Case Report
Polymicrobial Bacteremia Involving Comamonas testosteroni

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Comamonas spp. are uncommon isolates in microbiology laboratories and have been rarely observed as an infectious agent in clinical practice. They have widespread environmental distribution and have been isolated from water, soil, and plants as well as from some hospital devices such as intravenous catheters and water contained in humidifier reservoirs used in respiratory treatment. The genus Comamonas originally contained the following species: acidovorans, testosteroni, kerstersii, terrigena, denitrificans, and nitrativorans. It now contains 17 species, while acidovorans spp. have been reclassified as Delftia acidovorans. In spite of its uncommon human pathogenesis, there are few reports on the aggressive manner of it as an opportunistic pathogen, mostly related to testosteroni spp. We present a case of polymicrobial bacteremia involving Comamonas testosteroni. The aim of this case report is to alert clinicians to the potential diagnosis of bloodstream infections caused by uncommon pathogens.

1. Introduction

Comamonas testosteroni, formerly known as Pseudomonas testosteroni, is an aerobic, motile, nonspore forming, ubiquitous Gram-negative organism. Although it has low virulence potency, there are few reports on its involvement in human infections. It became clinically important after 1987 when reports began accumulating on human infections such as cellulitis [1], peritonitis [2], endocarditis [3], meningitis [4], endophthalmitis [5], tenosynovitis [6], and pneumonia [7]. However, cases of bloodstream infections caused by Comamonas testosteroni have been infrequently reported [2, 8–12].

The name of testosteroni was given due to its characteristic of utilizing carbon from the metabolism of testosterone [13], although this property has also been demonstrated by some Pseudomonas spp. and fungi [14]. Some strains of Comamonas testosteroni have acquired the plasmid-mediated blaNDM1 gene that confers broad spectrum antimicrobial resistance thus potentially hampering treatment options in the event where this bacterium causes human infections [15].

In this report, we present an uncommon case of polymicrobial bacteremia involving Comamonas testosteroni with a detailed review of the literature.

2. Case Report

An 80-year-old African American female, morbidly obese, was brought to emergency department complaining of generalized body aches (predominantly in shoulders) and undocumented fevers at home for about 1 week. Her past medical history was significant for systemic arterial hypertension, diabetes mellitus, hiatal hernia, osteoarthritis, and cholelithiasis (s/p cholecystectomy). On arrival to the hospital, her vital signs were as follows: blood pressure of 150/77 mmHg, a regular heart rate of 106 beats/minute, a respiratory rate of 21 breaths/minute, a temperature of 100.8°F, and an oxygen saturation of 96% breathing ambient air.

Remarkable laboratory findings on admission were a white blood cell (WBC) count of 14, 500/mm3 (4.8–10.8), a sodium level of 129 mmol/L (135–147), a potassium level of 5.5 mmol/L (3.5–5.3), and a creatinine level of 2.0 mg/dL (0.8–2.0). C-reactive protein (CRP) and erythrocyte sedimentation rate (ESR) levels were both elevated at 152.7 mg/L (1.0–4.0) and 130 mm/h (0–20), respectively. Her urine analysis showed positive leukoesterase and a WBC count of 10–15/hpf (0–5).

Coagulation and liver profiles were within normal limits. Bilateral shoulder X-ray demonstrated degenerative changes,
3. Discussion

This is one of fewer than 15 cases of Comamonas testosteroni bacteremia reported in the literature and the first reported case of Comamonas testosteroni and Staphylococcus aureus polymicrobial bloodstream infection, according to a MEDLINE search using the words “Comamonas” and “bacteremia” and “bloodstream infection.” Approximately 35 cases of infections involving Comamonas testosteroni have been reported in the medical literature. Majority of infections are community-acquired rather than nosocomial [3, 5], and most of the patients reported had some degree of immuno-suppression such as malignancy, chronic liver disease, and end-stage renal disease requiring hemodialysis [1, 11, 12]. Bacterial translocation from the gastrointestinal tract seems to play an important role in the pathogenesis of infections caused by Comamonas spp. [9, 16].

Based on results from previous reports, majority of the 14 reported patients with bacteremia caused by Comamonas testosteroni were males, their median age was 47 years (range, newborn–89), only one patient died, and predisposing factors for infection were not identified in two of them. Most of the patients responded well to appropriate antimicrobial therapy [12]. Comamonas spp. isolates are usually susceptible to aminoglycosides, fluoroquinolones, carbapenems, piperacillin-tazobactam, cephalosporins, and trimethoprim-sulfamethoxazole [1, 4].

The patient discussed in this report did not have any obvious source of infection. She did not have central venous catheters, and the abdominal imaging did not evidence any acute inflammatory process or mass. Her immunosuppressive state was limited to diabetes mellitus, which might have played a role as a possible predisposing factor. Therefore, the authors hypothesized that the most likely source might have been the right shoulder rotator cuff tendinitis. Although this hypothesis cannot be proven, it is supported by the fact that the patient reported shoulder pain, the CRP and ESR levels were markedly elevated, and the right shoulder imaging showed rotator cuff tendinitis. In 2000, Isotalo et al. reported a case of polymicrobial tenosynovitis, involving an organism most likely to be related to Comamonas spp. based on biochemical factors [6]. If proven, our patient may have been only the second case of tenosynovitis caused by Comamonas spp. reported in the literature. We also postulate that the second likely mode of bacteremia in this case could have been dissemination of Comamonas testosteroni from the bowel following gastrointestinal translocation, given the finding of extensive diverticulosis demonstrated in the abdominal imaging.

Polymicrobial bacteremia involving Comamonas testosteroni have been reported in association with Streptococcus parasanguis and Ralstonia pickettii [17]. Two other cases of polymicrobial bacteremia involving Comamonas spp. (Delftia acidovorans (formerly known as Comamonas acidovorans) and Comamonas kerstersii) were previously reported in association with Streptococcus agalactiae and Bacteroides fragilis, respectively [17]. To our knowledge, this is the first case of bloodstream infection involving Comamonas testosteroni and Staphylococcus aureus.

In this report, we have highlighted an unusual cause of bloodstream infection. Comamonas spp. should be kept in mind as a rare cause of bacteremia.

Conflict of Interests

The authors report no conflict of interests regarding the publication of this paper.

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