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# ORIGINAL RESEARCH

# Microbiology of Periprosthetic Hip and Knee Infections in Surgically Revised Cases from 34 Centers in Mainland China

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Correspondence: Xi-Sheng Weng Department of Orthopedics, Peking Union Medical College Hospital, CAMS & PUMC, No. I Shuaifuyuan, Wangfujing, Dongcheng District, Beijing, 100730, People's Republic of China Tel/Fax +86-10-69152710 Email drweng\_wxs1377@163.com **Background and Aim:** Periprosthetic joint infection (PJI) is one of the most devastating complications after total joint arthroplasty (TJA). However, the antibiotic resistance of infecting pathogens can significantly vary in different parts of the country. In the current study, we analyzed the demographic and microbiological profiles of knee and hip PJI over three years and compared the microbiological differences between them.

**Methods:** A multicenter retrospective study of PJI patients in 34 referral medical centers in mainland China from January 2015 to November 2017 was performed.

**Results:** A total of 925 PJI patients were recruited, 452 were identified as knee PJIs, and 473 were hip PJIs. The most common causative pathogens were *Staphylococcus aureus* (26.5%) and coagulase-negative staphylococci (14.3%). Methicillin-resistant staphylococci were involved in 25.6% (237/925) of all PJI cases. Mycobacterium and fungus only accounted for 6.5% (61) of all cases. Enteric gram-negative bacilli, anaerobes, and polymicrobial pathogens were more common in hip joint prostheses than in knee PJI (P = 0.014; P = 0.006; P = 0.002, respectively).

**Conclusion:** While the majority of causative pathogens in PJI cases are staphylococcal species, the prevalence of atypical organisms and resistant pathogens should also be given attention and warrant the need for empiric antibiotic treatment.

Keywords: arthroplasty, hips, knees, microorganism, prosthetic joint infection

#### Introduction

Prosthetic joint infection (PJI), while rare, is probably the most feared complication arising from lower extremity total joint arthroplasty (TJA), as it presents a significant therapeutic challenge and places a substantial socioeconomic burden on the health care system.<sup>1–3</sup> The incidence of PJI after knee or hip replacement is approximately 1-2%,<sup>4,5</sup> and the number of primary and revision TJA cases rapidly increases in aged people. Thus, the incidence of PJI is also expected to grow in the coming years.<sup>6</sup>

The use of appropriate antibiotics is the key to the successful treatment of PJI. Microbiological and epidemiological data can guide prophylactic and therapeutic antibiotic regimens, which can help reduce the risk of bacterial resistance to antibiotics.<sup>7</sup> Different causative pathogens of PJI can lead to different treatment outcomes. For instance, it has been reported that methicillin-resistant *Staphylococcus aureus* (MRSA) PJI has a poorer prognosis and a higher risk of treatment failure.<sup>8</sup> Furthermore, polymicrobial infections are associated with

Infection and Drug Resistance 2021:14 2411-2418

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2411

a higher risk of treatment failure, amputations, and infection-related mortality than infections caused by a single organism.<sup>9</sup> Many studies have found significant differences in the microorganismal epidemiology of PJI in different centers.<sup>10,11</sup> There are also notable microbiological differences depending on the location of the affected joint.<sup>1,7</sup>

An accurate understanding of the microbiology of post-TJA PJIs is required to identify the most appropriate empiric antibiotics and therapeutic strategies. However, microbiological data and a comparison of different microorganisms involved in knee and hip PJIs are limited, especially in mainland China. This study, therefore, aims to compare the microbiological profiles between knee and hip PJIs in mainland China.

## Methods

## Patients

This retrospective cohort study included patients who received surgical treatment for knee and hip PJIs in 34 referral medical centers in mainland China between January 2015 and November 2017. This study was approved by the Institutional Review Board of Peking Union Medical College Hospital (S-K993). The study was conducted in compliance with the Declaration of Helsinki. Data after desensitization (hide patient name and contact information) then were entered into a computerized data management system. Only the chief PI and local PI got the access to all data in the system.

The criteria for eligible hospitals were: 1) referral hospitals where arthroplasty is performed in high volumes routinely (>500 hip/knee arthroplasty cases per year); 2) hospitals with enough technical and professional capability; 3) hospitals that can provide local Institutional Review Board (IRB) approval. At least one hospital was enrolled in each province of the Chinese mainland. A total of 41 research hospitals were selected, accounting for 3% (41/ 1308) of the total number of Grade A tertiary hospitals in mainland China. To ensure the accuracy of the information, each research team appointed special personnel to check the data and contacted the investigation hospital to see if there were any empty entries or logical errors. All cases were followed for at least one year. Finally, 34 hospitals were enrolled in this retrospective study.

After the approval by the institutional review board, investigators searched the electronic databases in the hospital. They retrospectively reviewed patients' data with the "International Classification of Diseases, Ninth Revision, Clinical Modification" code. The ICD-9-CM procedure codes 81.53, 00.70, 00.71, 00.72, 00.73 were used to identify RTHAs, whereas the codes 81.55, 00.80, 00.81, 00.82, 00.83, 00.84 were used to identify RTKAs associated with a principal diagnosis of PJI based on ICD-9-CM diagnosis code 996.66. Two independent investigators reviewed the extracted data. Patients with primary septic arthritis or infection after internal fixation and open reduction were excluded. Due to the retrospective nature of the study, informed consent was waived.

## Definition of PJI

PJI was defined according to the following criteria, which was modified from the international diagnostic standards (MSIS2013 or ICM2018): (1) isolation in one intraoperative culture of microorganisms, with evidence of infection at the site of the knee or prosthesis (presence of purulence in an affected joint, high synovial neutrophil percentage and synovial white blood cell count, elevated serum C-reactive protein levels and serum erythrocyte sedimentation rate, or positive histological examination). (2) Isolation of the same microorganism from two or more cultures obtained from an infected prosthetic joint; (3) a sinus tract communicating with the prosthesis. PJI can be classified as early-onset (occurring within three months after surgery), delayed onset (occurring three months to two years after surgery), or late-onset (occurring two years after surgery) based on the time to infection.<sup>12</sup> Polymicrobial PJI was defined as more than one microorganism isolated from one or more cultures of synovial fluid or periprosthetic tissues.

## Data Collection and Analysis

The demographic characteristics, causative pathogens, time of infection, site of arthroplasty, and comorbidities were recorded. Where there was more than one infection episode on the same joint, only the first episode was recorded. The microbiological profiles of all infection cases were analyzed, and the difference in the proportion of causative pathogens between knee PJIs and hip PJIs was compared.

## Statistical Analysis

All statistical analyses were two-sided and performed using the SPSS version 23 (IBM). Categorical data were

analyzed by Fisher's exact test. A *P*-value of <0.05 was considered statistically significant.

## Results

#### **Demographical Characteristics**

A total of 925 first-time infection cases (452 knee PJIs and 473 hip PJIs) from January 2015 to November 2017 were included in the retrospective study. There were 207 male patients with knee PJI (46%) and 108 male patients with hip PJI (70%). The hip PJI group had a mean age of 60.8 years and a male-to-female ratio of 2.1:1, whereas the knee PJI group had a male-to-female ratio of 0.84:1 and a mean age of 68.6 years. Bilateral joint infection was experienced by 14 patients with knee PJIs and eight patients with hip PJIs.

The incidence of alcoholism in patients with hip PJI was higher than in the knee PJI group (24% vs 5%; P = 0.01). However, the incidence of hypertension in patients with knee PJI was higher than in hip PJIs (55% vs 33%; P < 0.001). No difference in smoking, diabetes mellitus, cirrhosis, or malignancy was observed between the two groups. Also, there was no difference in the development of early-onset infection between patients with hip PJIs and those with knee PJIs (22.6% vs 27.9%; P = 0.102) (Table 1).

## Microbiology

The most common causative pathogens were *S. aureus* (26.5%) and coagulase-negative staphylococci (14.3%) (Table 2). Staphylococci had a high resistance rate to oxacillin or methicillin (Table 3). Methicillin-resistant coagulase-negative staphylococci were isolated in 98 patients (10.9%), and MRSA was isolated in 97 cases (10.5%). Up to 76% of coagulase-negative staphylococci (CoNS) and 40% of *S. aureus* were methicillin-resistant. Methicillin-resistant staphylococci (MRS) was involved in 25.6% of all PJI cases.

It was found that the distribution of causative microorganisms in knee PJIs was 52% aerobic gram positive, 7.3% gram negative, 2.2% fungus, 3.7% mycobacterium, 5.5% polymicrobial, and 29.0% negative culture. In hip PJIs, the distribution was 43.5% aerobic gram positive, 6.3% anaerobic, 8.8% gram negative, 2.5% fungus, 13.8% polymicrobial, and 24.9% negative culture. The proportion of causative microorganisms (anaerobic pathogens and polymicrobial pathogens) in knee PJIs was lower than that in hip PJIs (anaerobic pathogens: P = 0.002;

| Table I    | Demographics | of | Hip | and | Knee | Prosthetic | Joint |
|------------|--------------|----|-----|-----|------|------------|-------|
| Infections |              |    |     |     |      |            |       |

|                   |           | 1      |           |      |                     |
|-------------------|-----------|--------|-----------|------|---------------------|
| Demographic       | Hip       | %      | Knee      | %    | Р                   |
| Characteristics   | (n=473)   |        | (n=452)   |      |                     |
| Sex               |           |        |           |      |                     |
| Male              | 320       | 70     | 207       | 46   | <0.001              |
| Female            | 153       | 30     | 245       | 54   | <0.001              |
| Age (years), mean | 60.8±15.0 |        | 68.6±11.9 |      | <0.001ª             |
| ±SD (range)       | (21–82)   |        | (32–80)   |      |                     |
| Comorbidities     |           |        |           |      |                     |
| Hypertension      | 156       | 33     | 248       | 55   | <0.001 <sup>a</sup> |
| Smoking           | 127       | 27     | 85        | 19   | 0.08                |
| Diabetes mellitus | 89        | 19     | 117       | 26   | 0.26                |
| Alcoholism        | 113       | 24     | 24 22     |      | 0.01 <sup>a</sup>   |
| Cirrhosis         | 17        | 4      | 18        | 4    | 0.09                |
| Malignancy        | 14        | 3 19   |           | 4    | 0.22                |
| Time to infection |           |        |           |      |                     |
| Early onset       | 107       | 22.6   | 126       | 27.9 | 0.102               |
| Delayed onset     | 170       | 35.9   | 35.9 150  |      | 0.939               |
| Late onset        | 196       | 41.5   | 176       | 38.9 | 0.113               |
| ASA grade         |           |        |           |      |                     |
|                   | 62        | 13.1   | 36        | 7.8  | 0.022               |
| 2                 | 321       | 67.9   | 316       | 69.9 | 0.508               |
| 3                 | 90        | 19 100 |           | 22.  | 0.0127              |

Note: <sup>a</sup>A p value of <0.05 was considered to be statistically significant.

Abbreviation: ASA grade, The American Society of Anesthesiologists physical Sstatus classification system.

polymicrobial pathogens: P = 0.01), whereas the ratio of mycobacterium in knee PJIs was higher than that in hip PJIs (P = 0.01). There was no significant difference in the distribution of fungus, gram-positive and gram-negative microorganisms, and negative culture between hip and knee PJIs.

In polymicrobial hip PJI cases, 77.2% had two causative microorganisms, 4.5% had three, 9.0% had four, and 9.0% had five. In polymicrobial knee PJI cases, 100% had two causative microorganisms. *S. aureus* was the most common microorganism in polymicrobial hip PJIs. The causative microorganisms in polymicrobial infections are shown in Table 4.

## Discussion

To our knowledge, this study is the most recent and most extensive cohort study that reported the microbiological data of post-TJA PJI, comparing the differences between knee and hip PJIs in mainland China.<sup>13,14</sup> The study found that staphylococcal species were the most common organisms causing PJI. MRS was prevalent in both knee and hip PJIs. The prevalence of PJI-causing organisms was found

| Pathogen                   | Hip and Knee<br>[n=925, n (%)] | Positive<br>Cultures<br>Rate<br>(n=676, %) | Hip<br>[n=473, n (%)] | Positive<br>Cultures<br>Rate<br>(n=355, %) | Knee<br>[n=452, n (%)] | Positive<br>Cultures<br>Rate<br>(n=321, %) | Ρ                  |
|----------------------------|--------------------------------|--|-----------------------|--|------------------------|--|--------------------|
| Aerobic gram-positive      | 441 (47.6)                     | 65.2                                       | 206 (43.5)            | 58.0                                       | 235 (52.0)             | 73.2                                       | 0.116              |
| Staphylococcus aureus      | 245 (26.5)                     | 36.4                                       | 111 (23.5)            | 31.1                                       | 134 (30.0)             | 43.2                                       | 0.169              |
| Coagulase-negative         | 132 (14.3)                     | 19.6                                       | 72 (15.2)             | 20.3                                       | 60 (13.3)              | 18.9                                       | 0.667              |
| Staphylococci              |                                |  |                       |  |                        |  |                    |
| Streptococcus species      | 40 (4.4)                       | 6.1  | 15 (3.1)              | 4.2  | 25 (5.5)               | 8.4  | 0.248              |
| Enterococcus faecalis      | 8 (0.8)                        | 0.9  | 3 (0.6)               | 0.8  | 5 (1.1)                | 1.1  | 0.907              |
| Listeria monocytogenes     | 8 (0.8)                        | 0.9  | 3 (0.6)               | 0.8  | 5 (1.1)                | 1.1  | 0.907              |
| Unspecified                | 8 (0.8)                        | 0.9  | 0 (0.0)               | 0.0  | 8 (1.8)                | 2.1  | 0.124              |
| Anaerobes                  | 32 (3.5)                       | 5.1  | 31 (6.3)              | 9.2  | I (0.002)              | 0.0  | 0.002 <sup>a</sup> |
| Propionibacterium species  | 16 (1.7)                       | 2.3  | 15 (3.1)              | 4.2  | I (0.002)              | 0.0  | 0.038 <sup>a</sup> |
| Peptostreptococcus species | 13 (1.4)                       | 1.9  | 13 (2.5)              | 3.4  | 0 (0.0)                | 0.0  | 0.064              |
| Prevotella species         | 3 (0.4)                        | 0.5  | 3 (0.6)               | 0.8  | 0 (0.0)                | 0.0  | 0.356              |
| Gram negative              | 75 (8.2)                       | 11.2                                       | 42 (8.8)              | 11.8                                       | 33 (7.3)               | 10.5                                       | 0.663              |
| Escherichia coli           | 26 (2.8)                       | 3.7  | 18 (3.8)              | 5.0  | 8 (1.7)                | 2.1  | 0.229              |
| Pseudomonas aeruginosa     | 16 (1.7)                       | 2.3  | 2 (0.4)               | 0.8  | 14 (3.0)               | 4.2  | 0.123              |
| Pseudomonas stutzeri       | 3 (0.3)                        | 0.5  | 0 (0.0)               | 0.0  | 3 (0.7)                | 1.1  | 0.277              |
| Serratia marcens           | 13 (1.4)                       | 1.9  | 6 (1.3)               | 1.7  | 7 (1.5)                | 2.1  | 0.869              |
| Klebsiella pneumoniae      | 9 (1.0)                        | 1.4  | 9 (1.9)               | 2.5  | 0 (0.0)                | 0.0  | 0.109              |
| Enterobacter cloacae       | 3 (0.3)                        | 0.5  | 3 (0.6)               | 0.8  | 0 (0.0)                | 0.0  | 0.356              |
| Salmonella enterica        | 3 (0.3)                        | 0.5  | 3 (0.6)               | 0.8  | 0 (0.0)                | 0.0  | 0.356              |
| Morganella morganii        | 3 (0.3)                        | 0.5  | 0 (0.0)               | 0.0  | 3 (0.7)                | 1.1  | 0.277              |
| Fungus                     | 21 (2.3)                       | 3.3  | 12 (2.5)              | 3.4  | 9 (2.2)                | 3.2  | 0.869              |
| Candida species            | 15 (1.7)                       | 2.3  | 9 (1.9)               | 2.5  | 7 (1.5)                | 2.1  | 0.789              |
| Unspecified                | 6 (0.6)                        | 0.9  | 3 (0.6)               | 0.8  | 2 (0.7)                | 1.1  | 0.907              |
| Mycobacterium              | 16 (1.7)                       | 2.3  | I (0.0)               | 0.0  | 15 (3.7)               | 5.3  | 0.014 <sup>ª</sup> |
| Mycobacterium tuberculosis | 2 (0.7)                        | 0.9  | 0 (0.0)               | 0.0  | 2 (1.5)                | 2.1  | 0.124              |
| M. tuberculosis complex    | 2 (0.7)                        | 0.9  | 0 (0.0)               | 0.0  | 2 (1.5)                | 2.1  | 0.124              |
| Mycobacterium fortuitum    | I (0.3)                        | 0.5  | 0 (0.0)               | 0.0  | I (0.7)                | 1.1  | 0.277              |
| Polymicrobial              | 90 (9.8)                       | 13.1                                       | 65 (13.8)             | 18.5                                       | 25 (5.5)               | 6.3  | 0.006 <sup>a</sup> |
| Negative cultures          | 249 (26.9)                     |  | 118 (24.9)            |  | 131 (29.0)             |  | 0.391              |
| Total                      | 925 (100.0)                    |  | 473 (100.0)           |  | 452 (100.0)            |  |                    |
| L                          | 1                              |  |                       |  |                        |  |                    |

| Table 2 | Microorganisms     | in Hi | D and Knee  | e Periprosthetic  | loint | Infection |
|---------|--------------------|-------|-------------|-------------------|-------|-----------|
| Table 1 | i nei ooi gamismis |       | p and renee | , i cripi osuicuc | Jonic | meetion   |

**Note:** <sup>a</sup>A p value of <0.05 was considered to be statistically significant.

to be different between infected hip and knee joints: enteric gram-negative bacilli, anaerobes, and polymicrobial pathogens were more likely to occur in hip prosthetic joints than in knee cases.

Due to the difference in population baseline data and operation conditions in mainland China and abroad, the value of PJI microbiological epidemiology results reported in foreign literature is limited in China. Most of the previous microbiological data of PJI cases reported in mainland China were from a single center<sup>15–17</sup> or studies with bias factors or a relatively small sample size. This caused a lack of recent PJI microbiological epidemic results in guiding the antibiotic treatment of PJI in mainland China. The present results will help improve this situation.

Several reports have shown that the microbiological profile may vary in different countries. In 2006, a retrospective case series analysis<sup>18</sup> of 34 primary hip arthroplasties and 41 primary knee arthroplasties revised for PJI showed that staphylococci species were implicated in 61% of cases (*S. aureus* = 25%, coagulase-negative *Staphylococcus* = 36%). The mean time to diagnosis was 1.15 years. In 2001, it was found that the predominant infectious organisms were gram positive in 6489 knee replacements (Group B *Streptococcus, Staphylococcus* 

 Table 3 Methicillin Resistance of Staphylococcal Periprosthetic
 Joint Infection

| Staphylococci        | Hip<br>[n=206, n (%)] | Knee<br>[n=235, n (%)] | Р    |
|----------------------|-----------------------|------------------------|------|
| MSSA <sup>a</sup>    | 72 (35.0)             | 107 (45.5)             | 0.71 |
| MRSA <sup>b</sup>    | 51 (24.8)             | 68 (28.9)              | 0.75 |
| MS-CoNS <sup>c</sup> | 15 (7.3)              | 10 (4.3)               | 0.47 |
| MR-CoNS <sup>d</sup> | 68 (33.0)             | 50 (21.3)              | 0.12 |
| Total                | 206 (100.0)           | 235 (100.0)            |      |

**Notes:** <sup>a</sup>Methicillin-sensitive *Staphylococcus aureus*. <sup>b</sup>Methicillin-resistant *S. aureus*. <sup>c</sup>Methicillin-sensitive coagulase-negative staphylococci. <sup>d</sup>Methicillin-resistant coagulase-negative staphylococci.

Abbreviations: MSSA, methicillin-sensitive *Staphylococcus aureus*; MRSA, methicillin-resistant *Staphylococcus aureus*; MS-CoNS, methicillin-sensitive coagulasenegative staphylococci; MR-CoNS, methicillin-resistant coagulase-negative staphylococci.

*epidermidis*, and *S. aureus*).<sup>19</sup> In 2009, a large European registry study investigated 426 infected knee arthroplasties recorded in the Swedish Knee Arthroplasty Register between 1986 and 2000.<sup>10</sup> CoNS and *S. aureus* were the most common organisms (27.5% and 30.5% of all cases, respectively). Staphylococci were involved in around 58% of PJI cases, which was lower than what we reported in the present study. They also found a significant increase in staphylococcal resistance to methicillin over the duration.

In 2014, Aggarwal et al<sup>11</sup> performed a retrospective study of deep PJI in two high-volume infection referral centers in the US and Europe. They studied intraarticular tissue cultures taken at the revision of primary knee and hip arthroplasty for indications of PJI. The study enrolled 921 patients with hip replacements and 749 patients with knee replacements. There was a significant difference in microorganism epidemiology between the two centers: the incidence of *S. aureus* infection was significantly higher in the US center (31.0%) than in the European center (13.0%). CoNS infections, however, were seen particularly more frequently in the European center (39.3%) than in the US center (20.2%). More recently (2016), Holleyman et al<sup>20</sup> found that *Staphylococcus* was the most common organism isolated after the revision of a primary implant for infection (in 72% of all cases).

Certain differences in the proportions and types of pathogens have been observed with regard to the joint that is infected. The Mayo Clinic database shows that patients with hip arthroplasty have a lower frequency of *S. aureus* than coagulase-negative staphylococcal infection compared with those with infected knees, where the two types of staphylococci are relatively equal.<sup>7</sup> Moreover, a previous study comprising 159 hip PJI and 135 knee PJI patients showed that anaerobic bacteria, enteric gram-

negative bacilli, and polymicrobial pathogens are more frequently identified in hip than in knee infections,<sup>14</sup> which was consistent with our findings. The possible reasons that may lead to the differences remain unknown. Further investigations are needed to explore the pathological mechanisms of pathogen infection at different joint types.

In the present study, methicillin-sensitive *S. aureus* (MSSA) and MS-CoNS were found to be potent pathogens and common causes of PJI. The antibiotic standard of care therapy for MSSA infections includes penicillinase-resistant penicillin (flucloxacillin/oxacillin/nafcillin), with first-generation cephalosporin and cefazolin as an alternative.<sup>21,22</sup> The third- or fourth-generation cephalosporins or cephalosporin may be used for MSSA infections for penicillin-allergic patients.<sup>23</sup>

However, several reports have shown an increase in the prevalence of resistant organisms.<sup>5,24</sup> The present study showed that the occurrence of MRS was high, with 76% of CoNS and 40% of S. aureus being methicillin-resistant. In 2005, a retrospective case series analysis from Hong Kong<sup>16</sup> characterized bacteria isolates between 1995 and 2003. The authors found no multiple-drug resistant isolates between 1995 and 1996; however, they observed a change in later years of the study. McLawhorn et al<sup>25</sup> found that MRSA and MRSE combined accounted for 18.1% of PJI pathogens in the US Aggarwal et al<sup>11</sup> showed that methicillin resistance was significantly more prevalent in the US than in Europe. These differences indicate geographical variability of organism prevalence, further addressing the need for registry data analysis at the population level. Owing to the high incidence of MRS in both infected hip and knee periprosthetic joints, physicians may need to select empiric antibiotics with broader coverage while waiting for culture results.

The incidence rates of polymicrobial infections in hip and knee prostheses were significantly different (65 vs 25 cases, P = 0.006), which is in accordance with previous studies.<sup>9,14</sup> Varying incidences (6 to 37%) of polymicrobial infection have been reported.<sup>11,26</sup> The literature also consistently demonstrated that patients with polymicrobial infections had inferior treatment outcomes. Tan et al<sup>27</sup> showed that patients with polymicrobial PJI had a higher failure rate (50.5%), as well as a higher rate of arthrodesis (odds ratio [OR] = 11.06), amputation (OR = 3.80), and mortality (OR = 7.88) compared with those with monomicrobial PJI. A series

| Table 4 | 4 Causative | Microorganism | is in Cases | of First-Epi | isode Poly | microbial Hip | and Knee Per | iprosthetic | oint Infection |
|---------|-------------|---------------|-------------|--------------|------------|---------------|--------------|-------------|----------------|
|         |             | 0             |             |              |            | I             |              |             |                |

| Pathogen                         | Hip<br>(n =65) | Isolates<br>Rate<br>(n= 161) | Polymicrobial<br>Infections<br>Rate (n=65) | Knee<br>(n=25) | Isolates<br>Rate<br>(n=50) | Polymicrobial<br>Infections<br>Rate (n=25) | Р                 |
|----------------------------------|----------------|------------------------------|--|----------------|----------------------------|--|-------------------|
| Aerobic gram-positive            |                |                              |  |                |                            |  |                   |
| Staphylococcus aureus            | 35             | 21.8                         | 53.8                                       | 14             | 28.0                       | 56.0                                       | 0.81              |
| Coagulase-negative staphylococci | 15             | 9.1                          | 22.7                                       | 4              | 8.0                        | 8.0  | 0.93              |
| Streptococcus species            | 9              | 5.5                          | 13.6                                       | 8              | 16.0                       | 0.0  | 0.18              |
| Enterococcus faecalis            | 9              | 5.5                          | 13.6                                       | 4              | 8.0                        | 0.0  | 0.70              |
| Enterococcus faecium             | 3              | 1.8                          | 4.5  | 4              | 8.0                        | 0.0  | 0.64              |
| Corynebacterium species          | 6              | 3.6                          | 9.1  | 4              | 8.0                        | 0.0  | 0.48              |
| Unspecified                      | 0              | 0.0                          | 0.0  | 0              | 0.0                        | 0.0  | 0.03 <sup>a</sup> |
| Anaerobes                        |                |                              |  |                |                            |  |                   |
| Peptostreptococcus species       | 15             | 9.1                          | 22.7                                       | 0              | 0.0                        | 0.0  | 0.28              |
| Clostridium species              | 3              | 1.8                          | 4.5  | 0              | 0.0                        | 0.0  | 0.64              |
| Bacteroides fragilis             | 3              | 1.8                          | 4.5  | 0              | 0.0                        | 0.0  | 0.64              |
| Prevotella species               | 3              | 1.8                          | 4.5  | 0              | 0.0                        | 0.0  | 0.64              |
| Fusobacterium nucleatum          | 3              | 1.8                          | 4.5  | 0              | 0.0                        | 0.0  | 0.64              |
| Veillonella species              | 3              | 1.8                          | 4.5  | 0              | 0.0                        | 0.0  | 0.64              |
| Gram-negative                    |                |                              |  |                |                            |  |                   |
| E. coli                          | 9              | 5.5                          | 13.8                                       | 0              | 0.0                        | 0.0  | 0.41              |
| K. pneumoniae                    | 12             | 7.3                          | 18.4                                       | 0              | 0.0                        | 0.0  | 0.34              |
| Enterobacter species             | 6              | 3.6                          | 9.6  | 4              | 8.0                        | 12.0                                       | 0.48              |
| Proteus vulgaris                 | 3              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Salmonella enterica              | 3              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Pseudomonas aeruginosa           | 3              | 1.8                          | 4.6  | 4              | 8.0                        | 16.0                                       | 0.23              |
| Acinetobacter baumannii          | 3              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Aeromonas hydrophila             | 3              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Moraxella species                | 0              | 0.0                          | 0.0  | 4              | 8.0                        | 16.0                                       | 0.03ª             |
| Unspecified                      | 2              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Fungus                           |                |                              |  |                |                            |  |                   |
| Candida species                  | 3              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Aspergillus versicolor           | 3              | 1.8                          | 4.6  | 0              | 0.0                        | 0.0  | 0.64              |
| Mycobacterium                    |                |                              |  |                |                            |  |                   |
| Mycobacterium chelonae           | 0              | 0.0                          | 0.0  | 0              | 0.0                        | 0.0  | 0.00              |
| Total isolates                   | 161            | 100.0                        |  | 50             | 100.0                      |  |                   |

Note:  $^{a}A p$  value of <0.05 was considered to be statistically significant.

analysis of 77 polymicrobial PJIs also demonstrated that<sup>28</sup> the infection-free rate after two years was 67.6% for polymicrobial infections as compared with 87.5% for monomicrobial infections. Additionally, Marculescu et -  $al^{29}$  reported that the two-year cumulative probability of success of monomicrobial PJIs was 72.8%, compared with 63.8% for polymicrobial PJIs. These findings offer three possible explanations for the increased failure rate in polymicrobial PJIs: (1) association with a soft tissue defect or a sinus tract; (2) frequent presence of difficult-to-treat organisms;<sup>27,29</sup> (3) increased comorbidities.

Moreover, the total negative culture rate was high (26.9%) in this study. A possible explanation for this might be the administration of antibiotics before obtaining culture samples. Many of the referral patients had already received antibiotic treatments, and therefore the medicines might not be held for at least two weeks when culture samples were obtained. Another possible reason is that culture samples obtained at the time of surgery were limited: some cases, for example, only had one swab culture sample. A minimum of three specimens for culture is recommended to maintain the same level of certainty in detecting PJIs.<sup>21,30,31</sup> A prospective study<sup>32</sup> examined 297

revision TJA procedures using multiple detection methods, and it was recommended that five or six specimens be collected to diagnose an underlying infection accurately. At least three samples should yield the growth of the underlying microorganism for adequate diagnosis of infection. According to the current department protocol, we collected at least three tissue samples, aspirated the synovial fluid, and injected it into blood culture bottles, improving the positive culture rate.

Despite the overall results obtained in the study, it had some limitations. First, there is a collection bias due to the retrospective design. As PJI cases that did not undergo revision surgeries were not included, data of this group infected by low-virulence pathogens may be missing. Second, we only included the first episode of PJI. Finally, although the study was performed at 34 referral hospitals in mainland China, some patients received primary arthroplasty and partial treatment for PJI at other institutions, which may have also caused selection bias. Hence, it is necessary to establish a national joint arthroplasty registry.

## Conclusion

This study found that the primary organisms causing post-TJA PJI are from the *Staphylococcus* genus. The prevalence of atypical organisms and resistant pathogens may warrant the need for empiric antibiotic therapy with broader coverage. The negative culture rate was high, so the current protocol on cultures needs improvement to maintain the same level of certainty. Fungal and mycobacterial PJI cases were found to be rare. Further work regarding PJI organism profiles is required, as it may confer significant clinical implications.

## Acknowledgments

The authors are particularly grateful to all the panelists for their participation. Doctor Wei Huang, M.D., Yi-He Hu, M. D., Wan-Chun Wang, M.D., Tang Liu, M.D., Jin-Cheng Wang, M.D., Jian-Lin Zuo, M.D., Shu-Qing Tao, M.D., Qi-Rong Qian, M.D., Ying-Zhen Wang, M.D., Tao Li, M.D., Zhi-Qiang Zhang, M.D., Ya-Yi Xia, M.D., Yong-Qing Xu, M.D., Rui He, M.D., Wei-Hua Xu, M.D., Xiao-Bin Tian, M. D., Jun Xiao, M.D., Wen-Bo Wang, M.D., Liao-Bin Chen, M. D., Qun-Hua Jin, M.D., Hong Yuan, M.D., Qing Jiang, M.D., Wen-Ming Zhang, M.D., Wei-Ming Liao, M.D., Zhan-Jun Shi, M.D., Jian-Zhong Xu, M.D. and Jian-Ning Zhao, M.D.

## Funding

The authors received financial support for the research from key projects of National Natural Science Foundation of China (NO. 81630064).

## Disclosure

The authors declare that they have no competing interests.

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