see Table S2 in the supplemental material). Programmed-cell-death protein 5 (PDCD5) accumulated to levels $> 1,000$ -fold higher in biofilm-infected explant media, consistent with explant cells undergoing apoptotic cell death. Accumulation of cytoskeletal proteins was also consistent with dermal cell apoptosis. Compared to uninfected explant media, higher quantities of cofilin-1 (90-fold), tropomyosin-1 (9-fold),

TABLE 1 Oxygen content in uninfected control explants^a

Explant	Oxygen level (mg/liter)	Effective diffusion coefficient (mm^2/s)
4 day	4.45 ± 1.17	$2.32 \times 10^{-3} \pm 8.31 \times 10^{-5}$
7 day	4.72 ± 1.12	$2.41 \times 10^{-4} \pm 5.96 \times 10^{-5}$
Dead ^b	$5.87 \pm 0.73^*$	$2.32 \times 10^{-3} \pm 8.31 \times 10^{-5}$

^a Values are means \pm SD, calculated from multiple data points from oxygen and diffusion coefficient depth profiles of three different explants. Comparisons of oxygen content and diffusion coefficients were carried out using ANOVA (*, $P < 0.05$).

^b Explants were killed in 2% paraformaldehyde.

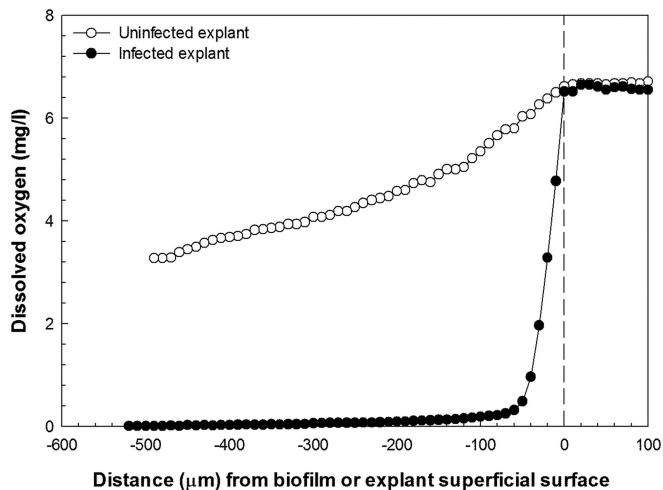


FIG 5 Typical dissolved-oxygen depth profiles of uninfected and *S. aureus*-infected explants. The biofilm or explant superficial surface begins at 0 μm , and the increasing negative distance refers to the increasing depth within the explant tissue. No measurable oxygen is present deep inside the tissue explants.

and LIM domain and actin-binding protein 1 (LIMA1) (337-fold) and smaller amounts of moesin (11-fold lower) were found in the spent media from infected explants. In addition, the antiapoptotic and proinflammatory cytokines macrophage migration inhibitory factor (58-fold) and interleukin 6 (9-fold) were also found in significantly larger amounts in the spent media from infected explants.

The *S. aureus*-origin proteins are given in Table S3 in the supplemental material, and the proteins that explain the *S. aureus* colonization and likely effect of biofilm on dermis explants are summarized in Table S4 in the supplemental material. In addition to known extracellular-matrix adhesins, the amino acid (proline and alanine)- and sugar (glucosamine)-metabolizing enzymes 1-pyrroline-5-carboxylate dehydrogenase (8,192-fold), alanine dehydrogenase (388-fold), and glucosamine-6-phosphate isomerase (1,024-fold) accumulated in high abundance in the spent medium from infected dermal tissue.

DISCUSSION

S. aureus is a common organism isolated from both chronic and acute wound infections, including surgical and combat-related wounds (38, 39). It has a number of virulence factors suitable for rapid growth and biofilm formation in wounds (40), and treatment is difficult, particularly for antibiotic-resistant strains. Despite its involvement in both acute and chronic wounds (41, 42), little is known about the role of *S. aureus* biofilms in wound pathogenesis. Three important factors affecting wound healing are oxygen availability, pH, and host cell survival. To examine these parameters in the context of biofilms, we first developed a porcine dermis explant model that resembles a dermal wound. Removal of epidermis exposes the extracellular matrix within the dermis, making it available for binding by *S. aureus*, which has a greater binding affinity for exposed extracellular matrix proteins (40, 43, 44). In addition, *S. aureus* is well adapted to survive and form biofilms in the extracellular matrix environment (see Table S3 in the supplemental material). We then demonstrated that biofilm-free explants remain largely viable for a week and that the partial

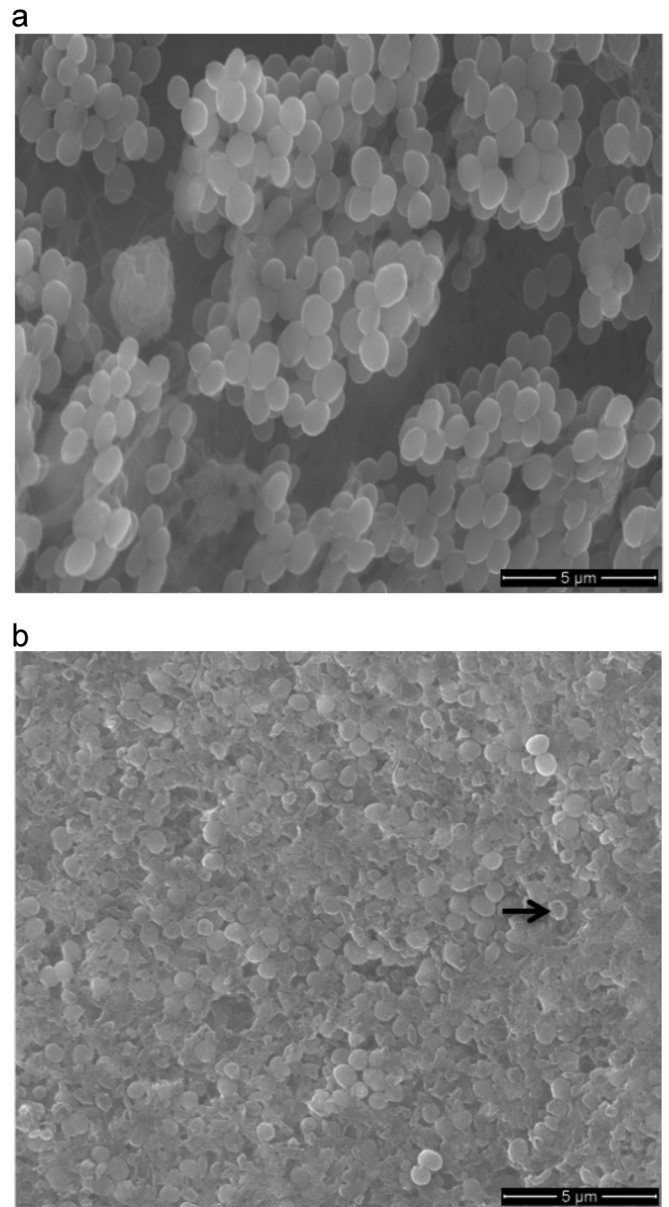


FIG 6 Environmental scanning electron microscopic images of 4-day (a) and 7-day (b) *S. aureus* biofilms on dermal explants. The cell clusters in 4-day biofilms appear healthy, with spatially separated cell clusters. The 7-day biofilms show a flat surface with lysed bacteria (arrow in panel b).

pressure of oxygen toward the deep surface is similar to that found in intact human dermis (45 to 65 mm Hg) (45). The explant model also permits the detection of soluble proteins that are released as a result of biofilm-explant interaction, which is impossible *in vivo*.

Oxygen is essential in wounds because it is required for ATP generation (46). The rate of diffusion of oxygen in tissue is directly proportional to the rate of diffusion of water molecules (30), where diffusion is a function of random movement of water molecules and is hindered by the compact fibrous nature of tissue, such as can be found in dermal explants. Compared to the diffusion coefficient of water in DMEM, the effective diffusion coefficient of water in explants decreased from 100% to about 29% at

TABLE 2 Differences between 4- and 7-day biofilm structural parameters

Structural parameter ^a	Value for ^b :	
	4-day biofilm	7-day biofilm
Biovolume (log μm^3)	7.31 \pm 0.06*	7.58 \pm 0.03*
Biofilm thickness (μm)	39.50 \pm 5.60*	86.0 \pm 2.50*
Biomass thickness (μm)	46.3 \pm 2.90*	88.2 \pm 3.20*
Biovolume-to-surface area ratio	1.7 \pm 0.07	1.9 \pm 0.08
Porosity	0.71 \pm 0.06	0.61 \pm 0.05
Diffusion distance (μm)	3.44 \pm 0.37*	5.14 \pm 0.31*

^a Biovolume is the total volume of the biofilm biomass. Biofilm thickness is the volume of the biofilm divided by the total surface area imaged. Biomass thickness is the volume of the biofilm divided by only that area of the substratum covered by the cells. Porosity is the ratio of total void volume to total biofilm volume. Diffusion distance is the average diffusion distance across all cell clusters in the biofilm.

^b Values are means \pm standard errors from 4 independent experiments. Values that differ significantly ($P < 0.05$) between 4- and 7-day biofilms are indicated with an asterisk. Comparisons were made using a *t* test (significance is based on the Bonferroni adjustment; $P < 0.008$).

the deep surface of the explants (Fig. 3). Dissolved-oxygen content also decreased from top to the bottom in both biofilm-free live and dead explants, but the overall oxygen content was always higher than 4.45 ± 1.17 mg/liter in viable explants (Table 1). Resting partial pressure of oxygen in most of the body tissues falls within the range we found in our explants (32), and therefore, the explant model reflects physiologically meaningful oxygen partial pressure. The oxygen content in 7-day explants was higher than that in the 4-day explants, consistent with reduced consumption of oxygen with decreasing explant viability, although the difference was not statistically significant in this case.

The relative effective diffusion coefficient increased significantly ($P < 0.0001$) in biofilm-infected explants (Fig. 3). This increase could primarily be due to cell death in the infected explants (Fig. 4). The degradation of extracellular matrix proteins by *S. aureus* could contribute to an increase in the average relative effective diffusion coefficient, but we did not observe any alterations in connective tissue based on histological examination, nor did we find any significant amounts of connective tissue matrix proteins in medium from *S. aureus*-infected dermal tissue (see Table S2 in the supplemental material). The increase in protein content in medium from *S. aureus*-infected explants was mainly of dermal cellular origin (see Table S2).

Despite the increase in relative effective diffusion coefficient, no measurable oxygen could be detected in the biofilm-infected explants. This indicated that changes in oxygen concentration in biofilm and dermal tissue are mainly caused by the oxygen consumption by biofilms rather than a mechanism that otherwise hindered diffusion. The *S. aureus* biofilms depleted oxygen throughout the full thickness (500 μm) of the explants (Fig. 5). This depletion was irrespective of changes in biofilm structure. We did not determine the minimum biomass of *S. aureus* biofilm that is needed to cause this degree of oxygen demand throughout an entire dermal explant.

Others have reported that hypoxic cells express more LDH, which results in a subsequent decrease in pH (33, 34). Although we observed more LDH activity in spent media from infected explants, the pH was higher in the biofilm, biofilm-infected explants, and the spent medium than in the uninfected explants. The reason for the higher LDH activity in infected-explant media

could partly be explained by substantial amounts of *S. aureus*-origin L-lactate dehydrogenase, Ldh2 (see Table S3 in the supplemental material). *S. aureus* can increase LDH expression under anaerobic conditions or in response to nitrosative stress (47, 48), but its role in the absence of lactate production (anaerobic metabolism) and nitrosative stress is not known. In severe cases of infection, *S. aureus* can cause lactic acidosis (49, 50). It is possible that in these cases, secretion of LDH by *S. aureus* could contribute to already existing lactic acidosis. Because it is an enzyme, small quantities of LDH can produce relatively large quantities of lactic acid.

S. aureus can increase biofilm and explant pH by consuming explant proteins and amino acids as a source of energy. Protein analysis of the spent medium identified some key *S. aureus*-origin enzymes involved in amino acid metabolism, including 1-pyrroline-5-carboxylate dehydrogenase (involved in proline catabolism [51]), glucosamine-6-phosphate isomerase (converts D-glucosa-

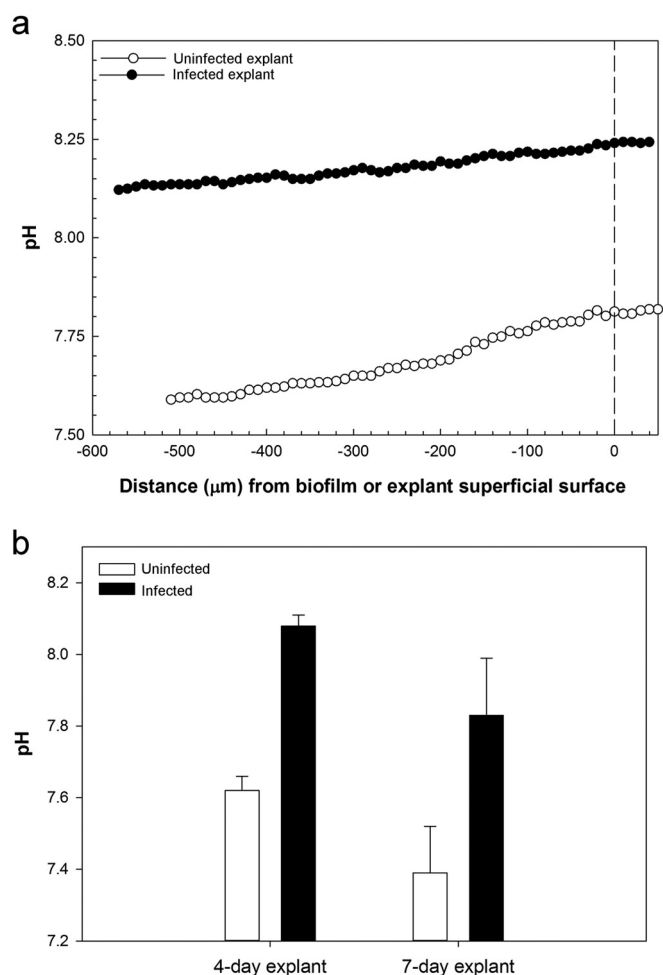


FIG 7 pH depth profiles in uninfected and *S. aureus*-infected explants. (a) Representative pH depth profile in uninfected and *S. aureus*-infected 4-day explants. (b) Histogram for mean pH \pm SD in uninfected and *S. aureus*-infected explants. Values were calculated by pooling measurements from every 10 μm from the top to the bottom of the explant from three different explants. The pH is significantly higher in *S. aureus*-infected than in uninfected explants (two-way ANOVA, Holm-Šidák method; $P < 0.002$; the model included explant age [4 day and 7 day] plus infection [uninfected and infected] plus interaction; the latter was not significant [$P = 0.97$]).

mine 6-phosphate to D-fructose 6-phosphate and ammonia [52]), and alanine dehydrogenase 1 (which catalyzes conversion of alanine and NAD⁺ to pyruvate, ammonium, and NADH [53]). The extracellular matrix, which is a binding site of *S. aureus* (40), contains abundant quantities of collagen, which is rich in proline and alanine, and hyaluronic acid; the latter contains D-glucosamine (54). The essential requirement of proline and alanine for *S. aureus* growth (55) indicates that *S. aureus* is well adapted for survival and multiplication in dermal tissue. The increase in pH could also be contributed to by arginine deiminase present in the arginine catabolic mobile element (ACME) of this strain. Arginine deiminase increases pH by producing ammonia and ATP from L-arginine (56). We also observed accumulation of alkaline shock protein in spent media from *S. aureus*-infected explants; alkaline shock protein expression is increased in *S. aureus* at elevated pH (57).

Wound pH can have a profound effect on enzyme function and on cells located at the site of injury. Acidic pH favors wound healing by enhancing fibroblast replication, early angiogenesis, and release of oxygen that is bound to hemoglobin (Bohr effect). This is in contrast with chronic wounds, where pH remains alkaline most of the time except during the epithelialization stage (11, 12, 20). *S. aureus* lives in acidic pH on skin surface (58, 59), but our model indicates that once established in wounds, *S. aureus* biofilms increase pH, and this is likely to be an impediment to wound healing.

Irrespective of the nature and type of the cells found in the dermis, *S. aureus* killed all the explant cells by day 4. The diffuse cell death combined with accumulation of large quantities of programmed cell death protein 5 (PDCD5) (more than 1,000-fold) and some apoptosis-associated cytoskeletal proteins indicated that apoptotic cell death may have occurred in the explants. PDCD5 (TFAR19) is a proapoptotic factor (60), and its expression is increased during apoptosis (61–63). It functions in apoptosis either through direct interaction with p53 (63) or via indirect acetylation of p53 (64). P53 plays a key role in cell cycle arrest and apoptosis (65, 66). PDCD5 could also act through impairment of chaperonin CCT (cytosolic chaperonin containing tailless complex polypeptide 1)-mediated beta-tubulin folding (67), thereby inducing apoptosis through an intrinsic pathway.

Previous reports indicate that *S. aureus* induces apoptosis in osteoblasts (through an extrinsic apoptotic pathway involving a tumor necrosis factor-related apoptosis-inducing ligand) and human umbilical vein and endothelial cell lines only when found intracellularly (68, 69). In our case, histological staining indicated no evidence that bacterial cells invaded the tissue explants, indicating that tissue cell death was driven by a mechanism that does not require intracellular invasion (but see below). *S. aureus* biofilms may instead induce apoptosis by depleting oxygen. In a hypoxic environment, hypoxia-inducible factor 1 (HIF-1), the key oxygen-sensitive transcription factor mediating cell response in hypoxia, induces high concentrations of proapoptotic genes and stabilizes p53 (70, 71). Under hypoxic conditions, bacterial biofilm products may potentiate the apoptotic effect.

Paradoxically, two proinflammatory and antiapoptotic cytokines (macrophage migration inhibitory factor [MIF] and interleukin 6 [IL-6] [72, 73]) accumulated to significantly larger amounts in biofilm-infected-explant media. Both MIF and IL-6 either inhibit or downregulate p53 and consequently should limit apoptosis (74–76). The expression of antiapoptotic cytokines lo-

cally at the site of infection, by either some or most cells (most likely during initial infection), may be the host's response to counter biofilm-induced apoptotic death, prolong cell survival, and aid in proliferation of infiltrating phagocytes and lymphocytes. In explants and actual wounds, *S. aureus* biofilms may induce both apoptosis and oncosis simultaneously, with apoptosis-induced cell death possibly predominating initially during the infection, when abundant ATP is available for energy-dependent apoptotic processes.

Recent work with a mouse model indicates that integrin-linked kinase (ILK) and Rac1 are requisite for *S. aureus* to invade mouse keratinocytes (77). *S. aureus* was able to penetrate skin explants (from the epidermis to dermis) of ILK-deficient mice with significantly greater efficiency, and consequently ILK is required for the epidermis to serve as an effective barrier against staphylococcal invasion. Once *S. aureus* breaches the epidermis, it likely penetrates the dermis through a combination of cell invasion and loss of tight junction integrity from protease activity. Our micrographs showed no evidence that *S. aureus* penetrated into the porcine dermis, and we hypothesize that cell death was driven by secreted toxins from the biofilm community. Nevertheless, our micrographs may have lacked sufficient analytic sensitivity to detect bacteria that might have been dispersed within the tissue. If they were present, their numbers were not great enough to detect with conventional staining techniques.

We studied the impact of *S. aureus* biofilms directly on dermal tissue because it contains multiple types of cells and closely resembles the natural wound surface. By using this model, we demonstrated that *S. aureus*-biofilms deplete oxygen and increase pH in tissues underlying the biofilms. During a host-pathogen interaction, bacterial pathogens may consume oxygen to the extent that host cells are deprived of this vital resource. Here we show that ~40- μ m-thick *S. aureus* biofilm growing on superficial surface of dermal tissue can increase the diffusion coefficient in underlying tissue while consuming oxygen so rapidly that nearly all oxygen is depleted throughout a full 500- μ m-thick dermal tissue explant. Hypoxia and alterations in pH have a global impact on gene expression, enzyme function, and cell behavior that adversely impact wound healing. We report that *S. aureus* biofilms kill dermal cells, most likely through an intrinsic pathway of apoptosis by inducing proapoptotic protein PDCD5. The intrinsic pathway of apoptosis can be activated by hypoxia, by bacterial products, or by a combination of both. In response to biofilm infection, we found substantial production of an important proinflammatory and antiapoptotic cytokine, MIF, which probably counters apoptosis induced by *S. aureus* biofilms for at least part of the infectious period. Our findings indicate that *S. aureus* biofilms can counter the host immune response far below the actual biofilm surface (500 μ m). We therefore propose that it is important to consider how bacterial biofilm-host interaction impacts the physical, biochemical, and immunological state of tissues underlying the actual surface of bacterial colonization.

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