Outbreak of *Pseudomonas Oryzihabitans*

Pseudobacteremia Related to Contaminated Equipment in an Emergency Room of a Tertiary Hospital in Korea

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*Pseudomonas ozyihabitans* is frequently found in various sites within hospital settings, including sink drains and respiratory therapy equipment. Although it rarely causes human infections, *P. ozyihabitans* has recently been considered a potential nosocomial pathogen, especially in immunocompromised hosts. We report our experience of an outbreak of *P. ozyihabitans* pseudobacteremia, presumably due to faulty aseptic preparation of a saline gauze canister.

**Key Words:** *Flavimonas ozyihabitans*; Pseudobacteremia; Environment contamination
Between October 7 and October 11, 2011, 8 patients visited the emergency room of a tertiary hospital and were diagnosed with blood cultures positive for *P. oryzihabitans*. When these patients visited our emergency room (ER) complaining of febrile sensations in spite of ambiguous clinical signs and symptoms, blood cultures were ordered to test for systemic infections. Antimicrobial susceptibility testing was performed using standard techniques. Blood samples were placed in culture bottles (Vital AER or Vital ANA, bioMérieux; Marcy-l’Etoile, France) and incubated for 24–48 h; *P. oryzihabitans* was identified in 10 samples from 8 patients using a Vitek system (bioMérieux). Antimicrobial susceptibility testing was also performed using the same system. All but 2 patients had a single positive blood culture result, and no microorganism was isolated from other sites. The characteristics of the affected patients, including demographic data, medical history, and presence of a central venous catheter, are shown in Table 1. None of the patients with blood cultures positive for *P. oryzihabitans* showed clinical signs of sepsis; this cluster of cultures positive for an unusual pathogen was suggestive of a pseudoepidemic. Review of microbiology laboratory records from the previous 12 months found no instances of *P. oryzihabitans*. During the same period, *P. oryzihabitans* was not isolated from clinical specimens from other wards. The ER environment was investigated to determine potential sources of infection and to implement control measures to stop the outbreak. Bacteriological cultures of ER environmental objects were performed.

A total of 22 environmental samples from the ER were cultured to test for a *P. oryzihabitans* cluster. Sites examined included the disinfectant, forceps, saline gauze cans, and gloves used for dressing wounds as well as hand-washing soap, tap water, tables, beds, chairs, and linens in the ER.

Among these environmental samples, only 1, from a saline gauze canister, was culture-positive for *P. oryzihabitans*. This culture result was compared to results of 16S rRNA sequencing of the same sample. Sequencing also revealed *P. oryzihabitans*. Antimicrobial susceptibility tests of the organism isolated from the saline gauze canister showed the same pattern as the patient samples. All isolates were sensitive to imipenem, gentamicin, penicillin, ciprofloxacin, ceftazidime, ampicillin, amoxicillin/clavulanic acid, trimethoprim/sulfamethoxazole, cefotaxime, and ceftazidime. All isolates were sensitive even to ampicillin, to which some *Pseudomonas* species have a known endogenous resistance. Saline-soaked cotton gauzes stored in canisters were used in our hospital to remove dirt from intact skin. However, these gauzes could have been used to clean disinfectant from the skin for access to veins and mistaken for cotton materials soaked in 70% ethanol, which were stored in similar canisters and were of the same color. Consequently, careful investigation of the above results revealed the outbreak to be pseudobacteremia; the most likely etiology was environmental contamination of the saline gauze can, which contaminated several blood cultures during collection of specimen.

After the outbreak, the infection control unit instituted infection control measures including removal of saline gauze cans from the ward, including the ER. After instituting control measures and monitoring blood culture results daily, no additional outbreaks have been documented in the ER. Therefore, we could conclude that we had found the focus of the *P. oryzihabitans* contamination and pseudobacteremia outbreak and that the outbreak was over.

Several reports have shown evidence of pseudobacteremia outbreaks caused by *P. oryzihabitans*. Eom et al. [11] reported an outbreak of *P. oryzihabitans* pseudobacteremia, presumably arising from faulty aseptic preparation of a blood culture set. In that report, genotypically identical isolates were identified based on pulsed-field gel electrophoresis (PFGE) patterns of 4 *P. oryzihabitans* isolates from patients with no signs or symptoms consistent with *P. oryzihabitans* bacteremia. Kim et al. [12] reported *Enterobacter nimipressuralis* pseudobacteremia from the use of contaminated saline-soaked cotton at the sites of venipuncture blood draw. Because *Enterobacter* clinical isolates could not be identified at the species level by 16S rRNA sequencing, the organism was accurately identified by sequencing of the hsp60 gene and infrequent-restriction

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**Table 1. Patients demographic data**

| Variables                      | Patients (N = 9) |
|-------------------------------|-----------------|
| Sex (female/male)             | 4/5             |
| Age (years)                   | 46.2 (2–79)     |
| Body temperature (°C)         | 36.7 (36–37.7)  |
| Medical condition             | Gallbladder empyema, Colon cancer, Plasma cell myeloma, Asthma, Febrile seizure, Myocardial infarction, Urethral injury, Colon cancer, Meningoencephalitis<sup>b</sup> |
| Presence of central venous catheter | No |

<sup>a</sup> Data are given as median (range).

<sup>b</sup> Meningoencephalitis may be caused by Klebsiella pneumoniae.
site polymerase chain reaction (IRS-PCR). In most reported instances of pseudobacteremia outbreaks, automated culture systems and molecular typing were used to find that hospital environmental contaminants, such as coagulase-negative cocci, Bacillus spp., Corynebacterium spp., Micrococcus spp., Propionibacterium spp., gram-positive bacilli, or Clostridium perfringens were the most common causes of outbreaks [13, 14]. Recently, Neulier et al. [15] reported a pseudo-outbreak of Pseudomonas putida respiratory infections. Investigation of the microbiology laboratory where 5 samples were processed by the same device found that P. putida isolates were obtained from a reusable container in this device. All isolates presented identical antibiotic susceptibility patterns; therefore, the authors declared this pseudo-outbreak of P. putida to be over without additional testing such as random amplified polymorphic DNA (RAPD) typing or PFGE. Similarly, we did not investigate the clonality of the environmental isolates in this study by 16S rRNA sequencing, RAPD typing, or PFGE. Although the clinical significance of the P. oryzihabitans outbreak presented here remains unclear, P. oryzihabitans is an infrequent cause of infection and was isolated from patients with no clinical signs of sepsis. In addition, antimicrobial susceptibility test results were the same for P. oryzihabitans isolated from both patients and the contamination source. We suggest that P. oryzihabitans from a contaminated saline gauze canister was an etiology of pseudobacteremia outbreak in our ER; however, molecular typing of the environmental and patient samples was not performed.

In conclusion, we found the focus of P. oryzihabitans contamination to be a saline gauze canister and could declare the pseudobacteremia outbreak of P. oryzihabitans to be over after daily monitoring of blood culture results. Our results confirm previous reports that pseudobacteremia is usually caused by hospital environmental contaminants and emphasize the importance of hospital surveillance to reduce outbreaks and protect other patients at risk.

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