Propionibacterium acnes and Staphylococcus lugdunensis Cause Pyogenic Osteomyelitis in an Intramedullary Nail Model in Rabbits

Abhay Deodas Gahukamble,a Andrew McDowell,b* Virginia Post,a Julian Salavarrieta Varela,a Edward Thomas James Rochford,a Robert Geoff Richards,a Sheila Patrick,b Thomas Fintan Moriartya

AO Research Institute Davos, Davos, Switzerland; Centre for Infection and Immunity, Queen’s University, Belfast, United Kingdom

Propionibacterium acnes and coagulase-negative staphylococci (CoNS) are opportunistic pathogens implicated in prosthetic joint and fracture fixation device-related infections. The purpose of this study was to determine whether P. acnes and the CoNS species Staphylococcus lugdunensis, isolated from an “aseptically failed” prosthetic hip joint and a united intramedullary nail-fixed tibial fracture, respectively, could cause osteomyelitis in an established implant-related osteomyelitis model in rabbits in the absence of wear debris from the implant material. The histological features of P. acnes infection in the in vivo rabbit model were consistent with localized pyogenic osteomyelitis, and a biofilm was present on all explanted intramedullary (IM) nails. The animals displayed no outward signs of infection, such as swelling, lameness, weight loss, or elevated white blood cell count. In contrast, infection with S. lugdunensis resulted in histological features consistent with both pyogenic osteomyelitis and septic arthritis, and all S. lugdunensis-infected animals displayed weight loss and an elevated white blood cell count despite biofilm detection in only two out of six rabbits. The differences in the histological and bacteriological profiles of the two species in this rabbit model of infection are reflective of their different clinical presentations: low-grade infection in the case of P. acnes and acute infection for S. lugdunensis. These results are especially important in light of the growing recognition of chronic P. acnes biofilm infections in prosthetic joint failure and nonunion of fracture fixations, which may be currently reported as “aseptic” failure.

Implant-related osteomyelitis is a serious complication of joint replacement and fracture fixation procedures. In the majority of cases, these infections are caused by bacterial species that are either permanent or transient members of the human microbiota. Of the different bacterial species present on the human skin, only a small number are frequently associated with implant-related osteomyelitis. One of the most common causative microorganisms is Staphylococcus aureus, which causes an acute infection characterized by fever, localized swelling, and osteolysis (1, 2). In contrast, the exact role played by other less virulent members of the skin microbiota in medical device-related infections, including the Gram-positive anaerobic-to-aerotolerant bacterium Propionibacterium acnes and coagulase-negative staphylococci (CoNS), has historically been less clear; these organisms were previously dismissed as contaminants when cultured from clinical specimens (3–5).

“Aseptic” implant failure is thought to result from periprosthetic osteolysis driven by wear debris arising from the implant materials (6). There is, however, a growing recognition that bacteria are an underestimated cause of subacute and chronic bone infections, aseptic loosening of implants, and arthrodesis after osteosynthesis (7–11). In particular, P. acnes is being increasingly recognized as a primary pathogen in relation to prosthetic shoulder implant infection (12), but it has also been linked to other medical device-related infections, including those associated with prosthetic hip joints and heart valves, as well as central nervous system shunts (13–15). In keeping with this pathogenic capacity, whole-genome sequencing has revealed the presence of multiple putative virulence determinants, including phase- and antigenically variable adhesins, as well as multiple cohomolysin CAMP factors (16). In contrast, Staphylococcus lugdunensis appears to be an unusually virulent CoNS species (17, 18) that has been linked to a wide range of acute infections, including those associated with prosthetic joints (19, 20), osteomyelitis (21, 22), septic arthritis (23), soft tissue infections (24), and infective endocarditis (25). Despite this, we still have a relatively limited understanding of the role and incidence of S. lugdunensis in human disease, which may lead to an understimation of its clinical relevance and, consequently, its importance in relation to implant-related osteomyelitis.

Although the pathogenic potentials of P. acnes and S. lugdunensis, especially in relation to medical device infections, are being increasingly recognized, there has been a limited number of preclinical experimental studies investigating the ability of P. acnes or S. lugdunensis to cause implant-related osteomyelitis. Therefore, the purpose of this study was to observe the effects of P. acnes and S. lugdunensis in establishing an established rabbit model (26) of implant-related osteomyelitis, in the absence of implant material wear debris, and characterize the resultant infections with respect to histological and microbiological outcomes.

MATERIALS AND METHODS

Ethics statement. The animal study was approved by the ethics committee of the Canton Grisons in Switzerland (approval no. 06/2008). The...
animal care and use protocol used for this study adheres to Swiss animal protection laws and its regulations. The bacterial isolates used in this study were obtained from adult patients who provided informed written consent to participate in a clinical study, which was approved by the Faculty of Medicine Research Ethics Committee at Queen’s University, Belfast, Northern Ireland. The P. acnes isolate was obtained as part of approval no. REF53/99, and the S. lugdunensis isolate was obtained as part of approval no. Ref86/01.

**Bacterial isolates.** The P. acnes strain LED2 was isolated after ultrasound treatment of a retrieved prosthetic hip joint, which was removed at Musgrave Park Hospital, Belfast, Northern Ireland, due to a supposed septic joint failure. A bacterial biofilm was also detected in the sonicate fluid by immunofluorescence microscopy (IFM) after labeling with a P. acnes-specific antibody, as previously described (11). The S. lugdunensis strain 010729 was isolated from an intramedullary (IM) nail-fixed tibial fracture at the Royal Victoria Hospital, Belfast, Northern Ireland. Although the fracture had united, the patient reported ankle pain. S. lugdunensis was also isolated from the nail plug and locking screws. Bacterial stock cultures were stored at 80°C in 20% (vol/vol) glycerol using the Protect (Microsynth AG, Balgach, Switzerland), visualized with UV light, and imaged using a GelDoc XR image analysis station (Bio-Rad, Reinach, Switzerland). The product sizes were estimated using BenchTop 100-bp and 1-kb DNA ladders (Promega) as molecular size markers. The PCR products were purified for sequencing using the PureLink quick gel extraction and PCR purification combo kit (Invitrogen), according to the manufacturer’s protocol.

**Nucleotide sequencing.** Automated sequencing was performed at Microsynth AG (Balghac, Switzerland) on an Applied Biosystems ABI 3730xl sequence analyzer 5.2 using the ABI BigDye system version 3.1. The sequences were compared with known sequences using BLAST (http://blast.ncbi.nlm.nih.gov).

**In vitro adhesion and biofilm formation.** In vitro adhesion of the bacteria to a range of orthopedic metals was measured using the preoperative contamination model of Rochford et al. (27). The metals used were electropolished stainless steel (SS), standard microrough titanium-aluminum-niobium (NS), and electropolished titanium-aluminum-niobium (NE). The adhesion studies were performed in phosphate-buffered saline (PBS) for S. lugdunensis and PBS supplemented with 0.05% (vol/vol) l-cysteine for P. acnes. Briefly, the bacterial suspension was adjusted to an optical density at 600 nm of 0.5 and diluted 10-fold to give a 1-liter suspension. The sequences were compared with known sequences using BLAST (http://blast.ncbi.nlm.nih.gov).

**In vivo animal model.** Custom-made IM nails composed of SS (ISO 5832-1), which were 2.5 mm in diameter and 85 mm in length, were used.

---

**TABLE 1 Primers used in this study**

<table>
<thead>
<tr>
<th>Target</th>
<th>Gene locus tag</th>
<th>Primer name 1</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>16S rRNA gene</td>
<td>fD1</td>
<td>CCGAATTTCGGACAACAGATTTTGATTCGAGCGCCATCTTGCCGTCAGCAG</td>
<td>60</td>
</tr>
<tr>
<td></td>
<td>rP2</td>
<td>CCCGGCGTAGCAGCTGGCTATTTGATTTTTGTTACGACTT</td>
<td>60</td>
</tr>
<tr>
<td>Von Willebrand factor</td>
<td>vwb1</td>
<td>TGGCGCGTAGATTTGGACGG</td>
<td>61</td>
</tr>
<tr>
<td>Fibronectin binding</td>
<td>fn</td>
<td>TGGCACACTCTTGCTCAGGTCAG</td>
<td>62</td>
</tr>
<tr>
<td></td>
<td>R1</td>
<td>TGGCACACTCTTGCTCAGGTCAG</td>
<td>62</td>
</tr>
<tr>
<td></td>
<td>fbl_check_F</td>
<td>CGTATTACCGCAATGAGCAACC</td>
<td>61</td>
</tr>
<tr>
<td></td>
<td>fbl_check_R</td>
<td>TCTCAGTGATGCCCAGATGC</td>
<td>61</td>
</tr>
<tr>
<td>Hemolytic peptides</td>
<td>slsh</td>
<td>GAGATTACTGGACAACAGACG</td>
<td>61</td>
</tr>
<tr>
<td>Tannase A</td>
<td>tanA</td>
<td>ACAGCACAACAAAACTTTACATCTCCA</td>
<td>61</td>
</tr>
<tr>
<td>Beta-hemolysin</td>
<td>SLGD_00006</td>
<td>TACCCCAATATACCGTGCTCCTTTT</td>
<td>61</td>
</tr>
<tr>
<td>Intercellular adhesion A</td>
<td>icaA</td>
<td>CACATCTTGCGTTGAG</td>
<td>61</td>
</tr>
</tbody>
</table>

**Molecular analyses.** Genomic DNA was prepared from single colonies using the Wizard Genomic DNA preparation kit (Promega, Düben-dorf, Switzerland), according to the manufacturer’s protocol. The primers used for PCR amplification are listed in Table 1. PCR amplification was carried out using an Eppendorf Mastercycler gradient thermal cycler (Vaudau-Eppendorf, Basel, Switzerland) in a total volume of 12.5 µl containing 10× PCR buffer (Invitrogen, Zug, Switzerland), 5 mM deoxyribonucleoside triphospate (dNTP) mix (Promega), 50 pmol of each primer (Microsynth, Balghac, Switzerland), 1 unit of Tag DNA polymerase (Invitrogen), and 10 to 50 ng template DNA. The products were separated by electrophoresis on 1% (wt/vol) agarose (Sigma-Aldrich) gels, stained with RedSafe nucleic acid staining solution (iNtRON Biotechnology, Basel, Switzerland), visualized with UV light, and imaged using a GelDoc XR image analysis station (Bio-Rad, Reinach, Switzerland). The product sizes were estimated using BenchTop 100-bp and 1-kb DNA ladders (Promega) as molecular size markers. The PCR products were purified for sequencing using the PureLink quick gel extraction and PCR purification combo kit (Invitrogen), according to the manufacturer’s protocol.
Prior to implantation, all 1M nails were washed twice in 70% (vol/vol) ethanol, followed by two periods of sonication in deionized water using an ultrasonic water bath operating at 40 Hz for 15 min. Each nail was then packaged individually and steam sterilized at 121°C for 20 min. Skeletally mature specific-pathogen-free female New Zealand white rabbits (Charles River Laboratories, Sulzfeld, Germany) were used in this study. Each rabbit was assessed by a veterinarian and determined to be healthy prior to inclusion in the study.

Prior to surgery, each rabbit was screened to determine whether the IM nail would be accommodated within the tibia. Under general anesthesia and using aseptic surgical techniques, the patellar ligament of the left tibia was divided and the anterior part of the tibia plateau exposed. The medullary cavity was opened with a 3.5-mm drill bit and evacuated with a 3.0-mm-diameter suction device. Using a catheter, the distal two-thirds of the medullary canal was inoculated with 50 µl of bacterial suspension in 25% (vol/vol) quarter-strength Ringer’s solution (QRS). The bacterial inocula were added to the implant site within 15 min of preparation. The implant was immediately inserted without hammering and the insertion site sealed with a water-soluble alkylene copolymer hemostasis material (Ostene; Ceremed, Inc. CA, USA). The patellar ligament and skin were closed in a routine manner. To minimize variability, the same surgeon operated on all rabbits.

The P. acnes inoculum for the animal study was prepared by subculturing a single colony of freshly grown LED2 from ABA into 50 ml MTL broth. The culture was incubated anaerobically for 16 h at 37°C. Immediately prior to surgery, an aliquot was taken and centrifuged at 2,500 rpm for 10 min, and the resulting pellet was resuspended in prereduced QRS containing the reducing agent t-cysteine (0.05% [vol/vol]). The inoculum density was adjusted to a target of 3 × 10^7 CFU/50 µl based on an optical density at 600 nm of 0.5. For the S. lugdunensis inoculum, one colony was taken from a young (<24 h) TSA culture and inoculated into 20 ml prewarmed TSB. The culture was incubated for 2 h in a shaking water bath at 37°C and 100 rpm. Prior to each surgery, an aliquot was taken and centrifuged at 2,500 rpm for 10 min, resuspended in QRS, and similarly adjusted to a target density of 600 nm of 0.6, approximately 3 × 10^7 CFU/50 µl.

A quantitative culture of each inoculum was performed immediately after preparation to determine the exact number of bacteria given to each animal. Serial 10-fold dilutions of the bacterial culture were performed in QRS and plated onto BA for 24 h (S. lugdunensis) or ABA plates for 14 days (P. acnes).

**Observation and euthanasia.** Upon completion of the surgical procedure, baseline radiographs were taken and the rabbits were returned to their individual hutches and monitored for 4 weeks. Each animal was monitored continuously throughout the observation period for signs of systemic infection. No antibiotics were administered to any animal during this study. Daily evaluations of attitude, appetite, and surgical incisions were carried out, while weight and body temperature were recorded weekly. Blood samples were collected preoperatively and weekly thereafter to determine the white blood cell (WBC) count (Vet abc; scil Animal care, Viernheim, Germany). Animal exclusion criteria were set at a weight loss of >10% of the initial body weight within 2 weeks, local infection with severe lameness, persistent swelling and discharge, or signs of systemic infection, such as fever, depression, and anorexia. After 28 days, each animal was humanely euthanized. Pilot surgeries were performed, whereby similar implants were placed, without any inoculation, in two animals. The weights, WBC counts, and histology images of these uninoculated animals were included for comparison.

**Postinfection bacteriological analysis.** All rabbits receiving an inoculum of either P. acnes or S. lugdunensis were processed for bacteriological analysis. After euthanasia, the skin was removed from the left leg and wiped down with 70% (vol/vol) ethanol. Swabs were taken from the tibial plateau at the point of insertion of the nail and streaked across BA plates. The tibia was then cut using an oscillating saw to expose the most distal end of the nail. The nail was removed through the distal end of the tibia and submerged in 6 ml of sterile QRS, gently agitated for 20 s to remove loosely adherent bacteria, and then transferred to a second bottle of sterile QRS. The nail was then vortexed for 30 s and subsequently placed in a sonicating water bath operating at a frequency of 40 kHz for 3 min before a final vortex for 30 s. The sonicates were immediately serially diluted in sterile QRS and 200-µl aliquots spread onto TSA, BA, and ABA plates for quantitative bacterial culture. The tibia was then cut just below the most proximal point previously occupied by the IM nail and the proximal portion fixed for histological processing as described below. A 2-cm section was also removed from the distal end of the tibia for histology. The remainder of the bone that previously surrounded the IM nail was then homogenized in 15 ml QRS with or without t-cysteine, as appropriate, using a Polytron PT 3100 homogenizer (Kinematica AG, Switzerland). Serially 10-fold diluted and duplicate 200-µl aliquots of the bone homogenate were added to TSA, BA, and ABA plates for incubation. All TSA and BA plates were incubated at 37°C aerobically, and the ABA plates incubated anaerobically, and the results were recorded at 24 h and reexamined after 48 h for S. lugdunensis- and 14 days for P. acnes-inoculated rabbits. In both groups, aerobic and anaerobic cultures was performed to ensure no secondary infection was present. The lower limit of detection of the bone samples was 75 CFU per bone fragment and 30 CFU for the IM nail. To be considered infected, either the bone or the IM nail was required to yield at least five colonies on both agar plates of the undiluted suspension. All bacterial growth was identified using the Remel RapID ANA II and the RapID Staph Plus system test kits, performed according to the manufacturer’s protocols (Remel, Switzerland).

**Histology.** The proximal and distal tibiae were fixed in 70% (vol/vol) methanol and subsequently decalcified with EDTA. Decalcified blocks were trimmed to include the entry point of the nail and the nail cavity and embedded in paraffin. Six-micrometer sections were cut and stained with hematoxylin and eosin for histomorphological scoring. Adjacent or nearby sections were stained with a modified Brown and Brenn stain to visualize the bacteria.

The regions of interest for histological scoring were the articular surface, entry site and tract of the nail, the medullary canal, cortical bone, and periosteum. Representative sections were graded by a histopathologist, who was blind to the rabbit treatments, for severity and characterization of the infection.

Electron microscopy was performed on representative sections to identify the location of the bacteria within the tissues. The sections were deparaffinized, sputter coated with gold palladium and then imaged using a Hitachi S-4700 scanning electron microscope (SEM). Images were taken in the secondary electron mode with an accelerating voltage of 5 kV.

**RESULTS**

**Strain characterization and adhesion properties.** P. acnes strain LED2 belongs to the type IB sequence type 5 (ST5) lineage (clonal complex 5 [CC5]), based on multilocus sequence typing (MLST) analysis of six core housekeeping genes and two putative virulence genes (16, 29, 30). It is hemolytic on horse and sheep ABA, has a demonstrable cohemolytic CAMP reaction, and expresses an abundance of the CAMP factor 1 protein as detected by SDS-PAGE and immunoblotting (31). In keeping with its identification as a type IB strain, it does not react with monoclonal antibodies specific for the phase-variable dermatan sulfate-binding adhesins present on the surface of type IA and IC isolates (32). S. lugdunensis 010729 is positive for the vwbI gene, which encodes the Von Willebrand factor, and fibronectin-binding proteins encoded by the fnm, fbl, and fbpA genes. Genes encoding beta-hemolysin and hemolytic peptides were also detected by PCR amplification. The intercellular adhesion gene icaA was also present.

In the bacterial adhesion assay, S. lugdunensis 010729 adhered in similar amounts to all of the metals tested (analysis of variance [ANOVA], P = 0.790), though the variation among the results...
was high (Fig. 1). The number of viable P. acnes cells retrieved from the materials was less than that of S. lugdunensis, although the amounts in the initial inocula were equivalent between groups. In addition, there were significantly fewer P. acnes adhering to NE than to SS and NS (ANOVA \(P = 0.050\)). The ability to form a biofilm in vitro was determined for LED2 and 010729 using the crystal violet staining method and classification scheme of Stepanović et al. (28). Under these conditions, S. lugdunensis produced a moderate biofilm within 24 h, with an optical density \(A_{600}\) and standard deviation (SD) of 0.12 \(\pm\) 0.01, while P. acnes produced a moderate biofilm with \(A_{600}\) and SD of 0.10 \(\pm\) 0.01 at day 7; however, only weak biofilm \((A_{600} < 0.05)\) formation was identified prior to day 7.

**Surgery and observation.** Postoperative radiographs showed good placement of the IM nail without any observable fractures. All rabbits tolerated the surgical procedure, and no animals were excluded from the study due to perioperative complications or postoperative sepsis. Postethical radiographs showed there was no migration of the implant, and no radiographic signs of severe osteolysis were observed (Fig. 2). Table 2 shows weight and WBC count values for P. acnes- and S. lugdunensis-infected animals and uninoculated rabbits throughout the study period. The uninoculated rabbits displayed a minor peak in WBC count and weight loss in the first postoperative week, both of which had returned to preoperative values prior to completion of the observation period, with overall weight gain observed by the completion of the study. The P. acnes-infected rabbits displayed minor weight loss in the first postoperative days, followed by full recovery and some weight gain, though less than that in the uninoculated animals. The WBC count also showed a minor spike in the first weeks, which was greater and more persistent than that observed in uninoculated animals. However, the WBC count returned to normal after the 4-week observation period. The S. lugdunensis-infected rabbits displayed, on average, a greater weight loss in the first 2 weeks, and by the end of the study, they had not gained any weight. The WBC count also showed a more pronounced and persistent elevation than that displayed by the P. acnes-infected animals, although the baseline WBC count was greater in the S. lugdunensis-infected group.

**Bacteriological outcomes.** The average intraoperative bacterial inoculum given to the rabbits that received P. acnes LED2 was \(1.67 \times 10^7\) CFU. Upon completion of the observation period, only 1/6 P. acnes-inoculated animals were culture positive from the joint swab. The bone cultures were positive for all six P. acnes-inoculated rabbits. Quantitative bacteriological analysis after sonication of the implant showed that all six P. acnes-inoculated animals were positive for biofilm on the IM nail. The TVC values of biofilm (sonication fluid and loosely adherent biofilm) and bone cultures (per gram) is shown in Fig. 3. On average, the total CFU counts cultured from the culture-positive bone samples and implants were \(2.6 \times 10^5\) CFU and \(3.3 \times 10^6\) CFU, respectively. In all cases, the microorganism cultured from the bone or the implants of the LED2-inoculated animals was confirmed to be P. acnes using Remel RapID diagnostic kits. IFM analysis with a monoclonal antibody (QUBPa3) (33) to P. acnes also confirmed the presence of the bacterium in sonicate samples (data not shown). No other bacterial species, based on colony morphology, were identified in any of the samples from these rabbits.

The average intraoperative bacterial inoculum given to the rabbits that received S. lugdunensis 010729 was \(2.49 \times 10^7\) CFU. At the completion of the 4-week observation period, 4/6 S. lugdunensis-inoculated rabbits were culture positive from the joint space swab. The bone cultures were also positive for the same rabbits and negative in the remaining two rabbits. No bacteria were isolated from animals recorded as culture negative. Quantitative bacteriological analysis after sonication of the implant showed that only 2/6 rabbits receiving the S. lugdunensis inoculum were positive for biofilm on the IM nail, both of whom were also culture positive from the bone. The TVCs of
the biofilm (sonication fluid and loosely adherent biofilm) and bone cultures (per gram) of each animal are shown in Fig. 4. On average, the total CFU counts cultured from the positive bone samples and implants were $5.1 \times 10^6$ CFU and $1.3 \times 10^7$ CFU, respectively. In all cases, the microorganism cultured from the bone or the implant of the S. lugdunensis-inoculated animals was confirmed to be S. lugdunensis using the Remel diagnostic kit. Sequence analysis of the 16S rRNA gene PCR fragment revealed 100% identity to the sequence of S. lugdunensis present in GenBank under accession no. N920143. As with P. acnes, IFM with a monoclonal antibody (QUBSe1) to P. acnes and S. lugdunensis also confirmed the presence of the bacterium in路过 samples (data not shown). No other bacterial species, based on colony morphology, were identified in any of the samples from these rabbits. All samples from the uninoculated rabbits were negative from the joint swab, bone samples, and sonicated samples (data not shown). No other bacterial species, staphylococci and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all

**Histology.** Characteristic features of osteomyelitis, such as micro- and macroabscesses, and the increased presence of neutrophils and other inflammatory cells were observed to various degrees in all six P. acnes-infected rabbits (Table 3). Focal inflammatory changes were seen in the subarticular region; however, the presence of fibrocartilage indicated the ulcerations were healing, and no subarticular macroabscesses were identified (Fig. 5). A higher-grade osteomyelitis was observed adjacent to the implant, where diffuse inflammation was seen, with some preservation of the microarchitecture (Fig. 6). These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, all specimens revealed the presence of bacteria, which primarily localized in small clusters located between adipocytes in the distal medullary cavity, which was confirmed by SEM imaging (Fig. 7).

The histological features of the S. lugdunensis infection were consistent with pyogenic osteomyelitis and septic arthritis, as summarized in Table 3. Focal inflammatory changes were seen in all animals with either micro- or macroabscesses in the subarticular region (Fig. 5 and 6). There was evidence of septic arthritis in all
six rabbits, with subarticular abscesses communicating with the adjoining joint in four of the animals. These pathologies were not observed in this model as a result of implantation of the IM alone (Fig. 5 and 6). After Brown and Brenn staining, bacteria were seen in all specimens examined (Fig. 7), including the two culture-negative animals. Bacteria were seen on the edges of the subarticular abscesses, while none or few were identified within the abscesses themselves. Small clusters were also located between adipocytes in the medullary cavity adjacent to activated erythropoietic and myelopoietic cells, indicative of early osteomyelitis. Some bacterial cells also appeared to have been phagocytosed by inflammatory cells in the bone marrow, although the viability of these bacteria was not determined (shown in Fig. 7). SEM analysis showed the scattered presence of S. lugdunensis throughout the tissue (Fig. 7), although an intracellular localization of S. lugdunensis could not be confirmed by SEM.

DISCUSSION
Historically, microorganisms, such as P. acnes and CoNS, have been considered contaminants from the skin rather than causative agents of osteomyelitis (3–5). Reports from the Patrick laboratory at Queen’s University Belfast in the late 1990s (11, 34) revealed the isolation and detection of these microbes in significant numbers from aseptically loosened failed implants by the use of ultrasound to dislodge adherent biofilms. These data suggested that these apparently low virulence commensal microorganisms might in fact be important pathogens capable of causing significant numbers of implant-related bone infections, leading to osteolysis and implant failure. Subsequent studies have supported these findings (10, 35–38), and further work at Queen’s University Belfast (QUB) also provided evidence of colonization of the bone adjacent to the implant (39). Despite this, the question of whether these bacteria are passive bystanders or drivers of prosthetic joint failure has been an ongoing subject of debate, with wear debris arising from implant materials still considered to be a major driver of implant failure (6).

While a large amount of research has been carried out to understand the pathogenic nature of S. aureus and Staphylococcus epidermidis, there is only limited information available for P. acnes...
and \textit{S. lugdunensis} and even fewer studies focusing on preclinical \textit{in vivo} models of infection (40–43). This lack of study also extends to the appropriateness of antimicrobial therapies for these microorganisms, the serological response to their infection, and the role of microbial virulence factors on the progression of infection. To address this issue, we established an experimental implant-related osteomyelitis model based on one described for \textit{S. aureus} (26), for the study of \textit{P. acnes}- and \textit{S. lugdunensis}-related infections. The model incorporates implanted IM nails, which, as with all indwelling biomaterials, provide enhanced colonization opportunities for professional pathogens and opportunistic pathogens alike. For our study, we used a strain of \textit{P. acnes} that was isolated from sonicate prepared from a failed prosthetic hip joint. This isolate had been previously characterized by MLST, which revealed that it belonged to the ST5 lineage within the type IB phylogenetic group (29). Isolates from this phylogroup are associated with healthy skin and are rarely recovered from acne vulgaris lesions. They have, however, been associated with soft tissue and medical device-related infections, although their exact clinical importance in these cases has remained unclear (16, 29). To date, a total of 13 distinct type IB STs have been described in the \textit{P. acnes} MLST database (http://pubmlst.org/pacnes/) based on the analysis of 69

### TABLE 3

Summary of the histological findings associated with \textit{P. acnes} LED2 and \textit{S. lugdunensis} 010729 infection

<table>
<thead>
<tr>
<th>Location and histological feature</th>
<th>\textit{S. lugdunensis} 010729</th>
<th>\textit{P. acnes} LED2</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Grade or score</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Subarticular</td>
<td>Medullary</td>
</tr>
<tr>
<td>Presence of ulceration</td>
<td>Yes/no</td>
<td>4</td>
</tr>
<tr>
<td>Presence of fibrocartilage, indicating healed ulceration</td>
<td>Yes/no</td>
<td>1</td>
</tr>
<tr>
<td>Presence of inflammatory cells or bacteria in the ligament, indicating septic arthritis</td>
<td>Yes/no</td>
<td>6</td>
</tr>
<tr>
<td>Presence of ulceration</td>
<td>Yes/no</td>
<td>4</td>
</tr>
<tr>
<td>Presence of fibrocartilage, indicating healed ulceration</td>
<td>Yes/no</td>
<td>1</td>
</tr>
<tr>
<td>Presence of inflammatory cells or bacteria in the ligament, indicating septic arthritis</td>
<td>Yes/no</td>
<td>6</td>
</tr>
<tr>
<td>Subarticular region, entry tract, and medullary cavity</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Diffuse increase in the cellularity of the marrow with near normal preservation of the marrow architecture (fat cells and precursors)</td>
<td>Grade 1 myelitis</td>
<td>1</td>
</tr>
<tr>
<td>Presence of ( \leq 5 ) microabscesses with or without a diffuse increase in cellularity similar to grade 1</td>
<td>Grade 2 myelitis</td>
<td>2</td>
</tr>
<tr>
<td>Presence of ( &gt;5 ) microabscesses with or without a diffuse increase in cellularity similar to grade 1</td>
<td>Grade 3 myelitis</td>
<td>2</td>
</tr>
<tr>
<td>Presence of macroabscesses defined as collection of neutrophils with a fibrous capsule OR a localized gross increase in cellularity</td>
<td>Grade 4 myelitis</td>
<td>1</td>
</tr>
<tr>
<td>Diffuse gross increase in neutrophils and inflammatory cells in the entire medullary canal inclusive of the periphery and center</td>
<td>Grade 5 myelitis</td>
<td>0</td>
</tr>
<tr>
<td>Presence of necrotic bone or sequestrum</td>
<td>Yes/no</td>
<td>1</td>
</tr>
<tr>
<td>Presence of fibrosis</td>
<td>Yes/no</td>
<td>6</td>
</tr>
<tr>
<td>Cortical bone</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Increase in the size of the lacunae with neutrophils</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
<tr>
<td>Presence of hyaline degeneration of vessels</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
<tr>
<td>Presence of vascular neutrophilia</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
<tr>
<td>Presence of empty lacunae</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
<tr>
<td>Fibrosis</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
<tr>
<td>Presence of necrotic bone or sequestrum</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
<tr>
<td>Inflammatory cells under the periosseum/subperiosteal abscess</td>
<td>Yes/no</td>
<td>NA</td>
</tr>
</tbody>
</table>

\( ^a \) NA, not applicable.

**FIG 5** Micrographs illustrating histological analysis of the point of entry of the nail and subarticular region for uninfected (A), \textit{P. acnes} LED2-infected (B), and \textit{S. lugdunensis} 010729-infected (C) rabbits. (A) The entry point has healed completely and no subarticular abscesses are observed. (B) Healing of the point of entry of the nail is also seen, and no signs of subarticular abscess formation were observed. Increased numbers of inflammatory cells were present deep in the medullary cavity adjacent to the implant site (arrow). (C) Macroabscesses were observed just below the articular surface (arrows) (grade 4). Hematoxylin and eosin staining was used.
type IB isolates, of which 11 belong to a single CC, with ST5 as the founder genotype (CC5) (bootstrap value, 99%), while two isolates are singletons (ST12 and ST51). As expected for a founding genotype, the ST5 clonal lineage is highly prevalent in the human population and is globally disseminated. *P. acnes* contains an expanded family of five CAMP factor homologues that appear to have arisen primarily as a result of horizontal gene transfer (HGT), as judged by previously described colocalization and sequence similarity criteria (16). Previous studies with five type IB *P. acnes* isolates (all subsequently identified as belonging to the ST5 lineage by MLST) demonstrated abundant production of the CAMP factor 1 protein in a comparison with strains from the type IA division; the secretion of large quantities of CAMP 1 was also observed with isolates from the large type II clade (29, 31). Currently, the exact role played by CAMP 1 and the other CAMP factor homologues is unclear, but they may play an important role in the survival of *P. acnes* within the human host as well as contributing to a pathogenic lifestyle (16). Recently, an MLST scheme and database for *S. lugdunensis* were described based on the analysis of seven core housekeeping genes (44). A clonal population structure with limited sequence diversity was revealed, and isolates recovered from hematogenic infections (blood or osteoarticular isolates) or from skin and soft tissue infections were not found to cluster in separate lineages. Currently, we do not have MLST data for *S. lugdunensis* 010729, but we are now analyzing the strain by whole-genome sequencing, which will facilitate the immediate assignment of an ST and CC when data for each of the MLST loci are complete.

![FIG 6 Micrographs illustrating histological analysis of the medullary region of uninfected (A), *P. acnes* LED2-infected (B), and *S. lugdunensis* 010729-infected (C) rabbits. (A) The medullary region displays normal physiology with abundant univacuolar adipocytes and the lack of cellular infiltration. (B) A diffuse marked increase in cellularity indicative of grade 3 osteomyelitis is seen (white arrow). (C) Focal inflammatory changes, such as microabscesses, are seen in the medullary region, with a marked decrease in adipocytes (white arrow). Hematoxylin and eosin staining was used.](http://jcm.asm.org/)

![FIG 7 Micrograph illustrating the localization of bacteria within infected animals. *S. lugdunensis* was primarily associated with abscesses (A), and an intracellular localization of *S. lugdunensis* was also observed (modified Brown and Brenn staining). (B) Scanning electron micrograph of bacterial cells in medullary area (arrows). (C) *P. acnes* cells were arranged in clusters (arrow) adjacent to adipocytes in the medullary cavity. (D) Scanning electron micrograph of bacterial cells in the medullary area. Note the pleomorphic rod morphology characteristic of *P. acnes* (arrows).](http://jcm.asm.org/)

![FIG 6 Micrographs illustrating histological analysis of the medullary region of uninfected (A), *P. acnes* LED2-infected (B), and *S. lugdunensis* 010729-infected (C) rabbits. (A) The medullary region displays normal physiology with abundant univacuolar adipocytes and the lack of cellular infiltration. (B) A diffuse marked increase in cellularity indicative of grade 3 osteomyelitis is seen (white arrow). (C) Focal inflammatory changes, such as microabscesses, are seen in the medullary region, with a marked decrease in adipocytes (white arrow). Hematoxylin and eosin staining was used.](http://jcm.asm.org/)
The experimental model used here requires the direct artificial inoculation of bacteria into the surgical field, which it must be noted will exceed the number of bacteria likely to contaminate a surgical wound during a primary elective procedure. However, if a failed prosthetic joint with a misdiagnosis of aseptic loosening is removed and a new sterile device is placed in the underlying infected site, it is very possible that higher numbers of bacteria will be present, particularly if the adjacent bone has been colonized. The bacterial inoculum we used (3 x 10^7 CFU) is well within the range of inocula previously applied in other experimental rabbit models of staphylococcal implant-related osteomyelitis (for example, 3 x 10^6 to 2 x 10^6 CFU [26, 45, 46]). Upon completion of the study, the total CFU counts recovered from both P. acnes- and S. lugdunensis-infected rabbits were quite varied, although the majority of culture-positive samples were between 1 x 10^6 and 1 x 10^7 CFU for P. acnes and 1 x 10^6 and 1 x 10^7 CFU for S. lugdunensis. The total numbers of bacteria quantitatively cultured from this study are similar to the numbers of S. aureus cultured from the previous model on which our study is based (26). The lower recovery of S. lugdunensis, in particular the lack of a consistent biofilm, is an interesting finding further discussed below.

The general histopathological features of osteomyelitis observed in this model included the presence of diffuse, focal, and multifocal inflammation characterized by infiltration of the medullary canal by neutrophils and the presence of subarticular and peri-implant abscesses with surrounding fibrosis. A histopathological scoring system, based upon similar studies using S. aureus (47–49), was developed to specifically focus upon the features of implant-related osteomyelitis as it occurs after intramedullary nailing. This scoring system was then used to determine whether there were differences in the characteristics of any infection caused by these two species of bacteria. In animals that underwent surgery but were un inoculated, the entry point of the nail healed uneventfully, and the medullary area displayed a physiological appearance, lacking in increased cellularity or significant fibrosis. A comparatively severe histopathology, characterized by macroabscess formation and septic arthritis, was observed after infection with S. lugdunensis. Although S. aureus-infected animals were not compared directly in this implant model, the histopathology of the S. lugdunensis-inoculated animals was comparable to that seen in a similar animal model using S. aureus (48). This is consistent with the association of S. lugdunensis with a clinical course more similar in severity to a typical S. aureus infection than to that caused by other CoNS (17). A complete-genome sequence of S. lugdunensis did not reveal the well-recognized virulence determinants that enable S. aureus to cause acute infection, but different strains of S. lugdunensis may carry other putative virulence determinant genes gained as a result of HGT (17, 50). Interestingly, the S. lugdunensis infection was not significantly associated with biofilm formation, as a biofilm was detected by culture on the IM nail of only 2/6 animals. This is despite in vitro biofilm formation by this S. lugdunensis isolate on a range of implant materials. However, the S. lugdunensis bacterium was observed in all histological sections after 28 days in vivo. The histological analysis revealed that S. lugdunensis was predominantly found adjacent to abscesses and in many cases appeared intracellularly within phagocytic cells in the abscess tissue. The viability of these intracellular bacteria is not known, but intracellular survival is a feature associated with S. aureus and S. epidermidis and is entirely in keeping with the intracellular survival of other CoNS (51). The intracellular location of S. lugdunensis is a potential explanation for the culture-negative animals observed in this study. Non-culture-based detection of staphylococci has also previously been described for culture-negative retrieved failed prostheses and biopsy specimens from human patients with long-bone nonunion of fractures (52). The culture-negative animals may also be due to an unavoidable sampling error, leading to a bacterial presence in the bone sample retained for histology yet absent from the biopsy specimens retained for bacterial culture. What is clear, however, is that the histopathology of the culture-negative S. lugdunensis-inoculated animals indicates an active infection in these animals. Further study is required to definitively characterize the natural progression of the infection in this model and determine whether this culture-negative finding indicates a subsidence of the infection.

In the in vivo model, P. acnes causes an infection characterized by biofilm formation and localized inflammation adjacent to the biofilm and implant. According to the histological scoring, P. acnes did not cause septic arthritis, and the entry wound from the insertion of the IM nail was found to be healing with fibrocortical-laginous tissue formation. The histological scoring of the medullary area adjacent to the implant and the biofilm was greater than that in the subarticular region, and even greater in severity than the S. lugdunensis data. These pathologies are not observed with implantation of the IM alone (Fig. 5 and 6). The P. acnes microcolonies were observed primarily adjacent to the adipocytes within the medullary cavity, some of which also showed regions with high-grade osteomyelitis. SEM analysis confirmed microcolony formation (Fig. 7) in the intercellular regions in the bone marrow, and there was no evidence of intracellular localization of P. acnes by SEM or light microscopy. The clinical signs, such as weight loss and white blood cell count, were less severe in the P. acnes-infected animals than in those infected with S. lugdunensis. Uninoculated animals gained more weight than P. acnes-infected animals, which supports the view that P. acnes causes chronic low-grade infections characterized by subtle clinical signs and symptoms and a significantly different histopathology than that from staphylococcal osteomyelitis.

Overall, both the in vitro and in vivo evaluations indicated that P. acnes differs in many respects to S. lugdunensis. Our in vitro study revealed that P. acnes adhered relatively poorly to a range of implant materials compared to S. lugdunensis. In vitro formation of biofilm and adhesion to orthopedic materials have been shown for P. acnes (31, 53–56), although a correlation between in vitro and in vivo studies is lacking. Our isolate, LED2, displayed weak biofilm formation in vitro, although this is clearly not representative of the clinical situation, or even for our experimental infection. Our in vivo data revealed that P. acnes formed a biofilm in all the animals examined; in contrast, S. lugdunensis appeared to have reduced ability to do so under these conditions. The relationship between biofilm formation and osteomyelitis therefore requires further investigation, particularly in the case of S. lugdunensis. Of high importance is the observation that more severe medullary osteomyelitis was observed in P. acnes-infected animals compared to those infected with S. lugdunensis. The most severe grade of intramedullary osteomyelitis, grade 5, was observed in three rabbits infected with P. acnes but in none infected with S. lugdunensis. The model used is a nonloaded IM nail without any interlocking bolts. As such, there is no risk of any wear-induced particles or fretting corrosion, and no such particles were observed in any histological section. Furthermore, osteomyelitis was not observed...
where the IM was implanted in the absence of either *S. lugdunensis* or *P. acnes*. This indicates that *P. acnes* can cause osteomyelitis in the absence of implant material wear debris and in the absence of overt clinical signs of infection. With regard to *S. lugdunensis*, the model confirms that this is a potentially pathogenic coagulase-negative staphylococcal species and that biofilm formation is not a prerequisite for infection.

These results have important implications in relation to distinguishing between true aseptic joint loosening driven solely by, for example, wear particle-mediated inflammation, and chronic infection-driven loosening, which may also be combined with wear debris involvement. These findings may also have wider implications in relation to the involvement of such bacteria in other conditions, such as dental infection (57) and synovitis-acnes-pustulosis-hyperostosis-osteitis (SAPHO) syndrome (58, 59). It would be interesting to follow the progression of both infections histologically over a longer time period and as it becomes more chronic in nature. It would also be valuable to investigate the effect of these pathogens on pseudarthrosis or delayed healing; however, currently, there is no standardized rabbit model of implant-related osteomyelitis with a fracture. Nevertheless, the model showed that the progression of implant-related osteomyelitis follows a bacterial species-specific course and highlights the potential of *P. acnes* and *S. lugdunensis* to cause significant implant-related osteomyelitis.

Based on the study, we conclude that both *P. acnes* and *S. lugdunensis* are pathogens capable of producing osteomyelitis in a rabbit model but exhibit different pathologies. The infection caused by *S. lugdunensis* was in keeping with a classically virulent microbe, with macroabscess formation and failure to heal entry wounds, but only a maximum grade 4 intramedullary osteomyelitis was observed in a single rabbit, with the remaining rabbits experiencing lower-grade osteomyelitis. Interestingly, the *S. lugdunensis* infection was not always characterized by biofilm formation, even when bacteria were present in the adjacent bone. This result may have important clinical implications with regard to the location and extent of tissue debridement required for implant revision/removal, which commonly occurs on implant removal only. In contrast, the infection caused by *P. acnes* was characterized by healing of the entry wound and low morbidity for the animals; however, there was extensive biofilm formation and, most significantly, evidence of grade 5 intramedullary osteomyelitis. This occurred in the absence of implant material wear debris. This experimental study therefore provides strong evidence for a potential key role of *P. acnes* in prosthetic joint infection and fracture nonunion in the absence of signs of classical infection and patient morbidity. *P. acnes* should therefore no longer be dismissed as an insignificant pathogen in the setting of failed retrieved implants; clinical diagnostic practice should be tailored to enable the efficient detection of *P. acnes*. Without this, there is the risk of an incorrect diagnosis of aseptic loosening, and subsequent patient treatment may be misinformed.

**ACKNOWLEDGMENTS**

This work was supported by a grant from AOTrauma.

We thank Iris Keller, Nora Goudouzian, Dirk Nehrbas, Ludovic Boure, Stephan Zetter, and Stephanie Neubert (AO Research Institute, Davos) for histological, surgical, and technical assistance. We thank the orthopedic surgical staff at Musgrave Park Hospital and the Royal Victoria Hospital in Belfast for facilitating the provision of implants to enable the original isolation of the strains and, in particular, orthopedic surgical trainee Michael McMullan for obtaining the retrieved intramedullary nail and the initial isolation of *S. lugdunensis*.

**REFERENCES**


