

**Microbial contamination of water sources for high-risk patients of a Pennsylvanian tertiary
care facility**

by

Akaash Ajay Patel

B.S. in Biotechnology, James Madison University, 2020

Submitted to the Graduate Faculty of the
Infectious Disease and Microbiology Department
Graduate School of Public Health in the partial fulfillment
of the requirements for the degree of
Master of Public Health

University of Pittsburgh

2021

UNIVERSITY OF PITTSBURGH
GRADUATE SCHOOL OF PUBLIC HEALTH

This essay is submitted

by

Akaash Ajay Patel

It was defended on

December 6th, 2021

and approved by

Committee Chair: Dr. Toan Ha, MD, PhD, Assistant Professor, Department of Infectious Diseases and Microbiology, University of Pittsburgh

Committee Chair: Dr. Ken Ho, MD, PhD, Clinical Professor, Department of Medicine, School of Medicine, University of Pittsburgh

Committee Chair: Dr. Ernesto T. A. Marques Jr., MD, PhD, Associate Professor, Infectious Diseases and Microbiology, University of Pittsburgh

Thesis Advisor: Mohamed Yassin, MD, PhD, Clinical Assistant Professor, Department of Medicine, School of Medicine, University of Pittsburgh

Copyright © by Akaash Ajay Patel

2021

Microbial contamination of water sources for high-risk patients of a Pennsylvanian tertiary care facility

Akaash Ajay Patel, MPH

University of Pittsburgh, 2021

Abstract

The present study aims to investigate the extent of microbial contamination of hospital water sources in UPMC Mercy hospital, a Pennsylvanian tertiary care facility through environmental sampling and microbiological culturing. Following interpretation of results, assess the hospital's use of mechanical filtration on unit floors and at distal points then provide recommendations for key stakeholders of the hospital so that actionable steps can be taken. Forty-five water samples were collected from the 8th floor general ICU, 4th floor trauma burn unit (TBU) ICU, 3rd floor medical surgical ICU, 4th floor unused ICU, 2nd floor endoscopy unit, and 5th floor hemodialysis unit. Water sources include filtered and unfiltered sources such as sinks, showers, hemodialysis boxes, nurses' water stations, and unit filtration systems. 35.6% (16/45) of water samples were positive for microbial growth as general ICU, medical-surgical ICU, and unused ICU were the main sources of the contamination. Patient sinks, showers, and hemodialysis boxes represent distal sites on the 8th floor general ICU that were contaminated even though the unit's double filtration system was negative for the water collected post filtration. In contrast, patient sinks and patient showers equipped with PALL filters on the 4th floor TBU ICU and hydrotherapy room showed no signs of microbial contamination. The results show that application of a double water filtration system on unit floors does reduce microbial contamination in the hospital water but has shown to not be effective in clearing microbial

contamination at distal sites of the water system. The preceding findings were presented to key stakeholders of the hospital resulting in infection preventionist led education for HCW staff on water flushing, a collaboration between engineers and hemodialysis unit to fit hemodialysis boxes with a hose connector to flush water, and whole genome sequencing of bacterial positive cultures. More research is needed to understand the extent of water-related HAIs to help create a comprehensive system that includes a water management program, preventative practices, and intervention strategies for water-related HAIs as they pose a substantial risk for patients in high risk settings.

Table of Contents

1.0 Introduction.....	1
1.1 Water source quality in tertiary care facilities	2
1.2 Understanding patient risks to water related HAIs	4
1.2.1 Increased HAI acquisition for patients in high risk settings.....	4
1.2.2 Antibiotic resistance of water pathogens	4
1.2.3 Water associated outbreaks in tertiary care facilities	5
1.3 Current standards for hospital water management.....	7
1.4 Specific aims.....	9
2.0 Methods.....	10
2.1 Research approach	10
2.2 Sample collection	11
2.3 Microbiological culturing and analysis	12
3.0 Results	13
3.1 Extent of positive water cultures in UPMC Mercy	13
4.0 Discussion.....	15
5.0 Conclusion and recommendations.....	18
Bibliography	19

List of Tables

Table 1 Sample collection schema for hospital water samples	11
Table 2 Distribution and frequency of positive water cultures across ICU units and hospital	13

1.0 Introduction

Healthcare-associated infections (HAIs) are infections that are acquired while receiving health care and appear 48 or more hours following admission into a hospital (CDC, 2021a). HAIs persist as a significant contributor to patient morbidity and mortality resulting in the direct medical costs of more than \$34 billion dollars each year to the U.S. hospital system (Scott et al., 2019). The hospital environment is a known reservoir for opportunistic pathogens with evidence of microbial contamination seen on high-touch patient surfaces, medical devices, and in hospital water systems (Facciola et al., 2019; Hayward et al., 2020; Russotto et al., 2015). Unnoticed microbial contamination in the hospital environment leads to prolonged HAI acquisition among susceptible patients found in high-risk settings such as intensive care units (ICU) (Eyre et al., 2018; Weber et al., 2010). The responsibility to reduce the microbial burden is centered on the relationship between healthcare workers (HCWs) and infection preventionist (IP), fostered by excellent interpersonal communication. The IP is charged with routine surveillance of microorganisms in the environment and to provide education on best practices for nurses and clinicians as well as education on the employment of evidence-based disinfection practices (Chirca, 2019). The burden then largely falls on HCWs to be receptive and committed to the conceived system and in providing patient-centered care.

The role of high-touch patient surfaces and medical devices as a reservoir and in the transmission of opportunistic pathogens is well characterized but is not well understood for pathogens residing in the hospital water (Hayward et al., 2020). Hospital water pathogens can be responsible for direct transmission events or serve as a selective driver of antibiotic resistance in gram-negative organisms (Weinbren, 2020). In a summary of internal CDC records of HAIs

between 2014 and 2017, researchers estimated 134 (21.6%) patient consultations involved water-related HAI events, implicating the substantial contribution of water-related HAIs to the overall growing problem of HAIs (Perkins et al., 2019).

1.1 Water source quality in tertiary care facilities

Commonly overlooked, hospital water distribution systems can be a potent source of opportunistic microorganisms in a hospital environment presenting a continual challenge for HCWs. Researchers frequently identified the opportunistic microorganisms as gram-negative bacterial species such as *Legionella spp.*, *Pseudomonas spp.*, and carbapenem-resistant *Enterobacteriaceae spp.* as well as gram-positive non-tuberculosis *Mycobacterium spp.* (D'Alessandro et al., 2015; Jamal et al., 2020; Millar & Moore, 2020; Montanari et al., 2009; Nakamura et al., 2020; Tang et al., 2020; Ziwa et al., 2019). All species represent opportunistic pathogens commonly seen in nosocomial environments.

Initially the municipal water supply that flows into the hospital water distribution system harbors opportunistic microorganisms as tertiary hospitals have shown that existent microbial contamination of the municipal water has led to persistent contamination of the hospital water distribution system when left untreated (D'Alessandro et al., 2015; Decker & Palmore, 2014; Vickers et al., 1987). Hospitals then use a process of chemical or mechanical treatment to reduce microbial water contamination (I Marchesi et al., 2020; I Marchesi et al., 2011).

Hospital water distribution systems can be contaminated in its proximal infrastructure being the central pipes supplying the water and at distal water outlets being point of use sites such as handwashing stations (Decker & Palmore, 2014). The water in the proximal

infrastructure faces structural malignancies that create areas of stagnant water and differential heat, presenting ideal conditions for microbial growth of *Legionella spp.* and biofilm formation of *Pseudomonas spp.*(Chirca, 2019; Decker & Palmore, 2014). Resolving structural issues are costly and require administrative initiative but are feasible long-term solutions to microbiological contamination seen in the hospital water. The frequency of distal water outlets in a hospital presents a unique challenge in that its infrequent use or poor maintenance result in an opportunity for microbial colonization. Evidence of contamination is seen in patient sinks and showers that can go unused across different intensive care units (ICUs) as they are not necessary in the daily provision of care (Jamal et al., 2020; Montanari et al., 2009; Tang et al., 2020; Ziwa et al., 2019). Distal sites then serve as a risk for patient microbial colonization given their proximity to the patient and improper management. Other distal sites include nurse water stations and ice machines that have been colonized by non-tuberculous mycobacteria despite disinfective practices (Millar and Moore 2020). In the provision of care, water utilizing medical devices such as hemodialysis machines and hydrotherapy baths have been shown to be colonized by opportunistic pathogens in the dialysis filtrate and drains, respectively (Montanari et al., 2009; Tang et al., 2020; Ziwa et al., 2019). The significance of existent microbial contamination in the hospital water distribution system is the potential for patient colonization or infection through direct and indirect pathways offering multiple modes of transmission for the water pathogens.

1.2 Understanding patient risks to water related HAIs

1.2.1 Increased HAI acquisition for patients in high risk settings

In the past year, the CDC annual progress report revealed an increase in CLABSI, CAUTI, VAE, and MRSA bacteremia compared to 2019 (Monegro et al., 2021). The increasing incidence of HAIs challenges the current provision of care to susceptible patients seeking care in adult ICUs. Patient demographics established before admission to an ICU create an opportunity to estimate patient vulnerability to HAIs. Univariate risk factor analyses demonstrate that patient age, immunosuppression status, history of antibiotic use, diagnosis of cardiovascular disease, and diagnosis of diabetes mellitus implicate a higher risk of an HAI for patients (Despotovic et al., 2020; Rodríguez-Acelas et al., 2017). Following admission into an ICU, the care the patient receives can implicate higher risks for HAI acquisition being the duration of stay in the ICU, use of mechanical ventilation, intubation of central lines, and reoperation (Despotovic et al., 2020; Rodríguez-Acelas et al., 2017). Considering both pre and post admission risk factors for HAI acquisition in adult ICUs can help create a conceptual framework for understanding the risks that individual patients face.

1.2.2 Antibiotic resistance of water pathogens

The hospital environment is a driver of antimicrobial resistance with clinical practices of over-prescribing antibiotics and inadequate sanitation and infection control practices being the selective promoters (McEwen & Collignon, 2018). The sequelae is the increasing presence of antibiotic resistant organisms that complicate patient treatment regimens and lead to more severe

outcomes during a patient infection. To combat antibiotic resistance, hospitals have adopted antimicrobial stewardship programs that demonstrated substantial benefits in clinical and economic outcomes in the past twenty years (Nathwani et al., 2019). The practices, although effective, exclude the presence of antibiotic resistant organisms in hospital water sources as a targetable intervention (Hayward et al., 2020).

A recent systematic review summarized water-associated HAI outbreaks in the U.S. in which researchers determined antibiotic resistant waterborne pathogens *P. aeruginosa*, *Mycobacterium spp.*, *Legionella spp.*, *S. aureus*, and carbapenem resistant Enterobacteriaceae (CRE) as the responsible agents (Hayward et al., 2020). Even with evidence of waterborne multidrug resistant organisms (MDROs), hospitals neglect the water sources in favor of supported antimicrobial stewardship programs. Theoretically, gram-negative pathogens can spread antibiotic resistance through mobile genetic elements in contaminated hospital water that could cause outbreaks of MDROs such as CRE. Interactions involving the spread of antibiotic resistance in hospital water sources are seen in the wastewater of healthcare facilities that show a significant number of antibiotic resistant water-borne pathogens in comparison to municipal water (Eda et al., 2021; Hassoun-Kheir et al., 2020). The interactions between the gram-negative bacteria could be a factor in increasing MDRO spread but there is no literature to support this.

1.2.3 Water associated outbreaks in tertiary care facilities

Gram-negative bacteria in water are associated with multiple reported infections in high risk patient settings of tertiary care facilities. Patient infections commonly occur in clusters from aerosol transmission from the water source, water splash events, exposure of implanted devices to water such as central venous catheters, contaminated water in medical devices, and from

HCW hands (Decker & Palmore, 2014; Vincenti et al., 2014; Yassin et al., 2020). At the National Institute of Health (NIH) there were multiple cases of *Sphingomonas* bacteremia and sepsis related to faucet water splash and contamination of medical and surgical equipment in an interdisciplinary ICU (Johnson et al., 2018). Investigators were only able to establish a genetic similarity between the hospital water reservoir of *Sphingomonas* and clinical isolates of *Sphingomonas* (Johnson et al., 2018). Continual transmission from water source to patient occurs and was seen in a German interdisciplinary ICU in which prolonged infections with multidrug resistant *P.aeruginosa* was the result of HCWs emptying ultra-filtrate bags used during hemofiltration in patient sinks leading to person to person transmission (Salm et al., 2016). There are numerous cases of bacteremia, endocarditis, and sepsis related to waterborne pathogens that follow the contamination of wounds, injection drug use, and contamination of dialysis access with faucet water (CDC, 1998; Chotikanatis et al., 2011).

Biofilm forming water pathogens are known to cause outbreaks or “pseudo-outbreaks” in dialysis centers and in patient ventilators or heart lung machines (Montanari et al., 2009). The formation of the biofilm leads to persistent colonization of pathogens that are difficult to treat, often requiring replacement of the biofilm contaminated source (Donlan & Costerton, 2002). In a Pittsburgh tertiary care facility there is an ongoing, large pseudo-outbreak of *Delftia acidovorans* associated with biofilm contamination of the reverse osmosis water in the hemodialysis machines (Yassin et al., 2020). *D. acidovorans* is a waterborne pathogen residing in the Pittsburgh municipal water and can infect susceptible patients directly. *P.aeruginosa*, another opportunistic pathogen known for its biofilm formation, was also found in the reverse osmosis water of hemodialysis machines at a different tertiary care facility, exhibiting high resistance rates to antibiotics (Vincenti et al., 2014). Both investigations into the pseudo-outbreaks recognized the

replacement of the reverse osmosis filters and hemodialysis machines as the cost-effective method for eliminating the contamination (Vincenti et al., 2014; Yassin et al., 2020).

The outbreaks highlight that microbial contamination of the hospital water system in its proximal or distal infrastructure can lead to patient infections directly or indirectly in adult ICUs. Current practices in patient treatment that involve an exposure to contaminated water may result in a water-related outbreak that necessitates an immediate intervention for the patient and causative source.

1.3 Current standards for hospital water management

The Joint Commission and CDC oversee water management programs across the U.S. hospital system. The CDC provides guidelines in the form of toolkits to create water management programs in which The Joint Commission sets and enforces standards for hospitals and water management programs (CDC, 2021b; The Joint Commission, 2021b). A new standard issued by The Joint Commission, Standard EC 02.05.02 goes into effect January 1st, 2022 and requires hospitals, critical care facilities, and nursing care centers to create a water management program that addresses *Legionella* and other water pathogens (The Joint Commission, 2021a).

The main purpose of the new standards is to protect patient and HCW health using a systems approach to water treatment. Previous standards were broad and lack specific guidance on targeted water treatment strategies, creating variation in water treatment strategies in U.S. hospitals with different tertiary care facilities exploring novel solutions for microbiological contamination within their hospital water distribution systems (CDC, 2021b; The Joint Commission, 2021a). Currently tertiary care facilities flush distal outlets for at least 5 minutes at

65°C called superheating to decontaminate sites with *L. pneumophila*, but this straightforward strategy has proven ineffective in eliminating long-term contamination (Chen et al., 2005). Longitudinal studies of different control strategies for *L. pneumophila* in tertiary care facilities identified hyperchlorination (>10mg/L of free chlorine at point of delivery), the use of electric boilers, and point of use filters as the cheapest, most-effective strategies for reducing systemic contamination of *L. pneumophila* in the hospital water system (Isabella Marchesi et al., 2020; I Marchesi et al., 2011). Although only applied to *L. pneumophila*, these water treatment strategies can reduce the microbial burden of other pertinent bacterial species.

UPMC Mercy, a tertiary care facility in Pittsburgh, Pennsylvania uses a two-pronged approach of systemic chemical treatment and mechanical filtration to reduce the microbial burden in the hospital water distribution system. The water flowing through the proximal infrastructure is treated with monochloramine, successfully eradicating *Legionella* from their water system (Kandiah et al., 2013). High-risk patient settings including the general ICU, trauma burn ICU, hydrotherapy rooms, and hemodialysis units have a double filtration system on the floor to supply filtered water into distal sites. Theoretically, the double filtration system works to further reduce the microbial burden of patient accessible water. The trauma burn ICU equipped all patient sink and shower heads with PALL filters to prevent continued colonization at distal water sites (Loveday et al., 2014). Oversight of the water distribution system involves IP, nurses, and cleaning staff in maintaining distal water sources through environmental rounding or employing point of use guidance such as letting patient showers run before use. Even with the establishment of a complex water management program, evidence of microbiological contamination of hospital water continues to emerge within the hospital.

1.4 Specific aims

1. Describe the extent of positive water cultures in high-risk patient environments at a tertiary care hospital through environmental sampling and microbiological culturing
2. Assess the hospital's use of mechanical filtration on unit floors and at distal points.
3. Provide recommendations and implication for the key stakeholders and hospital to take actionable steps

2.0 Methods

2.1 Research approach

The cross-sectional study was conducted using environmental water samples collected during the month of October 2021 in a tertiary care hospital in Pittsburgh, Pennsylvania. Forty-five water samples were collected from filtered and unfiltered water sources in the hospital. The sample distribution is as follows: nineteen samples from 8th floor general ICU, seven samples from 4th floor trauma burn unit (TBU) ICU, four samples from 4th floor unused ICU, four samples from the 2nd floor endoscopy unit, seven samples from the 3rd floor medical surgical ICU, and four samples from the 5th floor hemodialysis unit.

The maintenance team collected environmental water samples from the hemodialysis unit and floor unit water filtration systems. The distribution of collected water sources and origins is outlined in **Table 1**.

Table 1 Sample collection schema for hospital water samples

Patient Unit	Sample Source	Samples Collected	Collective Total
General ICU 8th Floor			19
Sinks	Patient Room	5	
Showers	Patient Room	5	
Hemodialysis Boxes	Patient Room	5	
Hemodialysis Pre-Post Filter	Floor Unit	4	
TBU ICU 4th Floor			26
Sinks	Patient Room	3	
Nurse water station	Floor Unit	2	
Pre-Post filter hydrotherapy room	Floor Unit	2	
MS ICU 3rd Floor			33
Sinks	Patient Room	4	
Nurse water station	Floor Unit	2	
Nurse sink	Floor Unit	1	
Endoscopy Unit 2nd Floor			37
Sinks	Patient Room	4	
Unused General ICU 4th Floor			41
Sinks	Patient Room	4	
Hemodialysis Unit 5th Floor			45
Hemodialysis Sink	Floor Unit	1	
Hemodialysis Unit	Floor Unit	1	
Hemodialysis Pre-Post Filter	Floor Unit	2	

2.2 Sample collection

Water was collected from unit sinks, showers, nurse water stations, hemodialysis boxes, and unit filtration systems. The water was allowed to run for 10 seconds, and then 100mL of water was collected into sterile 100mL collection tubes. The water was stored at 37°C and within 48 hours samples were cultured.

2.3 Microbiological culturing and analysis

100mL of the water sample was vacuum filtered using 0.2-micron Nalgene membranes. The Nalgene membrane was then aseptically transferred to a TSA plate using sterile tweezers. TSA plates were incubated at 36°C for 72 hours. Following incubation, results of bacterial growth were determined and recorded. Positive bacterial growth was recorded as <10 colonies, 10 to 100 colonies, and too numerous to count (TNTC).

All positive TSA cultures were subsequently cultured on MacConkey agar and Blood agar plates. Single colonies from positive TSA cultures were picked and streaked onto MacConkey and Blood Agar plates. MacConkey and Blood agar plates were incubated at 36°C for 48 hours. Following incubation, results of bacterial growth were determined and recorded. Positive cultures represent bacteria that show countable growth on TSA and Blood agar plates.

3.0 Results

3.1 Extent of positive water cultures in UPMC Mercy

Table 1 displays the distribution of forty-five environmental water samples collected at distal and proximal sites of the UPMC Mercy water distribution system (**Table 1**).

Table 2 Distribution and frequency of positive water cultures across ICU units and hospital

Unit Type	Positive Samples (%)
ICU Units	16/37 (41.7)
Sinks and showers	13/22 (59.1)
Hemodialysis Boxes	5/5 (100)
Nurse water stations	2/4 (50)
Unit pre-post filters	1/6 (16.7)
Endoscopy Unit 2 nd Floor	
Sinks	0/4 (0)
Hemodialysis Unit	
Sink	0/1 (0)
Units	0/3 (0)
Filtered vs. Unfiltered water	16/45 (35.6)
Filtered	8/31 (25.8)
Unfiltered	8/14 (57.1)

The results show that 35.6% (16/45) of the water samples were positive for microbial growth (**Table 2**). Distal sites in the 8th floor general ICU and 3rd floor medical surgical ICU represent the main source of microbial contamination of the sampled water sources being that 59.1% of sinks and showers, 100% of hemodialysis boxes, and both the ice and water of the nurse water station were positive for microbial growth (**Table 2**). Proximal sites of the adult ICUs include the unit double filtration systems located on the 8th floor general ICU and 4th floor TBU ICU. Both filtration units showed no evidence of microbial contamination in the filtered water although the pre-filtered water collected from the 8th floor general ICU filtration system was positive for microbial growth. ICU units that do not have a unit filtration system include the 3rd floor medical-surgical ICU and the 4th floor unused ICU. In comparison of filtered and unfiltered sources of water, 25.8% of filtered water sources were contaminated and 57.1% of unfiltered water sources were contaminated (**Table 2**).

4.0 Discussion

The present study shows evidence for microbial contamination within the hospital water system in high-risk patient settings. In comparison of filtered and unfiltered water sources, 25.8% (8/31) of filtered and 57.1% (8/14) of unfiltered water sources were contaminated. The main sources of microbial contamination of filtered water sources are seen at distal sites of the 8th floor general ICU as 61.5% (8/13) of sinks, showers, and hemodialysis boxes sampled from patient rooms were positive for microbial growth. The proximal infrastructure of the water system being the 8th floor double filtration system shows evidence of contamination prior to being filtered but the water collected post filtration was negative for microbial contamination.

The unexplained distal contamination seen on the 8th floor general ICU can potentially be attributed to the local water distribution system on the floor that supplies the water from the double filtration system to the patient rooms. PALL filters used in patient sinks and showers as well as the hydrotherapy room on the 3rd floor TBU ICU showed no signs of microbial contamination, demonstrating their effectiveness as a point of use filtration system in comparison to unit filtration systems. 80% (8/10) of patient sinks on the 3rd floor medical-surgical ICU and the 4th floor unused ICU were contaminated. Both units do not have a double filtration system so there was an expectation of microbial contamination, given these are unfiltered water sources. Additionally, the nurses' water station of the 3rd floor medical-surgical ICU was contaminated in both the water and ice it provided that might need to be replaced if standard disinfection practices do work.

The application of a double water filtration system on unit floors does reduce microbial contamination in the hospital water but has shown to not be effective in clearing microbial

contamination at distal sites of the water system. Particularly, the 8th floor general ICU and 3rd floor medical-surgical ICU shows evidence of widespread contamination of patient accessible water sources implicating a higher risk for patient transmissible events. Although no direct transmission events were observed in both units, those water sources serve as potential outbreaks for the ICU units as this is seen in the extended outbreak *K. oxytoca* in a medical-surgical ICU in which outbreak associated clones of *K. oxytoca* were ubiquitous in patient sinks of the ICU (Lowe et al., 2012).

The preceding findings and conclusions were presented to relevant stakeholders at UPMC Mercy including the director of critical care, the lead engineer, infectious disease clinicians and IPs. The discussion resulted in actionable steps that are as follows. The IP personnel will provide education to nursing staff located on the 8th floor general ICU to routinely flush and flush the water before use for all distal water sources located in patient rooms. The hemodialysis boxes within patient rooms are locked and given that they are meant to connect to a direct water line there's no way to flush the water without flooding the patient room. To address this, the hemodialysis unit and engineers will work to unlock all hemodialysis boxes and to fit each unit with a hose connector that allows the water to be easily flushed and accessible to the nursing staff. Lastly, the water samples that were successfully cultured will be sent to Phigenics, an independent water management company, to be sequenced using whole genome sequencing to identify the microorganisms present in the UPMC Mercy water.

General limitations of the study involve the research approach being that sample size was small. Collecting more water samples from unit floors, particularly the 8th floor general ICU and 3rd floor medical-surgical ICU, would paint a more accurate picture of the distribution and frequency of microbial contamination of hospital water. During sample collection, water was

only flushed for ten seconds prior to being collected, which does not adhere to the actual practice of water flushing that can last more than a minute. At the time of collection this seemed impractical, as collecting water from hemodialysis boxes would cause flooding of the patient room if it was flushed for more than ten seconds so to maintain consistency, all water sources were flushed for ten seconds. The simple error in approach could contribute to the high positivity of water cultures collected from patient showers and hemodialysis boxes on the 8th floor general ICU as these water sources could have gone unused for weeks, in which there could be a buildup of microbial contamination that needs to be flushed.

5.0 Conclusion and recommendations

The central limitation of the study is that it cannot address, if microbial contamination at distal water sites in high-risk patient settings be identified as a targetable intervention? Simply proposing there is microbial contamination at distal water sites can lead to costly and time-consuming interventions that result in no clear benefit to the patient. Instead, determining that there is persistent microbial contamination of the water source within the patient unit would be sufficient rationale for an intervention to prevent patient colonization and infections. Targeted interventions can be used such as the employment of point of use filters at distal water sites, which is a proven effective measure to reduce endemic infections and colonization of patients in high-risk settings (Cervia et al., 2010; Trautmann et al., 2008).

If continued microbial colonization is seen in patient water sources on the 8th floor general ICU and 3rd floor medical surgical ICU, PALL filters should be considered as a final means of eliminating microbial contamination without having to replace the distal infrastructure. Overall more research is needed to understand the extent of water-related HAIs to help create a comprehensive system that includes a water management program, preventative practices, and intervention strategies for water-related HAIs as they pose a substantial risk for patients in high risk settings.

Bibliography

- CDC. (1998, January 30). *Outbreaks of Gram-Negative Bacterial Bloodstream Infections Traced to Probable Contamination of Hemodialysis Machines -- Canada, 1995 United States, 1997; and Israel, 1997*. Outbreaks of Gram-Negative Bacterial Bloodstream Infections Traced to Probable Contamination of Hemodialysis Machines -- Canada, 1995 United States, 1997; and Israel, 1997.
<https://www.cdc.gov/mmwr/preview/mmwrhtml/00051244.htm>
- CDC. (2021a). *The NHSN Patient Safety Component Manual* .
https://www.cdc.gov/nhsn/pdfs/pscmanual/pcsmanual_current.pdf
- CDC. (2021b, March 25). *Legionella Water Management Programs Overview* / CDC.
<https://www.cdc.gov/legionella/wmp/overview.html>
- Cervia, J. S., Farber, B., Armellino, D., Klocke, J., Bayer, R. L., McAlister, M., Stanchfield, I., Canonica, F. P., & Ortolano, G. A. (2010). Point-of-use water filtration reduces healthcare-associated infections in bone marrow transplant recipients. *Transplant Infectious Disease*, 12(3), 238–241. <https://doi.org/10.1111/j.1399-3062.2009.00459.x>
- Chen, Y., Liu, Y., Lee, S. S., Tsai, H., Wann, S., Kao, C., Chang, C.-L., Huang, W., Huang, T., Chao, H., Li, C., Ke, C., & Lin, Y. E. (2005). Abbreviated duration of superheat-and-flush and disinfection of taps for Legionella disinfection: lessons learned from failure. *American Journal of Infection Control*, 33(10), 606–610.
<https://doi.org/10.1016/j.ajic.2004.12.008>
- Chirca, I. (2019). The hospital environment and its microbial burden: challenges and solutions. *Future Microbiology*, 14, 1007–1010. <https://doi.org/10.2217/fmb-2019-0140>
- Chotikanatis, K., Bäcker, M., Rosas-Garcia, G., & Hammerschlag, M. R. (2011). Recurrent intravascular-catheter-related bacteremia caused by Delftia acidovorans in a hemodialysis patient. *Journal of Clinical Microbiology*, 49(9), 3418–3421.
<https://doi.org/10.1128/JCM.00625-11>
- D'Alessandro, D., Fabiani, M., Cerquetani, F., & Orsi, G. B. (2015). Trend of Legionella colonization in hospital water supply. *Annali Di Igiene : Medicina Preventiva e Di Comunita*, 27(2), 460–466. <https://doi.org/10.7416/ai.2015.2032>
- Decker, B. K., & Palmore, T. N. (2014). Hospital water and opportunities for infection prevention. *Current Infectious Disease Reports*, 16(10), 432.
<https://doi.org/10.1007/s11908-014-0432-y>
- Despotovic, A., Milosevic, B., Milosevic, I., Mitrovic, N., Cirkovic, A., Jovanovic, S., & Stevanovic, G. (2020). Hospital-acquired infections in the adult intensive care unit-

- Epidemiology, antimicrobial resistance patterns, and risk factors for acquisition and mortality. *American Journal of Infection Control*, 48(10), 1211–1215.
<https://doi.org/10.1016/j.ajic.2020.01.009>
- Donlan, R. M., & Costerton, J. W. (2002). Biofilms: survival mechanisms of clinically relevant microorganisms. *Clinical Microbiology Reviews*, 15(2), 167–193.
<https://doi.org/10.1128/CMR.15.2.167-193.2002>
- Eda, R., Maehana, S., Hirabayashi, A., Nakamura, M., Furukawa, T., Ikeda, S., Sakai, K., Kojima, F., Sei, K., Suzuki, M., & Kitasato, H. (2021). Complete genome sequencing and comparative plasmid analysis of KPC-2-producing *Klebsiella pneumoniae* isolated from hospital sewage water in Japan. *Journal of Global Antimicrobial Resistance*, 24, 180–182. <https://doi.org/10.1016/j.jgar.2020.12.007>
- Eyre, D. W., Sheppard, A. E., Madder, H., Moir, I., Moroney, R., Quan, T. P., Griffiths, D., George, S., Butcher, L., Morgan, M., Newnham, R., Sunderland, M., Clarke, T., Foster, D., Hoffman, P., Borman, A. M., Johnson, E. M., Moore, G., Brown, C. S., ... Jeffery, K. J. M. (2018). A *Candida auris* Outbreak and Its Control in an Intensive Care Setting. *The New England Journal of Medicine*, 379(14), 1322–1331.
<https://doi.org/10.1056/NEJMoa1714373>
- Facciola, A., Pellicanò, G. F., Visalli, G., Paolucci, I. A., Venanzi Rullo, E., Ceccarelli, M., D'Aleo, F., Di Pietro, A., Squeri, R., Nunnari, G., & La Fauci, V. (2019). The role of the hospital environment in the healthcare-associated infections: a general review of the literature. *European Review for Medical and Pharmacological Sciences*, 23(3), 1266–1278. https://doi.org/10.26355/eurev_201902_17020
- Hassoun-Kheir, N., Stabholz, Y., Kreft, J.-U., de la Cruz, R., Romalde, J. L., Nesme, J., Sørensen, S. J., Smets, B. F., Graham, D., & Paul, M. (2020). Comparison of antibiotic-resistant bacteria and antibiotic resistance genes abundance in hospital and community wastewater: A systematic review. *The Science of the Total Environment*, 743, 140804.
<https://doi.org/10.1016/j.scitotenv.2020.140804>
- Hayward, C., Ross, K. E., Brown, M. H., & Whiley, H. (2020). Water as a Source of Antimicrobial Resistance and Healthcare-Associated Infections. *Pathogens (Basel, Switzerland)*, 9(8). <https://doi.org/10.3390/pathogens9080667>
- Jamal, A. J., Mataseje, L. F., Brown, K. A., Katz, K., Johnstone, J., Muller, M. P., Allen, V. G., Borgia, S., Boyd, D. A., Ciccotelli, W., Delibasic, K., Fisman, D. N., Khan, N., Leis, J. A., Li, A. X., Mehta, M., Ng, W., Pantelidis, R., Paterson, A., ... Mulvey, M. R. (2020). Carbapenemase-producing Enterobacterales in hospital drains in Southern Ontario, Canada. *The Journal of Hospital Infection*, 106(4), 820–827.
<https://doi.org/10.1016/j.jhin.2020.09.007>
- Johnson, R. C., Deming, C., Conlan, S., Zellmer, C. J., Michelin, A. V., Lee-Lin, S., Thomas, P. J., Park, M., Weingarten, R. A., Less, J., Dekker, J. P., Frank, K. M., Musser, K. A., McQuiston, J. R., Henderson, D. K., Lau, A. F., Palmore, T. N., & Segre, J. A. (2018).

- Investigation of a Cluster of *Sphingomonas koreensis* Infections. *The New England Journal of Medicine*, 379(26), 2529–2539. <https://doi.org/10.1056/NEJMoa1803238>
- Kandiah, S., Yassin, M. H., & Stout, J. (2013). Monochloramine use for prevention of *Legionella* in hospital water systems. *Infectious Disorders Drug Targets*, 13(3), 184–190. <https://doi.org/10.2174/1871526511313030006>
- K Thet, M., Pelobello, M. L. F., Das, M., Alhaji, M. M., Chong, V. H., Khalil, M. A. M., Chinniah, T., & Tan, J. (2019). Outbreak of nonfermentative Gram-negative bacteria (*Ralstonia pickettii* and *Stenotrophomonas maltophilia*) in a hemodialysis center. *Hemodialysis International. International Symposium on Home Hemodialysis*, 23(3), E83–E89. <https://doi.org/10.1111/hdi.12722>
- Loveday, H. P., Wilson, J. A., Kerr, K., Pitchers, R., Walker, J. T., & Browne, J. (2014). Association between healthcare water systems and *Pseudomonas aeruginosa* infections: a rapid systematic review. *The Journal of Hospital Infection*, 86(1), 7–15. <https://doi.org/10.1016/j.jhin.2013.09.010>
- Lowe, C., Willey, B., O'Shaughnessy, A., Lee, W., Lum, M., Pike, K., Larocque, C., Dedier, H., Dales, L., Moore, C., McGeer, A., & Mount Sinai Hospital Infection Control Team. (2012). Outbreak of extended-spectrum β -lactamase-producing *Klebsiella oxytoca* infections associated with contaminated handwashing sinks(1). *Emerging Infectious Diseases*, 18(8), 1242–1247. <https://doi.org/10.3201/eid1808.111268>
- Marchesi, I., Marchegiano, P., Bargellini, A., Cencetti, S., Frezza, G., Miselli, M., & Borella, P. (2011). Effectiveness of different methods to control legionella in the water supply: ten-year experience in an Italian university hospital. *The Journal of Hospital Infection*, 77(1), 47–51. <https://doi.org/10.1016/j.jhin.2010.09.012>
- Marchesi, Isabella, Paduano, S., Frezza, G., Sircana, L., Vecchi, E., Zuccarello, P., Oliveri Conti, G., Ferrante, M., Borella, P., & Bargellini, A. (2020). Safety and effectiveness of monochloramine treatment for disinfecting hospital water networks. *International Journal of Environmental Research and Public Health*, 17(17). <https://doi.org/10.3390/ijerph17176116>
- McEwen, S. A., & Collignon, P. J. (2018). Antimicrobial resistance: a one health perspective. *Microbiology Spectrum*, 6(2). <https://doi.org/10.1128/microbiolspec.ARBA-0009-2017>
- Millar, B. C., & Moore, J. E. (2020). Hospital ice, ice machines, and water as sources of nontuberculous mycobacteria: Description of qualitative risk assessment models to determine host-Nontuberculous mycobacteria interplay. *International Journal of Mycobacteriology*, 9(4), 347–362.
- Monegro, A. F., Muppidi, V., & Regunath, H. (2021). Hospital Acquired Infections. In *StatPearls*. StatPearls Publishing.
- Montanari, L. B., Sartori, F. G., Cardoso, M. J. de O., Varo, S. D., Pires, R. H., Leite, C. Q. F., Prince, K., & Martins, C. H. G. (2009). Microbiological contamination of a hemodialysis

- center water distribution system. *Revista Do Instituto de Medicina Tropical de Sao Paulo*, 51(1), 37–43. <https://doi.org/10.1590/s0036-46652009000100007>
- Nakamura, I., Amemura-Maekawa, J., Kura, F., Kobayashi, T., Sato, A., Watanabe, H., & Matsumoto, T. (2020). Persistent Legionella contamination of water faucets in a tertiary hospital in Japan. *International Journal of Infectious Diseases*, 93, 300–304. <https://doi.org/10.1016/j.ijid.2020.03.002>
- Nathwani, D., Varghese, D., Stephens, J., Ansari, W., Martin, S., & Charbonneau, C. (2019). Value of hospital antimicrobial stewardship programs [ASPs]: a systematic review. *Antimicrobial Resistance and Infection Control*, 8, 35. <https://doi.org/10.1186/s13756-019-0471-0>
- Perkins, K. M., Reddy, S. C., Fagan, R., Arduino, M. J., & Perz, J. F. (2019). Investigation of healthcare infection risks from water-related organisms: Summary of CDC consultations, 2014-2017. *Infection Control and Hospital Epidemiology*, 40(6), 621–626. <https://doi.org/10.1017/ice.2019.60>
- Rodríguez-Acelas, A. L., de Abreu Almeida, M., Engelman, B., & Cañon-Montañez, W. (2017). Risk factors for health care-associated infection in hospitalized adults: Systematic review and meta-analysis. *American Journal of Infection Control*, 45(12), e149–e156. <https://doi.org/10.1016/j.ajic.2017.08.016>
- Russotto, V., Cortegiani, A., Raineri, S. M., & Giaratano, A. (2015). Bacterial contamination of inanimate surfaces and equipment in the intensive care unit. *Journal of Intensive Care*, 3, 54. <https://doi.org/10.1186/s40560-015-0120-5>
- Salm, F., Deja, M., Gastmeier, P., Kola, A., Hansen, S., Behnke, M., Gruhl, D., & Leistner, R. (2016). Prolonged outbreak of clonal MDR Pseudomonas aeruginosa on an intensive care unit: contaminated sinks and contamination of ultra-filtrate bags as possible route of transmission? *Antimicrobial Resistance and Infection Control*, 5, 53. <https://doi.org/10.1186/s13756-016-0157-9>
- Scott, R. D., Culler, S. D., & Rask, K. J. (2019). Understanding the Economic Impact of Health Care-Associated Infections: A Cost Perspective Analysis. *Journal of Infusion Nursing*, 42(2), 61–69. <https://doi.org/10.1097/NAN.0000000000000313>
- Tang, W., Mao, Y., Li, Q. Y., Meng, D., Chen, L., Wang, H., Zhu, R., & Zhang, W. X. (2020). Prevalence of opportunistic pathogens and diversity of microbial communities in the water system of a pulmonary hospital. *Biomedical and Environmental Sciences : BES*, 33(4), 248–259. <https://doi.org/10.3967/bes2020.034>
- The Joint Commission. (2021a). *The Joint Commission* . New Water Management Requirements. <https://www.jointcommission.org/standards/prepublication-standards/new-water-management-requirements/>
- The Joint Commission. (2021b, January 1). *The Joint Commission*. <https://www.jointcommission.org/standards/about-our-standards/>

- Trautmann, M., Halder, S., Hoegel, J., Royer, H., & Haller, M. (2008). Point-of-use water filtration reduces endemic *Pseudomonas aeruginosa* infections on a surgical intensive care unit. *American Journal of Infection Control*, 36(6), 421–429. <https://doi.org/10.1016/j.ajic.2007.09.012>
- Vickers, R. M., Yu, V. L., Hanna, S. S., Muraca, P., Diven, W., Carmen, N., & Taylor, F. B. (1987). Determinants of *Legionella pneumophila* contamination of water distribution systems: 15-hospital prospective study. *Infection Control : IC*, 8(9), 357–363. <https://doi.org/10.1017/s0195941700067412>
- Vincenti, S., Quaranta, G., De Meo, C., Bruno, S., Ficarra, M. G., Carovillano, S., Ricciardi, W., & Laurenti, P. (2014). Non-fermentative gram-negative bacteria in hospital tap water and water used for haemodialysis and bronchoscope flushing: prevalence and distribution of antibiotic resistant strains. *The Science of the Total Environment*, 499, 47–54. <https://doi.org/10.1016/j.scitotenv.2014.08.041>
- Weber, D. J., Rutala, W. A., Miller, M. B., Huslage, K., & Sickbert-Bennett, E. (2010). Role of hospital surfaces in the transmission of emerging health care-associated pathogens: norovirus, *Clostridium difficile*, and *Acinetobacter* species. *American Journal of Infection Control*, 38(5 Suppl 1), S25–33. <https://doi.org/10.1016/j.ajic.2010.04.196>
- Weinbren, M. J. (2020). Dissemination of antibiotic resistance and other healthcare waterborne pathogens. The price of poor design, construction, usage and maintenance of modern water/sanitation services. *The Journal of Hospital Infection*. <https://doi.org/10.1016/j.jhin.2020.03.034>
- Yassin, M. H., Abramovitz, B., Hariri, R., McKibben, L., & Pinevich, A. J. (2020). Delftia acidovorans pseudo outbreak in portable reverse osmosis machines: Interventions to ensure safe and cost-effective hemodialysis. *American Journal of Infection Control*, 48(3), 304–308. <https://doi.org/10.1016/j.ajic.2019.11.027>
- Ziwa, M., Jovic, G., Ngwisha, C. L. T., Molnar, J. A., Kwenda, G., Samutela, M., Mulowa, M., & Kalumbi, M. M. (2019). Common hydrotherapy practices and the prevalence of burn wound bacterial colonisation at the University Teaching Hospital in Lusaka, Zambia. *Burns*, 45(4), 983–989. <https://doi.org/10.1016/j.burns.2018.11.019>