



## White paper

## The Paradox of Prosthetic Joint Infection and the Microbiome: Are Some Bacteria Actually Helpful?

Ayesha Abdeen, MD, FRCS(C)<sup>a, \*</sup>, Craig J. Della Valle, MD<sup>b</sup>, Daniel Kendoff, MD<sup>c</sup>,  
Antonia F. Chen, MD<sup>d</sup><sup>a</sup> Department of Orthopaedic Surgery, Boston Medical Center, Boston, MA, USA<sup>b</sup> Midwest Orthopaedics at Rush, Chicago, IL, USA<sup>c</sup> HELIOS Klinikum Berlin-Buch, Berlin, Germany<sup>d</sup> Department of Orthopaedic Surgery (Affiliated with Harvard Medical School), Brigham and Women's Hospital, Boston, MA, USA

## ARTICLE INFO

## Article history:

Received 16 August 2021

Accepted 30 November 2021

Available online xxx

## Keywords:

Microbiome

Periprosthetic joint infection

Dysbiosis

Arthroplasty

## ABSTRACT

Periprosthetic joint infection (PJI) is a potentially catastrophic complication of total joint arthroplasty of the lower extremity. PJI is associated with significant burden of illness and economic cost. There are a number of well-established modifiable risk factors for PJI. Myriad perioperative protocols are used with the intent of reducing the incidence of PJI. However, it remains unclear why infections occur despite correction of modifiable risk factors and/or adherence to prophylactic protocols. There is emerging evidence that the microbiome—the diverse population of commensal microorganisms that inhabit the human body—may play a role in the pathogenesis of musculoskeletal infections. The impact of the microbiome on PJI warrants further investigation and may change how we conceptualize, prevent, and treat PJI.

© 2021 Published by Elsevier Inc. on behalf of The American Association of Hip and Knee Surgeons. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

## Introduction

Total joint arthroplasty (TJA) of the hip and knee is among the most successful surgical innovations of the past century given the restoration of function and reduction of pain. The problem of the greatest magnitude in TJA is perhaps one that we understand the least, periprosthetic joint infection (PJI). PJI is conceivably the most devastating complication after TJA as it is associated with potentially catastrophic outcomes, such as loss of limb or life [1]. PJI is the primary mode of failure of total knee arthroplasty (TKA) and the 3rd most common reason for revision in total hip arthroplasty (THA) [2]. Although PJI occurs in less than 1%–2% of primary TJA cases, the impact is substantial given the high volume of TJA cases performed nationally [2]. By 2030, THA and TKA cases performed annually in the US are projected to grow to 1.26 million and 935,000 cases, respectively [3], and hence, the

absolute number of infections will unfortunately only rise over time.

## Problem statement

Complications of PJI including local recurrence of infection and systemic sepsis pose a significant burden of illness. The treatment of chronic PJI often involves revision surgery and, in some instances, requires permanent implant removal, fusion or amputation, or prolonged antimicrobial therapy [4–6]. The cost associated with these treatments including patient time lost from work and productivity, along with the impact on family members and friends, all amounts to a substantial economic burden. It is estimated that the combined annual hospital costs related to PJI of the hip and knee in the US will be \$1.85 billion by 2030 [2].

Multiple patient and surgical factors play a role in the pathogenesis of PJI. A number of modifiable risk factors for PJI have been identified, including smoking, obesity, diabetes, vitamin D deficiency, and malnutrition, and these are now the central focus of perioperative optimization and infection prevention [7–9]. Surgical strategies to reduce PJI pertain to reducing bacterial bioburden at the surgical site, which has led to a number of potential strategies including preoperative methicillin-resistant and methicillin-

\* Corresponding author. Boston Medical Center, Department of Orthopaedic Surgery, 850 Harrison Avenue, Dowling 2 North, Boston, MA, USA, 02118. Tel.: (617) 414-0684.

E-mail address: [ayasha.abdeen@bmc.org](mailto:ayasha.abdeen@bmc.org)

sensitive *Staphylococcus aureus* screening and decolonization, appropriately timed and selected intravenous (IV) antibiotic prophylaxis, preoperative skin decolonization, and using alcohol-based skin preparation solutions [10–13]. Other methods involve attempts to decrease bacterial contamination in the operating room (OR) environment, including limiting OR traffic, utilizing laminar flow, and donning exhaust suits [14,15]. Many of these techniques remain controversial and are not universally used. Despite implementation of patient and surgical optimization techniques, and extensive efforts to reduce infection, PJI still occurs, indicating that current modes of prevention are insufficient.

## Proposed solution

### *A new paradigm of the role of bacteria in the pathogenesis of PJI*

The continued threat of PJI, often in the absence of known patient risk factors or deviation from preventative protocols, calls into question the current paradigm of the pathogenesis and prevention of these infections. The human body is inhabited by a host of commensal microbes in the skin, respiratory tract, genitourinary tract, oral cavity, and gastrointestinal (GI) tract, also known collectively as the “microbiome” [16]. Colonization of the microbiome with bacteria, fungi, protists, and archaea, collectively called “microbiota,” starts at birth and in the first year of life whereby a newborn is exposed to maternal and environmental microbes that initiate the gut microbiota [16].

It is estimated that 500–1000 distinct species live in the human body at any given time [16,17]. The composition of the microbiome is affected by host genotype and early bacterial exposure that remains relatively stable in an individual; however, the microbiome can be altered by other environmental factors such as antibiotic use and diet [16–18]. The unique, symbiotic interplay between microbiome and host is thought to play a major role in the immune function and overall health of the host.

In 2007, the National Institute of Health launched the Human Microbiome Project to evaluate human microbiota [17]. This endeavor led to a number of findings pertaining to disease pathogenesis and treatment for a variety of disease states. The microbiota of the gut comprises the majority of the human microbiome [16,18]. Disruption of these symbiotic relationships, called “dysbiosis,” is attributed to a broad array of disease states including inflammatory bowel disease (Crohn’s disease and ulcerative colitis), upper respiratory disease, autoimmune disorders, liver disease, obesity, cancers, and major depressive disorder [16,19,20]. Conditions specifically associated with an increased risk of PJI of the hip and knee, including obesity and diabetes mellitus, are related to an imbalance of the microbiome [21,22]. Alterations in the microbiome have been linked to other musculoskeletal conditions including increased incidence of bone fragility [18,23].

Hence, we are presented with a paradox: some bacteria may actually help our cause. More than 100 trillion organisms inhabit the human GI tract, rendering it the primary site of interactions between microbes and the immune system [24]. More than 70% of the body’s lymphocytes are located in the GI tract, making it the largest immune organ [24]. The microbiota of the GI tract stimulates and enhances the function of the immune system [18]. Strains of commensal bacteria isolated from the human gut have been shown to confer immunomodulatory capabilities, such as regulation of a range of cytokines including interleukin (IL)-10, IL-17a, IL-22, and interferon gamma [25]. Disruptions of the gut microbiome have been demonstrated to reduce the number and effectiveness of macrophages, thus rendering the host less able to respond to pathogenic bacteria [26]. The link between dysbiosis and increased infection risk has been established; reduced diversity of the gut

microbiota has been shown to increase the risk of infection after hematopoietic stem cell transplantation [27,28]. A paucity of diversity of the gut microbiome has also been linked to PJI in pre-clinical studies, as Hernandez et al. demonstrated in a murine model that a change in the microbiome that decreased the presence of normal gut flora increased the incidence of PJI and impaired the immune response to infection [18].

Imbalance in the microbiome may also explain the seemingly incongruous connection between malnutrition and obesity with PJI. It is well established that morbid obesity increases the risk of PJI [8]. However, the mechanism for this is not well understood; it remains speculative whether the increased adipose layer results in incisional complications that lead to superficial infections then become deep PJIs. Inexplicably, patients who undergo bariatric surgery and weight loss remain at a high risk for PJI after primary and revision TJA [29]. Conversely, undernourished patients with low body mass index are also at an increased risk of PJI [30]. The unifying link between underweight patients, overweight patients, and those who underwent weight reduction surgery that results in increased PJI risk may be nutritional deficits [31]. Malnutrition and obesity are both conditions that are linked to dysfunction of the microbiome, as obese and undernourished patients both have reduced diversity of the gut microbiota [32]. Perturbations in the gut microbiome have a profound effect on nutrient absorption [33]. This common finding in obese and undernourished patients paired with the increased risk of PJI in these patients suggests that dysbiosis may be involved in the pathogenesis of PJI.

It remains unclear why microflora of one region, such as the gut, appears to be protective against PJI, whereas other commensal bacteria are pathogenic. *Staphylococcus aureus* that colonizes the nares and skin of 20% of the population is the target of decolonization protocols before TJA [10]. In rare instances, microbiota of the oral cavity or genitourinary tract has been identified as pathogens in osteoarthritis and PJI [34,35]. There are at least 2 case reports of PJI with *Fusobacterium nucleatum*, an anaerobic gram-negative bacillus of the oral microbiota, and 2 cases of hip PJI involving *Gardnerella vaginalis* of the genitourinary tract [36,37]. In these unusual cases, PJI was treated successfully without reported recurrence, suggesting that these commensal bacteria may have resulted in a less virulent form of PJI. Further investigation is needed to distinguish which commensal bacteria are pathogenic vs beneficial in PJI.

It is well known that prolonged antibiotic therapy, which eradicates helpful commensal organisms, inadvertently allows pathogens to thrive including *Candidiasis* or *Clostridium difficile* that can lead to bowel infections [38,39]. Short-term prophylactic antibiotics are unlikely to cause dysbiosis, whereas prolonged or suppressive antibiotic therapy has been demonstrated to reduce the diversity of the microbiome, resulting in increased susceptibility to infection [40]. It is thus possible that prolonged antimicrobial therapy may likewise increase the risk of PJI. While hematogenous spread of an infection from one location (such as oral abscess or recurrent urinary tract infection) to a TJA is a cause for concern, it is possible that the antibiotics used to treat that infection can create an additional risk owing to disruption of the microbiome. Paradoxically, treatment for PJI with prolonged antimicrobials and long-term suppressive antibiotics used for the treatment of PJI may also pose a risk of recurrence or new infection by virtue of disrupting the microbiome. Further study is required to determine the extent and duration of dysbiosis after antimicrobial therapy and whether this poses a risk for PJI during that timeframe.

## Future directions and long-term focus

If dysfunction within a microbiome can result in disease, it follows that interventions to restore the microbiome could potentially

prevent or treat the disease state. An expert panel was convened in October 2013 by the International Scientific Association for Probiotics and Prebiotics to discuss the field of probiotics [41]. The Food and Agriculture Organization of the United Nations and the World Health Organization definition of probiotics is “live microorganisms which when administered in adequate amounts confer a health benefit on the host” [41]. Probiotics have been demonstrated to enhance innate immunity and suppress the growth of pathogenic organisms. The mechanisms by which this is accomplished include inhibition of IL-10 and stimulation of secretory IgA and reduction of inflammatory cytokines [42].

Probiotics (beneficial bacteria), prebiotics (fiber), and a combination of both (synbiotics) have been found to reduce the incidence of surgical site infections after colorectal and hepatobiliary surgery and have been shown to confer incisional wound healing properties after surgery [43,44]. A meta-analysis by Skonieczna-Zydecka et al. identified 35 randomized control trials investigating the preoperative and/or postoperative use of probiotics, prebiotics, and/or synbiotics compared with controls (placebo or standard of care) after a variety of surgical procedures, including liver transplant, colectomy, esophagectomy, pancreaticoduodenectomy, hepatectomy, and other major abdominal surgery. This analysis revealed that the use of probiotics and synbiotics conferred a significant reduction in superficial surgical site infections and deep-space surgical infections [43]. The use of probiotics and synbiotics also reduced surgery-related complications, including pneumonia and urinary tract infections, and was associated with shorter length of stay in hospital. The authors concluded the effect of perioperative treatment with probiotics and synbiotics to be a complex one that was also associated with a reduction in post-surgical serum inflammatory markers, including C-reactive protein (CRP) and IL-6 [43].

Based on preclinical studies that suggest an altered microbiome plays a role in PJI development and clinical studies that have shown a reduction in SSI with correction of dysbiosis in other surgical fields, the microbiome's role in PJI is a promising and worthy area of further investigation [18,43]. The incidence of dysbiosis in patients with PJI needs to be evaluated. If found to play a significant role, treatments for dysbiosis such as probiotics or fecal transplant may prove effective in the prevention of PJI.

## Recommendations

We as arthroplasty surgeons ascribe to an ingrained dogma that all bacteria are bad. Without a doubt, impeccable sterility in the OR environment remains non-negotiable. However, we may need to broaden our thinking about microorganisms and their varying roles in PJI. The role of probiotics and synbiotics in the prevention of PJI warrants further investigation. The current notion that all bacteria should be annihilated with broad-spectrum antibiotics administered intravenously, topically, and in antibiotic cement may ultimately need to be abandoned for a more laser-focused approach, whereby antibiotics and other treatments that target specific pathogens while preserving symbiotic microbial species are developed and used. [40] The cornerstone of PJI prevention may ultimately involve coexisting with and even promoting “good” bacteria to fight the “bad” bacteria.

## Conflicts of interest

The authors declare that there are no conflicts of interest.

## References

- [1] Zmistowski B, Karam JA, Durinka JB, Casper DS, Parvizi J. Periprosthetic joint infection increases the risk of one-year mortality. *J Bone Joint Surg Am* 2013;95(24):2177.
- [2] Premkumar A, Kolin DA, Farley KX, et al. Projected economic burden of periprosthetic joint infection of the hip and knee in the United States. *J Arthroplasty* 2021;36(5):1484.
- [3] Sloan M, Premkumar A, Sheth NP. Projected volume of primary total joint arthroplasty in the U.S., 2014 to 2030. *J Bone Joint Surg Am* 2018;100(17):1455.
- [4] Chalmers BP, Matrkra AK, Sems SA, et al. Two-stage arthrodesis for complex, failed, infected total knee arthroplasty. *Bone Joint J* 2020;102-B(6\_Supple\_A):170.
- [5] Mahmoud SS, Sukeik M, Alazzawi S, Shaath M, Sabri O. Salvage procedures for management of prosthetic joint infection after hip and knee replacements. *Open Orthop J* 2016;10:600.
- [6] Osmon DR, Berbari EF, Berendt AR, et al. Diagnosis and management of prosthetic joint infection: clinical practice guidelines by the Infectious Diseases Society of America. *Clin Infect Dis* 2013;56(1):e1.
- [7] Alamanda VK, Springer BD. The prevention of infection: 12 modifiable risk factors. *Bone Joint J* 2019;101-B(1\_Supple\_A):3.
- [8] Lubbeke A, Zingg M, Vu D, et al. Body mass and weight thresholds for increased prosthetic joint infection rates after primary total joint arthroplasty. *Acta Orthop* 2016;87(2):132.
- [9] Gonzalez AI, Luime JJ, Uçkay I, et al. Is there an association between smoking status and prosthetic joint infection after primary total joint arthroplasty? *J Arthroplasty* 2018;33(7):2218.
- [10] Hadley S, Immerman I, Hutzler L, Slover J, Bosco J. Staphylococcus aureus decolonization protocol decreases surgical site infections for total joint replacement. *Arthritis* 2010;2010:924518.
- [11] Ribau AI, Collins JE, Chen AF, Sousa RJ. Is preoperative Staphylococcus aureus screening and decolonization effective at reducing surgical site infection in patients undergoing orthopedic surgery? A systematic review and meta-analysis with a special focus on elective total joint arthroplasty. *J Arthroplasty* 2021;36(2):752.
- [12] Wyles CC, Hevesi M, Osmon DR, et al. 2019 John Charnley Award: increased risk of prosthetic joint infection following primary total knee and hip arthroplasty with the use of alternative antibiotics to cefazolin: the value of allergy testing for antibiotic prophylaxis. *Bone Joint J* 2019;101-B(6\_Supple\_B):9.
- [13] Driesman A, Shen M, Feng JE, et al. Perioperative chlorhexidine gluconate wash during joint arthroplasty has equivalent periprosthetic joint infection rates in comparison to betadine wash. *J Arthroplasty* 2020;35(3):845.
- [14] Panahi P, Stroh M, Casper DS, Parvizi J, Austin MS. Operating room traffic is a major concern during total joint arthroplasty. *Clin Orthop Relat Res* 2012;470(10):2690.
- [15] Hooper GJ, Rothwell AG, Frampton C, Wyatt MC. Does the use of laminar flow and space suits reduce early deep infection after total hip and knee replacement?: the ten-year results of the New Zealand Joint Registry. *J Bone Joint Surg Br* 2011;93(1):85.
- [16] Gilbert JA, Blaser MJ, Caporaso JG, et al. Current understanding of the human microbiome. *Nat Med* 2018;24(4):392.
- [17] Turnbaugh PJ, Ley RE, Hamady M, et al. The human microbiome project. *Nature* 2007;449(7164):804.
- [18] Hernandez CJ, Yang X, Ji G, et al. Disruption of the gut microbiome increases the risk of periprosthetic joint infection in mice. *Clin Orthop Relat Res* 2019;477(11):2588.
- [19] Weingarden AR, Vaughn BP. Intestinal microbiota, fecal microbiota transplantation, and inflammatory bowel disease. *Gut Microbes* 2017;8(3):238.
- [20] Lee JT, Kim CM, Ramakrishnan V. Microbiome and disease in the upper airway. *Curr Opin Allergy Clin Immunol* 2019;19(1):1.
- [21] Chen X, Devaraj S. Gut microbiome in obesity, metabolic syndrome, and diabetes. *Curr Diab Rep* 2018;18(12):129.
- [22] Lee CJ, Sears CL, Maruthur N. Gut microbiome and its role in obesity and insulin resistance. *Ann N Y Acad Sci* 2020;1461(1):37.
- [23] Guss JD, Horsfield MW, Fontenele FF, et al. Alterations to the gut microbiome impair bone strength and tissue material properties. *J Bone Miner Res* 2017;32(6):1343.
- [24] Takiishi T, Fenero CIM, Camara NOS. Intestinal barrier and gut microbiota: shaping our immune responses throughout life. *Tissue Barriers* 2017;5(4):e1373208.
- [25] Geva-Zatorsky N, Sefik E, Kua L, et al. Mining the human gut microbiota for immunomodulatory organisms. *Cell* 2017;168(5):928.
- [26] Khosravi A, Yáñez A, Price JG, et al. Gut microbiota promote hematopoiesis to control bacterial infection. *Cell Host Microbe* 2014;15(3):374.
- [27] Taur Y, Jenq RR, Ubeda C, van den Brink M, Pamer EG. Role of intestinal microbiota in transplantation outcomes. *Best Pract Res Clin Haematol* 2015;28(2–3):155.
- [28] Taur Y, Pamer EG. Microbiome mediation of infections in the cancer setting. *Genome Med* 2016;8(1):40.
- [29] Lee GC, Ong K, Baykal D, Lau E, Malkani AL. Does prior bariatric surgery affect implant survivorship and complications following primary total hip arthroplasty/total knee arthroplasty? *J Arthroplasty* 2018;33(7):2070.

- [30] Blevins K, Aalirezaie A, Shohat N, Parvizi J. Malnutrition and the development of periprosthetic joint infection in patients undergoing primary elective total joint arthroplasty. *J Arthroplasty* 2018;33(9):2971.
- [31] Man SL, Chau WW, Chung KY, Ho KKW. Hypoalbuminemia and obesity class II are reliable predictors of peri-prosthetic joint infection in patient undergoing elective total knee arthroplasty. *Knee Surg Relat Res* 2020;32(1):21.
- [32] de Clercq NC, Groen AK, Romijn JA, Nieuwdorp M. Gut microbiota in obesity and undernutrition. *Adv Nutr* 2016;7(6):1080.
- [33] Hernandez CJ, Guss JD, Luna M, Goldring SR. Links between the microbiome and bone. *J Bone Miner Res* 2016;31(9):1638.
- [34] Corona PS, Lung M, Rodriguez-Pardo D, et al. Acute periprosthetic joint infection due to *Fusobacterium nucleatum* in a non-immunocompromised patient. Failure using a Debridement, Antibiotics + Implant retention approach. *Anaerobe* 2018;49:116.
- [35] Ehrlich GD, Hu FZ, Sotereanos N, et al. What role do periodontal pathogens play in osteoarthritis and periprosthetic joint infections of the knee? *J Appl Biomater Funct Mater* 2014;12(1):13.
- [36] Thomas M, Zeller V, Heym B, et al. *Gardnerella vaginalis*, from the vaginal microbiota to prosthetic joint infection. *J Bone Joint Infect* 2019;4(4):189.
- [37] Shi TB, Fang XY, Wang CX, et al. Rare occurrence of acute hematogenous periprosthetic joint infection due to *Fusobacterium nucleatum* in the background of a dental procedure: a case report. *Orthop Surg* 2020;12(6):2026.
- [38] Lagunes L, Rello J. Invasive candidiasis: from mycobiome to infection, therapy, and prevention. *Eur J Clin Microbiol Infect Dis* 2016;35(8):1221.
- [39] Song JH, Kim YS. Recurrent clostridium difficile infection: risk factors, treatment, and prevention. *Gut Liver* 2019;13(1):16.
- [40] Blaser MJ. Antibiotic use and its consequences for the normal microbiome. *Science* 2016;352(6285):544.
- [41] Hill C, Guarner F, Reid G, et al. Expert consensus document. The International Scientific Association for Probiotics and Prebiotics consensus statement on the scope and appropriate use of the term probiotic. *Nat Rev Gastroenterol Hepatol* 2014;11(8):506.
- [42] Jeppsson B, Mangell P, Thorlacius H. Use of probiotics as prophylaxis for postoperative infections. *Nutrients* 2011;3(5):604.
- [43] Skonieczna-Zydecka K, Kaczmarczyk M, Łoniewski I, et al. A systematic review, meta-analysis, and meta-regression evaluating the efficacy and mechanisms of action of probiotics and synbiotics in the prevention of surgical site infections and surgery-related complications. *J Clin Med* 2018;7(12):556.
- [44] Zaharuddin L, Mokhtar NM, Muhammad Nawawi KN, Raja Ali RA. A randomized double-blind placebo-controlled trial of probiotics in post-surgical colorectal cancer. *BMC Gastroenterol* 2019;19(1):131.