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Early implant-associated osteomyelitis results in a peri-implanted bacterial reservoir

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Implant-associated osteomyelitis (IAO) is among the most severe orthopedic conditions (1). The absolute number of IAO cases is increasing, owing to the growing number of patients with bone implants (2). In the United States of America, the infection rate is 5–15% in fracture fixation devices and 0.3–5% in joint prosthesis (3, 4). Treatment of IAO can include surgical debridement, removal of implants and long-lasting antimicrobial therapy, and calls for a multidisciplinary approach (1). Nevertheless, treatment failure is common. Despite removal of the infected device and extensive debridement, there is a high risk of re-infection and prolonged use of postoperative antibiotics (5). In two different retrospective studies, treatment failure rates of IAO have been estimated to 41.8 and 58.2 percent, respectively (6, 7). An explanation for the infections and re-infections of IAO has been suggested to be bacterial survival in the peri-implanted bone tissue (8).

Insertion of orthopedic implants is an equipment requiring process that involves drilling and often the use of bone cements. A by-product of these procedures is the generation of heat resulting in osteonecrosis (9). The necrotic osteocytes lose their inhibitory effects on osteoclasts leading to increased osteoclast activity and thereby bone resorption (10). Aside from resorption of dead bone, the process of osseointegration of an implant is especially dependent on the ingrowth of osteoblasts and mesenchymal stem cells (11). Along with osteonecrosis, bone resorption and osseointegration, a foreign body response also occurs around the implant (12). All these cellular changes make the peri-implanted bone tissue a perfect locus resistentiae minoris for bacterial infection (13).
The aim of this study was to answer the following question: how deep is a bacterial infection going in the peri-implanted bone tissue, in a case of IAO, within a week? The question was addressed in a porcine model of IAO in which evaluation of the entire infected bone is possible after a fixed period of time, compared to surgical biopsies from humans. Furthermore, it is highly relevant to use pigs for modeling of infectious diseases in humans, like IAO, as the porcine immune system shows a higher degree of similarity when compared to rodents and rabbits (14).

MATERIAL AND METHODS

Study design

The study is a descriptive study based on mainly microscopic observations in bone tissue and on orthopedic implants, obtained from a porcine model of IAO. The model is based on tibial insertion of a small steel implant combined with inoculation of Staphylococcus aureus bacteria or sterile saline. Twelve pigs of 30 kg body weight, obtained from a specific pathogen-free (SPF) herd, were divided into three groups based on their time of killing (Table 1). The Danish Animal Experimental Inspectorate approved the protocol (license No. 2013/15-2934-00946).

Experimental surgery and inoculum

Animals were anesthetized (15) and a tibial implant was inserted (K-wire 2 × 20 mm) 1 cm below the growth plate of the right tibia. The procedure was recently described (16). The inoculating S. aureus strain was a pathogenic porcine strain, previously used in porcine models of osteomyelitis (17). The strain was prepared as described by Johansen et al. (18) and diluted with 0.9% sterile isotonic saline to obtain an inoculation dose of 10⁶ colony forming units (CFU)/kg BW in a final volume of 10 μL.

Postoperative care of pigs

The pigs were daily monitored throughout the experiment by skilled personal. A body temperature above 41 °C, impaired ability to stand and anorexia were set as human endpoints. The pigs received intramuscular injections (0.1 mg/kg BW) of buprenorphine (Temgesic 0.3 mg/mL, Schering-Plough, Heist-op-den-Berg, Belgium) every 6–8 h. The pigs did not receive local or systemic antibiotic treatment, which is applied to human patients under therapy, as this would have hampered the focus of the study, that is, the ability to spread within the bone of the inflammation and infection.

Pathology

Following killing after 2, 4 and 6 days, the implant was removed from the implant cavity using a sterile lancet and collected for scanning electron microscopy (SEM) and peptide nucleic acid fluorescence in situ hybridization (PNA FISH). From all pigs, the right tibia was dissected free and decalcified. However, in group C animals, the right tibial bone was sagittal sectioned through the implant cavity before decalcification. Following decalcification, the proximal end of right tibial bone, containing the implant cavity, was cut into five sagittal pieces 3–4 mm each. Afterward, the bone pieces were processed routinely and embedded in paraffin wax. Sections (4–5 μm) were stained with hematoxylin and cosin (HE) and special stained with phosphotungstic acid hematoxylin (PATH) and Masson’s trichrome for demonstration of fibrin and collagen, respectively. Bone tissue with

| Table 1. Summary of design and results in a porcine model of implant-associated osteomyelitis |
|---|---|---|---|---|---|
| Animals | Inoculum | Time of killing PI | Microbiology of swab from implant cavity | IHC detection of S. aureus in PIBA | Detection of S. aureus on implant surface by PNA FISH | Detection of biofilm on implant surface by SEM | Score of neutrophils in PIBA<sup>1.<sup>3</sup></sup> |
| Group | CFU/Kg BW af S. aureus (Spa-type) | (Days) | (Spa-type) | | | | |
| A | Saline | 2 | S. aureus (not typed) | No | – | – | 3+ |
| | Saline | 2 | S. aureus (t1333) | No | No | – | 2+ |
| | 10⁴ (t1333) | 2 | S. aureus (t1333) | Yes | No | – | 3+ |
| | 10⁵ (t1333) | 2 | S. aureus (t1333) | Yes | – | Yes | 3+ |
| B | Saline | 4 | S. aureus (not typed) | No | – | – | 2+ |
| | Saline | 4 | S. aureus (t1333) | Yes | No | – | 3+ |
| | 10⁴ (t1333) | 4 | S. aureus (t1333) | Yes | No | – | 3+ |
| | 10⁵ (t1333) | 4 | S. aureus (t1333) | Yes | – | Yes | 3+ |
| C | Saline | 6 | S. aureus (not typed) | No | – | – | 2+ |
| | Saline | 6 | S. aureus (t1333) | No | – | Yes | 2+ |
| | 10⁴ (t1333) | 6 | S. aureus (t1333) | Yes | – | Yes | 3+ |
| | 10⁵ (t1333) | 6 | S. aureus (t1333) | Yes | – | Yes | 2+ |

BW, body weight; PI, postinfection; IHC, immunohistochemistry; PIBA, peri-implanted pathological bone area; PNA FISH, peptide nucleic acid in situ fluorescence hybridization; SEM, scanning electron microscopy.
pathological changes, around the implant cavity, was defined as the peri-implanted pathological bone area (PIBA). The largest size of PIBA was measured perpendicularly to the implant cavity on the most representative section. On the same section, the scoring system developed by Pandey el al. (19) was used to define whether an infection occurred. Pandey found that the presence of 2+ or more (more than one neutrophil granulocyte per high power field (400) on average after examination of at least 10 high power fields) in periprosthetic tissue, distinguishing between septic and aseptic loosening of prosthesis and bone implants in humans. The scores were as follows: 0 = absent; 1+ less than 1 cell on average per high power field; 2+ = 1–5 cells on average per high power fields; 3+ = >5 cells on average per high power fields.

**Microbiology**

Cotton swabs were taken from the implant cavity after removal of the implant. Swabs were processed and characterized as previously described (20), and selected bacterial isolates were *Spa*-typed (21).

**Immunohistochemistry of bone tissue**

Tissue sections of 4 μm were prepared and processed for indirect *in situ* identification of *S. aureus* with immunohistochemistry. Primary *S. aureus*-specific antibodies (ab37644; Abcam, Cambridge, UK, diluted 1:1000 in 5% swine serum) were used (18). The size of the largest observed *S. aureus*-positive aggregates and their distance to the implant cavity were estimated on each section, by measuring the length directly on the immunohistochemistry (IHC) sections.

**Scanning electron microscopy and peptide nucleic acid fluorescence *in situ* hybridization of implants**

Selected implants were examined with scanning electron microscopy (SEM) and peptide nucleic acid fluorescence *in situ* hybridization (PNA FISH) (Table 1). The implants were placed in 2.5% glutaraldehyde or formalin for SEM and PNA FISH, respectively. The samples for SEM were rinsed three times in 0.15 M sodium phosphate buffer (pH 7.4); specimens were postfixed in 1% OsO4 in 0.12 M sodium cacodylate buffer (pH 7.4); specimens were postfixed in 1% OsO4 in 0.12 M sodium cacodylate buffer (pH 7.4) for 2 h. Following a rinse in distilled water, the specimens were dehydrated to 100% ethanol and critical point dried (Balzers CPD 030 instrument, Leica Microsystems, Wetzlar, Germany) using CO2. The specimens were subsequently mounted on stubs, using colloidal coal as an adhesive, and sputter coated with gold (Polaron SEM E5000 coating unit). Specimens were examined with a Philips FEI30 scanning electron microscope operated at an accelerating voltage of 2 kV. The samples for PNA FISH were carefully rinsed in sterile saline for 5 min. Samples were placed in a dish (depth 1 mm, diameter 10 mm) and covered by 100 μL PNA FISH-specific *S. aureus* probe (Advandx, Woburn, Massachusetts, USA). Samples were incubated at 55 °C for 90 min. Subsequently, dishes were submerged for 30 min in 55 °C warm wash buffer (4 mL 60× wash buffer (Advandx) to 240 mL milQ water). Afterward, the samples were air-dried in the dark followed by covering by 3 mM DAPI solution (Life Technologies, Carlsbad, California, USA) for 15 min at room temperature. Excess DAPI was removed by gently rinsing with PBS (the Substrate Department at the Panum Institute, Denmark). All observations were performed using a Zeiss LSM 710 confocal laser scanning microscope (Oberkochen, Germany).

**Statistics**

An unpaired t-test was used to analyze PIBA measurements between control and infected animals (GraphPad Software inc, version 7, LaJolla, California, USA).

**RESULTS**

**Clinical observations**

All animals were healthy when entering the study, but became lame on the operated leg after surgery and up until killing. The degree of lameness was low, as the animals were able to use the leg and walk around freely. One of the infected Group B animals had intermittent elevated body temperature. All animals ate and drank normally before and during the experiment.

**Macroscopic pathology**

Thick purulent material was seen within the implant cavity in the infected animals following 4 and 6 days. In all control pigs, and pigs infected for 2 days, serohemorrhagic fluid was present. Following sagittal section of the inoculated tibial bone from the two infected Group C animals, signs of osteomyelitis were seen around the implants as purulent and sequestered trabecular bone tissue (Fig. 1).

**Histopathology**

The neutrophil granulocyte score counted inside PIBA and the size of PIBA is presented in Table 1 and Fig. 2A, respectively. The size of PIBA showed no marked difference between infected and control animals after 2 days from surgery with a mean difference of 0.7 mm (Fig. 2A). However, in Groups B and C, the mean differences between infected and control animals were 4.3 mm and 4.4 mm, respectively (Fig. 2A). The p-value of the difference in PIBA size between control (n = 6) and infected animals (n = 6) was 0.0014 (Fig. 2B). The largest registered PIBA value for day 4 (Group B) and day 6 (Group C) was 6.0 mm and 6.21 mm, respectively (Fig. 2A).

The following description of pathomorphological changes is correlated with the trabecular tissue only. In Group A, PIBA was a mix of erythrocytes, bone marrow leukocytes, necrotic trabecular tissue...
(with empty lacuna), neutrophils and fibrin exudation (Fig. 3A). However, more neutrophils and extensive fibrin exudation were seen in the infected animals. At 4 and 6 days after inoculation (Groups B and C), a cellular layer of elongated fibroblasts, neutrophils, macrophages and giant cells were seen toward the implant cavity (Fig. 3B). This layer was surrounded by osteonecrotic trabecular bone intermingled with the same cell types. The most pronounced pathomorphological findings within PIBA of infected Groups B and C animals were massive extension of the cellular layer and active osteoclasts seen in resorptions lacuna of the necrotic bone trabecular (Fig. 3C,D). In the periphery of PIBA, fibroblasts and associated new collagen could be seen, although more pronounced in infected animals compared to controls. Generally, almost no bone tissue was present within PIBA of infected Group C animals.

**Localization of bacteria in bone tissue**

Bacteria were not identified by IHC in cortical bone tissue of any animal. Immunopositive *S. aureus* bacteria were present in all infected pigs of Groups A, B and C within both the exudate of the implant cavity and within PIBA (Table 1 and Fig. 4A). Additionally, *S. aureus* bacteria were also detected in one control animal of Group B within BIPA (Table 1 and Fig. 4B). IHC-positive *S. aureus* bacteria were not identified in the control animals of Groups A and C. On the tissue section representing the center of PIBA, the size of the largest *S. aureus*-positive aggregate stayed between 25 and 84 μm and was located from 25 to 2000 μm within PIBA regardless of the time of inoculation. Moreover, bacteria were found on two consecutively sections from distant bone pieces (2–3 mm each), not covering the implant cavity, in one infected animal of Groups A and B, each. All bacterial aggregates...
inside PIBA were seen within the cellular zone or adjacent to osteonecrosis.

**Microbiology**

All swab and Spa-typing results are presented in Table 1.

**Implants SEM and PNA FISH**

By SEM, coccoid bacteria were observed attached to the surface of all implants from both control and infected animals in the form of biofilm. The areas of biofilm were defined to isolated islands on the implant surface. In general, the bacteria were found embedded in a biomass of extracellular matrix, leukocytes and erythrocytes (Fig. 4C). Attached leukocytes appeared to be more dominant on implants of infected animals. By PNA FISH, *S. aureus* could be identified on one implant (Table 1 and Fig. 4D).

**DISCUSSION**

This study shows that pathomorphological changes and infecting bacteria of IAO can go up to 6 mm into the surrounding bone tissue within a week, in a porcine model (Fig. 2). The surrounding bone tissue was described as PIBA. Within PIBA, *S. aureus* bacteria could be identified both adjacent to and distant from the implant. However, despite...
bacteria in PIBA (shown with IHC), in the exudate within the implant cavity (shown with microbiology of swabs) and within the biofilm on the implants (shown with PNA FISH and SEM), bacterial aggregates could not be demonstrated laying on the surface of PIBA, that is, facing the implant cavity.

The present observations of peri-implanted bacteria are supported by a study of subcutaneous implant-associated infections (IAI) in a mouse model, showing that bacteria (*Staphylococcus epidermidis*) could be found within the tissue at a certain distance from the implant (22, 23). In orthopedic infections, bacteria are usually discussed in the context of biofilm formation directly on the implant (24). However, case reports of IAO have also confirmed that peri-implanted bone tissue may contain pathogenic bacteria (25, 26). Recently, by the same method as in this study (27), the size of bacterial colonies within peri-implanted bone tissue from patients with osteomyelitis has been estimated to range between 5 and 50 µm. In general, the maximal size of an *in vivo* biofilm has been estimated to be 200 µm (27). All measurements of *S. aureus* colonize within peri-implanted bone tissue of the present porcine model were below 84 µm. Therefore, the size of the observed bacterial colonies can be accepted as discriminative to human cases.
The Infectious Disease Society of America (IDSA) has recently proposed five criteria for the diagnosis of periprosthetic infections (28). The infected pigs from this study fulfilled criteria 2 (pus around the implant), 3 (histopathological evidence of inflammation) and 5 (positive intraoperative cultures). Additionally, the control animals also fulfilled criteria 3 and 5. However, Spa-typing results indicated that the control animals contaminated and infected themselves. The Spa types observed among control pigs isolates are typically seen in commensal porcine S. aureus strains (29) and differed from the Spa type used for inoculation (17). Self-contamination of the control animals thus highlights the hypothesis of the peri-implanted area as a locus resistentiae minoris; that is, a minimal number of bacteria can colonize an implant and the surrounding tissue. As S. aureus is the most common pathogen associated with IAO, and an independent risk factor of treatment failure (6), this bacterium was used. The selected porcine S. aureus strain was chosen, due to its ability to induce bone infection and inflammation (17, 18). The dose of 10^4 CFU/kg BW was estimated to be among the lowest infective dose, based on a former dose–response study using the same porcine S. aureus strain (18).

All animals scored 2+ or 3+ in number of neutrophils. The scoring system for the number of neutrophils was originally used on tissue section from cases of chronic low-grade IAO in humans (19). Specific neutrophil number cutoffs for high-grade IAO, to which infected animals of this study may belong, has not been established (30).

Although the number of infected and control animals killed on each time point is limited, grouping of pigs as either being infected (n = 6) or control (n = 6) animals allowed a statistical comparison of PIBA size (Fig. 2B). As significance was observed between these two groups, the group size was acceptable, that is, including more pigs could be a waste of animals. It was not possible to do statistical calculations between infected and control animals of each time points. However, the values from each time points can give an estimate of the variance and effect size in power/sample size calculations. This might be relevant for further experiments aiming to study effects of surgical or medical treatments in the porcine model. The histopathological examinations of all animals clearly demonstrated trabecular osteonecrosis around the implant cavity due to drilling of the bone prior to insertion of the implant. Additionally, the injection of S. aureus bacteria resulted in noticeable pathomorphological changes between infected and control animals, visualized by the size of PIBA (Fig 2A). The most obvious changes within infected animals were bacterial aggregates and accumulations of inflammatory cells, giant cells, active osteoclasts and fibroblasts. Thus, the present injection of bacteria, during insertion of the bone implants, resulted in biofilm formation on the implants and a peri-implanted bacterial reservoir directing bone pathomorphology. A peri-implanted bacterial reservoir might hamper a fast diagnosis and therapeutic debridement in cases of IAO. Furthermore, when located in the peri-implanted bone tissue, S. aureus bacteria can form biofilm (24), small colony variants (31) and be internalized by osteoblasts (31), three situations which favor persistence of the bacteria and infection.

The present porcine model of IAO showed that bacterial aggregation occurred on the surface of implants and within the surrounding bone tissue within 6 days of infection. Thus, it is important to notice that peri-implanted bone tissue shortly after implantation can serve as a reservoir for infecting bacteria. This is important knowledge for optimizing outcomes of surgical debridement in IAO. In clinical situations of acute IAO, that is, lasting for less than 3–4 weeks, simple lavage has been recommended (28). However, as this study shows, this approach might be problematic if the bacteria already within 1 week can be located half a centimeter into the surrounding bone tissue.

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