Abstract. Background/Aim: Prosthetic joint infection (PJI) remains a serious complication of total joint arthroplasty. To effectively treat PJI, it is essential to identify the microorganism causing it and be able to combine correct surgical and anti-infective treatments. This cannot always be achieved with the currently employed diagnostic methods. The aim of this study was to evaluate a semi-automated tissue biopsy bead milling method (Ultra-TurrAX, Axonlab AG; Reichenbach, Germany) based on the hypothesis that the results are more sensitive for microbe detection and less prone to contamination. Materials and Methods: We included 35 consecutive patients undergoing 38 hip or knee arthroplasty revisions in this study. In addition to manually processed biopsies, we processed tissue specimens harvested intraoperatively using a semi-automated method. The sensitivity and specificity of both methods were calculated using MSIS criteria and sonication results as gold standards. Results: For total hip arthroplasty samples were evaluated separately based on MSIS criteria as the reference standard, Ultra-TurrAX processing yielded 81% (62-100%) sensitivity and 100% specificity. Using sonication as the gold standard, a sensitivity of 80% (60-100%) and specificity of 100% (45-100%) were calculated. In total knee arthroplasty, Ultra-TurrAX processing yielded 27% (1-54%) sensitivity and 57% (20-94%) specificity when using MSIS criteria as the gold standard. Using sonication as the gold standard, a sensitivity of 60% (17-100%) and specificity of 77% (54-100%) were calculated. Conclusion: This is the first study to analyze bead mill processing in total hip and knee arthroplasty revisions in a consecutive patient series. The method’s sensitivity was comparable to and its specificity superior to regular sample processing results reported in the literature with respect to hip arthroplasties and to both hip and knee arthroplasties collectively. With respect to total knee arthroplasties, the method fared worse in our collective, most likely due to the small number of patients in the sample. Integrating the method into the clinical workflow allowed for speedier and more efficient sample handling and processing. The theoretical advantage of a lower risk of contamination because of fewer manual processing steps is, in our opinion, valid.

Prospective Analysis of a Sterile, Semi–automated Tissue Biopsy Homogenization Method in the Diagnosis of Prosthetic Joint Infections

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Prosthetic joint infection (PJI) remains a serious complication of total joint arthroplasty, and is likely to become even more important in the coming decades with the projected increase in life expectancy, population age and the number of arthroplasties performed. PJI poses a challenge not only due to the complexity of its surgical and medical treatment but also due to the difficulties encountered in establishing the correct diagnosis.

Currently, detecting infectious agents in synovial fluid aspirates or tissue biopsies is the mainstay of diagnosing PJI (1, 2). However, the reported accuracy of synovial fluid analysis varies in the literature, likely due to difficulties in detecting sessile organisms attached to biofilms (1, 3, 4). Growing microorganisms from tissue specimens collected for diagnostic purposes or during prosthesis exchange surgery is reported to lead to a higher detection rate (5).

To increase diagnostic accuracy and establish a common diagnostic standard, a set of diagnostic criteria was defined by the Musculoskeletal Infection Society (MSIS) (6, 7). However, these criteria do not necessitate a direct identification of the infectious agent and its susceptibility to certain antibiotics. Identifying the optimal anti-infective...
treatment to accompany a one- or multi-stage surgical exchange is, however, essential for overall success.

Consequently, we believe that improving tissue biopsy yields is indispensable to achieving higher detection rates and detection accuracy in diagnosing PJI.

Toward this end, a sterile, semi-automated tissue homogenization method using magnetic beads has been introduced (8, 9). As opposed to widely used manual trituration, the tissue specimen is transferred directly into a sealed, sterile vessel in the operating room. In the microbiology laboratory, the tissue, still in the sterile vessel, is then milled via the acceleration of metal beads also contained in the vessel in a magnetic field without having to open the vessel. In theory, this technique decreases the risk of contamination by minimizing manipulation of the sample. Furthermore, it increases the number of pathogens prone to detection, whereas with commonly used methods for processing biopsies, there is a certain number of pathogens situated in inaccessible parts of the specimen.

Thus, our hypotheses for this study were: 1. Sterile, semi-automated homogenization of tissue samples has superior sensitivity and specificity in the detection of PJI; 2. This method reduces the contamination rate. The aims of this study were: 1. To calculate the validity of microbiological detection after semi-automized homogenization, with MSIS-criteria and sonication, each defined as gold standards. 2. To compare the contamination rates of the two sampling methods.

**Materials and Methods**

In this study, we included consecutive patients of legal age to give informed consent who had any indication for one- or multi-stage exchange of a total hip or knee arthroplasty. Patients with acute DIs (both early postoperative and late onset hematogenous) and patients who did not give informed consent, or were unable to for medical reasons, were excluded.

**Preoperative diagnostic criteria.** For each joint, a preoperative diagnostic evaluation for PJI was performed according to our institution’s standards. Evaluations included a radiographic analysis for signs of loosening, serological assessment of the C-reactive protein (CRP) level, white blood cell (WBC) and neutrophil count, erythrocyte sedimentation rate (ESR) measurement, and synovial fluid aspirate culture. In cases with inconsistent findings for these assessments, our institution’s protocol calls for arthroscopic sampling for a tissue culture and histological analysis (10).

One or multi-stage prosthesis exchange was performed depending on whether the preoperative diagnostic findings suggested an infection or other types of failure.

**Postoperative diagnostic criteria.** As per institutional standards, we collected 3-6 tissue samples intraoperatively, that were sent to the microbiology laboratory for regular microbiological workup and semi-automated homogenization. They were allocated to either workup method as evenly as seen fit by the operating surgeon.

**Explanted prosthesis components were sonicated** (11), and the sonication results were used as the gold standard for postoperative sample assessment. Biopsy results inconsistent with the sonication results were deemed to be due to contamination if they only occurred in a single sample from a joint, if the microorganism was cultivated solely after enrichment, and if the histological analysis of the synovia-like interface membrane (SLIM) was not suggestive of infection (12). However, if a positive biopsy met the necessary criteria for PJI based on MSIS standards, the case was considered a PJI.

**Semi-automated homogenization.** For sample storage and transport, irradiated plastic vials with a sample volume capacity of up to 15 ml, containing 2 ml of water and 10 stainless-steel beads (Ultra-TurrAX, Axon Lab AG; Reichenbach, Germany), were used (Figure 1). Each sample was transferred into a vial, which was then sealed. For milling, the sealed vial was put in a dispersion device (Ultra-TurrAX Dispersion Device, Axon Lab AG; Reichenbach, Germany), which accelerated the beads magnetically for 3 min (Figure 2). For samples containing bone fragments or other tough tissues, the acceleration and duration can be adjusted. After homogenization, the sample can be used for standard microbiological culture and can be kept sealed and frozen for later analysis.

**Data analysis.** Binary parameters (MSIS criteria met vs. not met, sonication positive vs. negative, contamination vs. no contamination) were analyzed using chi-squared tests. We calculated the sensitivity and specificity, positive and negative predictive values and likelihood ratios for conventional tissue culturing and sterile homogenization, using sonication results and MSIS criteria as gold standards.

**Results**

Over a time period of 11 months (July 2015-May 2016), we included 35 (16 male, 19 female) patients with a total of 38 examined total hip and knee arthroplasties (20 hip, 18 knee arthroplasties) in this study. Average patient age was 70.3 (range=36-88) years.

Twenty-seven patients met the MSIS criteria for PJI, and 11 were classified as negative for PJI. Twenty patients had microorganisms recovered from their sonication fluid, and 18 patients did not.

Arthroscopy for the tissue sample workup before prosthesis exchange surgery was performed in 6 instances (3 hip, 3 knee arthroplasties). Two of these samples were classified as debris-induced (type 1) membranes, in one of which, the sonication result was positive and the MSIS criteria were negative. In the other, both the sonication result and MSIS criteria were negative. One histological sample was judged to show neither infection nor debris-induced synovitis (type 4 membrane), and both the sonication result and MSIS criteria were negative.

Two arthroscopies showed purulence in the joints in instances of recurrent acute infection; therefore, after collecting samples for microbiology workup, no further histological samples were collected, and the prostheses were
explanted in the same session.

**Semi-automized homogenization in total hip arthroplasty.** In the 20 total hip arthroplasties examined, the combination of regular tissue-sample workup and homogenization with the MSIS criteria as the reference standard yielded a sensitivity of 88% (71-100%) and a specificity of 75% (33-100%). Positive and negative predictive values for tissue biopsies overall were 93% (81-100%) and 60% (17-100%), respectively. The positive likelihood ratio was 3.5 (5.95-1.06), and the negative, 0.17 (-0.03-0.88).

With sonication results as the reference standard, the sensitivity and specificity of all tissue samples combined were 87% (69-100%) and 60% (17-100%), respectively. The positive predictive value was 87% (69-100%), and the negative predictive value was 60% (17-100%). The positive likelihood ratio was 2.17 (0.84-35.31), and the negative, 0.22 (0-0.88).

Evaluated separately and with MSIS criteria as the reference standard, Ultra-TurrAX-processed samples yielded an 81% (62-100%) sensitivity and 100% specificity, a positive predictive value of 100% and a negative predictive value of 57% (20-94%), with a negative likelihood ratio of 0.19 (0-0.38). Regular tissue samples yielded an 88% (71-100%) sensitivity, a 75% (33-100%) specificity, a positive predictive value of 93% (81-100%), a negative predictive value of 60% (17-100%), a positive likelihood ratio of 3.5 (1.06-5.95) and a negative likelihood ratio of 0.17 (0-0.88).

With sonication as the reference standard, a sensitivity of 80% (60-100%) and specificity of 80% (45-100%) were calculated for Ultra-TurrAX samples, along with positive and negative predictive values of 92% (78-100%) and 57% (20-94%), respectively, and positive and negative likelihood ratios of 4 (1.09-6.6) and 0.25 (0-0.9), respectively.

For regular samples, we calculated an 87% (69-100%) sensitivity, 60% (17-100%) specificity, 87% (69-100%) positive predictive value, and 60% (17-100%) negative predictive value, with a positive likelihood ratio of 2.17 (0.84-35.31) and a negative ratio of 0.22 (0-1.79). For an overview of the results with MSIS criteria as a gold standard, see Table I. For an overview of the results with sonication as a gold standard, see Table II.

**Semi-automated homogenization in total knee arthroplasty.** In the 18 total knee arthroplasties examined, the combination of regular tissue-sample workup and homogenization with MSIS criteria as the reference standard yielded a sensitivity of 27% (1-54%) and a specificity of 57% (20-94%). The positive and negative predictive values for tissue biopsies overall were 50% (10-90%) and 33% (7-60%), respectively. The positive likelihood ratio was 0.64 (0.01-8.65), and the negative, 1.27 (0.49-4.84).

With sonication results as the reference standard, the sensitivity and specificity of both methods combined were 60% (17-100%) and 77% (54-100%), respectively. The positive predictive value was 50% (10-90%), and the negative predictive value was 83% (62-100%). The positive likelihood ratio was 2.6 (0.37-5.93), and the negative, 0.52 (0-1.54).

Evaluated separately and with MSIS criteria as the reference standard, Ultra-TurrAX-samples yielded 27% (1-54%) sensitivity and 57% (20-94%) specificity, a positive
There was a total of 6 patients with suspected contamination of 1 or more samples.

Two patients had two contaminated regular tissue samples each: in the first case, the MSIS criteria were positive for infection, there was a type 3 periprosthetic membrane, and sonication, as well as regular and UltraTurrax, processing of biopsies yielded Propionibacterium granulosum in microbiological culture. One regular biopsy, however, was positive for Staphylococcus saprophyticus and another for Micrococcus luteus.

The second patient did not meet the MSIS criteria, histology was not suggestive of an infection (type 1 membrane), and sonication- and UltraTurrax-processed biopsies were negative. In two regular tissue specimens, however, Staphylococcus epidermidis was detected.

Three patients had likely contaminations in one UltraTurrax-processed biopsy each: The first patient had positive MSIS criteria and histological results, and sonication and regular processing of one tissue specimen yielded Staphylococcus aureus. One UltraTurrax-processed sample showed Staphylococcus epidermidis growth.

The second patient did not exhibit positive MSIS criteria, there was a type 1, debris-induced membrane, and regular tissue biopsies were negative. One UltraTurrax-processed sample showed Staphylococcus hominis growth.

predictive value of 50% (10-90%) and a negative predictive value of 33% (7-60%), with a positive likelihood ratio of 0.64 (0.01-8.65) and a negative likelihood ratio of 1.27 (0.49-4.84). Regular tissue samples yielded 60% (17-100%) sensitivity, 100% specificity, a positive predictive value of 100%, a negative predictive value of 87% (69-100%), and a negative likelihood ratio of 0.4.

With sonication as the reference standard, a sensitivity of 60% (17-100%) and specificity of 77% (54-100%) were calculated for Ultra-Turrax samples, along with positive and negative predictive values of 50% (10-90%) and 83% (62-100%), respectively. The positive and negative likelihood ratios were 2.6 (0.37-5.93) and 0.52 (0-1.54), respectively.

For regular samples, we calculated 60% (17-100%) sensitivity, 100% specificity, a 100% positive predictive value, an 87% (69-100%) negative predictive value, and a negative likelihood ratio of 0.4 (0-0.83). For an overview of the results with MSIS criteria as gold standard, see Table I. For an overview of the results with sonication as gold standard, see Table II.

Contaminations. There was a total of 6 patients with suspected contamination of 1 or more samples.
Discussion

In an experimental approach, bead mill tissue sample homogenization was superior to regular sample collecting and processing methods (8, 9). In theory, the described method has several advantages. First, it minimizes the number of manipulations in the sampling and evaluation process, which presumably reduces contamination rates. Second, homogenization theoretically enables an increased “bacteria yield” per sample as the homogenized tissue can be evenly distributed on culture media. In regular samples, microorganisms might persist in pouches or folds of tissue that is not thoroughly ground and be inaccessible. Third, in combination with reaching a consensus on tissue sample size, standardizing vessel size, the number of magnetic beads and milling time would present a further step towards a truly standardized approach for diagnosing PJI.

To our knowledge, there has only been one other study evaluating sterile semi-automated tissue homogenization in PJIs in a clinical setting (13). The authors reported a documentation rate for PJI of 83.7%, with a contamination rate of 8.7%. The patient collective consisted of 92 patients with known PJI. Our own data for hip and knee arthroplasties combined and for hip arthroplasties alone are comparable to those published by Roux et al., although our patient series did not consist only of patients with known PJI (13).

In comparison with other reported validations of tissue sampling, our data for knee arthroplasties show worse results, in respect to both regular and Ultra-TurrAX sampling methods (11, 14). The calculated validities for both methods in hip and knee arthroplasty and in hip arthroplasty infections alone, however, are comparable to previously published data (2, 11, 15-17).

Our first hypothesis, that semi-automated homogenization has superior specificity and sensitivity, was not supported. In total hip arthroplasties, the method had slightly lower sensitivity rates compared with the results of regular tissue sampling. However, specificity was considerably higher (100% vs. 75% with MSIS criteria as the gold standard, 80% vs. 60% with sonication results as the gold standard). The opposite was seen in total knee arthroplasties: here, the sensitivity and specificity achieved via Ultra-TurrAX-processing were invariably lower compared with the results of the regular sampling and processing method.

This inconsistency was not seen in a study by Schäfer et al., who were able to analyze hip and knee arthroplasty data as a common collective for a lack of differences (18). Our observations might be due to the small number of patients and the resulting lack of power of this study. Subdividing patients into hip and knee arthroplasty groups could have further accentuated the diverging trends in the results. A more homogenous patient collective (including only patients with known PJI) might have yielded more consistent data.

However, we used the trial Ultra-TurrAX vials made available to us in every clinical situation in which we collect tissue samples. While this choice of patient collective reduces the comparability of the results, one of our main interests was how well the method could be integrated into our workflow. In the personal opinion of the surgeons, OR personnel and laboratory technicians in our Microbiology Institute who were involved in the trial, sample handling and workflow using Ultra-TurrAX vials was speedier, more efficient and less prone to mistakes.

While the small number of patients and of documented contaminations prevents any statistical deductions, the number of contaminations was twice as high in regularly processed specimens. Whether this is proof of theoretical and anecdotal advantages of the method remains to be studied with a broader approach.

Conflicts of Interest

The Authors declare that there are no conflicts of interest.

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