TABLE 1. Clostridium difficile Culture Results: Effectiveness of Manual Cleaning Versus UV-C

	Post Bleach CFUs		Post UV-C CFUs	
Site	≥10 CFUs	<10 CFUs	≥10 CFUs	<10 CFUs
Over-bed table	13	41	0	54
Toilet seat	9	65	0	74
Computer keyboard	3	19	1	43
Bathroom doorknob	2	20	0	22
Faucet handles	2	39	0	41
Bed side rails	1	3	0	4
Bedside commode	1	11	0	12
Recliner chair table	1	6	0	7
Call light	0	2	0	2

NOTE. CFU, colony-forming units; UV-C, ultraviolet light at 254 nanometers.

detection of CFU following UV-C treatment (odds ratio [OR], 0.027; 95% confidence interval [CI], 0.0006-0.1664; P < .0001). The hospital-onset C. difficile cases decreased to 11.4 per 10,000 patient days from 12.9 for the same period the prior year (during which UV-C had not yet been utilized).

In rooms of patients with confirmed CDI, UV-C treatment significantly reduced the amount of *C. difficile* spores present on the surfaces tested. Manual bleach cleaning alone resulted in residual spores in 13% of high-touch-surface cultures. These cultures were obtained as part of daily routine cleaning without the knowledge of Environmental Services workers. The performance of our cleaning staff is regularly evaluated by VeriClean blacklight audits, another objective evaluation tool demonstrating areas potentially missed by the cleaning staff. The average pass rate was 90.8% during the study period. Data recently published by Wong et al<sup>7</sup> are consistent with our findings; they reported that 5 of 22 rooms (22%) were positive for C. difficile after terminal cleaning.<sup>7</sup> The hospital-onset C. difficile rate decreased as well, even without 100% compliance of UV-C treatment of discharge enteric contact isolation rooms.

In patient rooms of those with confirmed CDI, adding UV-C treatment to daily bathroom and terminal discharge cleaning reduces the amount of C. difficile spores present on frequently contaminated surfaces. UV-C disinfection represents an additional measure for room cleaning to avoid inadvertent transfer of *C. difficile* spores to hands or other surfaces.

## ACKNOWLEDGMENTS

We would like to thank Walters Arrey and Denise Webb for their help in successful implementation of UV-C disinfection.

Financial support: The Ohio State University Comprehensive Cancer Center provided the funds for the Clostridium difficile cultures.

Potential conflicts of interest: All authors report no conflicts of interest relevant to this article.

> Christina Liscynesky, MD;<sup>1,2</sup> Lisa P. Hines, MACPR, BS, RN, CIC;<sup>2</sup> Justin Smyer, MPH, MLS(ASCP)CM, CIC;<sup>2</sup>

Megan Hanrahan, MBOE, CLSSGB;<sup>3</sup> Robert C. Orellana, MPH;<sup>4,5</sup> Julie E. Mangino, MD, FSHEA<sup>1,4</sup>

Affiliations: 1. Division of Infectious Diseases, Department of Internal Medicine, The Ohio State University Columbus, Ohio; 2. Department of Clinical Epidemiology, Comprehensive Cancer Center, The Ohio State University, Columbus, Ohio; 3. Department of Environmental Services, Comprehensive Cancer Center, The Ohio State University, Columbus, Ohio; 4. Department of Clinical Epidemiology, The Ohio State University Wexner Medical Center, Columbus, Ohio; 5. Division of Epidemiology, College of Public Health, The Ohio State University, Columbus, Ohio.

Address correspondence to Christina Liscynesky, MD, N1147 Doan Hall, 410 West 10th Avenue, Columbus, OH 43201 (Christina.Liscynesky@osumc.edu).

Received February 17, 2017; accepted May 25, 2017; electronically published July 3, 2017

Infect Control Hosp Epidemiol 2017;38:1116-1117

© 2017 by The Society for Healthcare Epidemiology of America. All rights reserved. 0899-823X/2017/3809-0019. DOI: 10.1017/ice.2017.126

## REFERENCES

- 1. National and state healthcare-associated infections progress report (updated March 2016). Centers for Disease Control and Prevention website. http://www.cdc.gov/nhsn/datastat/. Updated September 27, 2016. Accessed November 18, 2016.
- 2. Weber DJ, Anderson DJ, Sexton DJ, Rutala WA. Role of the environment in the transmission of Clostridium difficile in healthcare facilities. Am J Infect Control 2013;41:S105-S110.
- 3. Goldstein EJC, Johnson S, Maziade P-J, et al. Pathway to prevention of nosocomial Clostridium difficile infection. Clin Infect Dis 2015;60:S148-S158.
- 4. Hota B. Contamination, disinfection, and cross-colonization: are hospital surfaces reservoirs for nosocomial infection? Clin Infect Dis 2004;39:1182-1189.
- 5. Nerandzic MM, Thota P, Sankar CT, et al. Evaluation of a pulsed xenon ultraviolet disinfection syst. Infect Control Hosp Epidemiol 2015;36:192-197.
- 6. Boyce JM, Farrel PA, Towle D, Fekieta R, Aniskiewicz M. Impact of room location on UV-C irradiance and UV-C dosage and antimicrobial effect delivered by a mobile UV-C light device. Infect Control Hosp Epidemiol 2016;37:667-672.
- 7. Wong T, Woznow T, Petrie M, et al. Postdischarge decontamination of MRSA, VRE, and Clostridium difficile isolation rooms using 2 commercially available automated ultraviolet-C-emitting devices. Am J Infect Control 2016;44:416-420.

Resolution of Carbapenemase-Producing Klebsiella pneumoniae Outbreak in a Tertiary Cancer Center; the Role of Active Surveillance

Carbapenem-resistant Enterobacteriaceae (CRE) are a source of healthcare-associated infections with high attributable mortality. Carbapenemase-producing CRE (CP-CRE) (eg, KPC, OXA-48, NDM, IMP or VIM) are more commonly acquired

exogenously than non–CP-CRE. The most common CP-CRE in the United States harbors the *Klebsiella pneumoniae* carbapenemase (KPC) gene ( $bla_{KPC}$ ). Here, we report our investigation of a cluster of KPC-producing *K. pneumoniae* (KPC-KP) in a large academic medical facility cancer center.

The Johns Hopkins Hospital is a 1,059-bed academic medical facility in Baltimore, Maryland with an NCIdesignated Comprehensive Cancer Center comprising 6 geographically separate units. Overall, 3 units were involved in the outbreak: units A and B (16- and 15-bed hematologyoncology units) and unit C (a 15-bed bone marrow transplant [BMT] unit). Patients on these units undergo admission and weekly routine active surveillance for CRE. In July 2016, the hospital epidemiology and infection control staff was alerted by the microbiology laboratory that 2 patients had phenotypically identical CRE K. pneumoniae with similar antimicrobial susceptibility profiles in perirectal surveillance cultures. Pulsed-field gel electrophoreses (PFGE) confirmed the same strain. Epidemiological investigation revealed that the patients had consecutively occupied the same patient room in unit B 2 months prior: Patient 2 was admitted to the room 3 days after patient 1 was discharged. Patients 1 and 2 had negative admission and weekly perirectal surveillance cultures preceding the surveillance cultures that grew KPC-KP (13 and 36 days after admission, respectively), indicating probable acquisition during their hospital stays. Outbreak investigation and risk mitigation strategies were initiated.

Case-finding strategies included provider notification to identify suspected cases, microbiology laboratory request to alert for all KPC-KP isolates, and point-prevalence survey. On units B and C, where case patients were located, perirectal swabs were obtained from 24 patients, urine was obtained from 3 patients, and wound cultures were obtained from 2 patients. Contact-based precautions were used for all patients until point-prevalence results were obtained to interrupt potential transmissions from undetected carriers. There were no positive results. However, 2 other cases of KPC-KP were identified through clinician and laboratory notification; patient 3, who was readmitted to unit C with a positive blood culture and patient 4, who had not been recently admitted but who had had routine perirectal surveillance with KPC-KP during a prior admission to unit A. PFGE again confirmed the same strain. Whole-genome sequencing of the 4 case-patient isolates revealed a clonal K. pneumoniae strain producing KPC-2, CTX-M-15, TEM-1B, SHV-28, and OXA-1 sequence type 15. Environmental samples were taken for culture (13 high-touch surfaces and 2 bathroom sink drains) from the rooms and bathrooms of patients 1 and 2. A molecularly identical KPC-KP grew in the culture from an IV pole sample in the room of patient 2. Multiple other gram-negative organisms were cultured from the drains in both patient bathrooms.

Mitigation strategies included reinforcement of strict compliance with infection control precautions, enhanced environmental cleaning and disinfection, universal contact

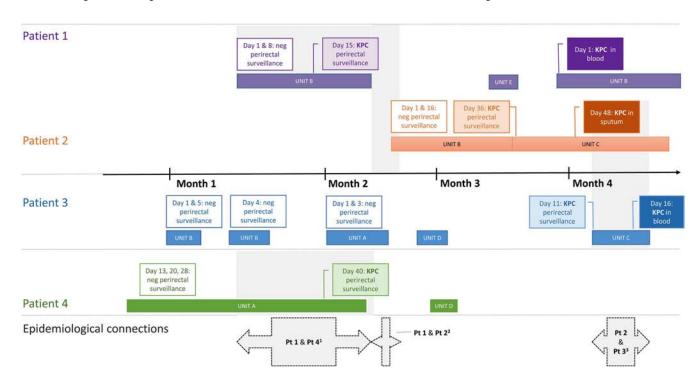


FIGURE 1. Timeline of KPC-KP patient inpatient stays and surveillance and clinical cultures. Patients are numbered in the order KPC-KP was identified. Solid lines denote inpatient stays. Inpatient days are numbered from the first inpatient day (ie, day 1) of each admission. Shaded areas indicate epidemiological connections: <sup>1</sup>Patients 1 and 4 were admitted to different units on the same floor. <sup>2</sup>Patient 2 occupied the same room after patient 1 was discharged. <sup>3</sup>Patients 2 and 3 were admitted to the same unit.

precautions for units with case patients pending negative point-prevalence surveys and discharge of all case patients, dedicated equipment for case patients, and cohorting patients with dedicated nursing staff. Enhanced environmental disinfection using hydrogen-peroxide vapor technology was used post discharge for rooms inhabited by a known case patient and, when feasible, for all other rooms and shared equipment on affected units. Patient 3, who had significant comorbidities, died from progressive multiorgan failure with neutropenic sepsis and KPC-KP bacteremia. Two patients died from unrelated causes and 1 patient proceeded to successful BMT. Figure 1 shows the timing of cultures and epidemiological connections. Patients 1 and 2 had consecutive stays in the same room; patients 2 and 3 were on the same unit at the same time. The link between patient 4 and the other patients is not immediately evident. Patient 4 was an inpatient at the cancer center at the same time as patient 1 but on a different unit. The surveillance cultures of these 2 patients first grew KPC-KP on the same day. However, equipment and staff are not confined to 1 unit in the cancer center.

There are reports of KPC-KP outbreaks persisting for up to 2 years.<sup>2-4</sup> In 2011, the NIH Clinical Center had a largely ICU-based outbreak involving 19 patients who became colonized or infected with blaKPC-positive K. pneumoniae, with 6 attributable deaths.<sup>2</sup> Patients with unrecognized CRE colonization may have served as undetected sources.<sup>5,6</sup> The interval between detection of colonization and infection in our outbreak ranged from 6 to 67 days. Active surveillance cultures at the cancer center allowed earlier identification and initiation of contact and other precautions rather than relying solely on clinical cultures, potentially contributing to the timely resolution of this outbreak. In addition, real-time molecular genotyping allowed for timely identification of full-genotype of  $\beta$ -lactamase genes, which was confirmed in additional patients.

This report highlights the importance of the hospital environment in the transmission of CRE as evidenced by 2 case patients having consecutive stays in the same hospital room and finding identical KPC-KP on the i.v. pole of a different room with a known infected patient. CRE have been found in patient rooms, including on equipment and sink drains.<sup>2,4,7</sup> Also, CRE can persist despite equipment and environmental cleaning and disinfection.<sup>2,4</sup> Our use of hydrogen peroxide vapor for shared equipment and once for each discharge room on affected units during the outbreak, rather than only rooms that had housed a known KPC-KP patient, may have shortened the outbreak period.

## ACKNOWLEDGMENTS

We acknowledge the contributions of The Johns Hopkins Hospital Microbiology Laboratory team; Shawna Lewis for processing the environmental cultures, Belita Opene for performing the molecular typing, and Tracy Howard for conducting the pulsed-field gel electrophoresis.

Financial support: No financial support was provided relevant to this article. Potential conflicts of interest: Drs Carroll, Maragakis, and Rock report grants outside the submitted work. All other authors report no conflicts of interest relevant to this article.

Clare Rock, MD, MS;1,2,3a Melanie S. Curless, RN, MPH, CIC;<sup>3a</sup> Maggie Cantara, MHS, CIC;<sup>3</sup> Seema Mehta, MD, MS;1 Kristen A. Marrone, MD;1 Karen C. Carroll, MD;4 Patricia Simner, PhD, D(ABMM);4b Lisa L. Maragakis, MD, MPH<sup>1,2,3b</sup>

Affiliations: 1. Division of Infectious Diseases, Department of Medicine, Johns Hopkins University School of Medicine, Baltimore, Maryland; 2. Armstrong Institute for Patient Safety and Quality, Johns Hopkins University School of Medicine, Baltimore, Maryland; 3. Hospital Epidemiology and Infection Control, The Johns Hopkins Hospital, Baltimore, Maryland; 4. Division of Medical Microbiology, Johns Hopkins University School of Medicine, Baltimore, Maryland.

<sup>a,b</sup>Authors of equal contribution, respectively.

Address correspondence to Clare Rock, MD, MS, Johns Hopkins University School of Medicine, Division of Infectious Diseases, 600 North Wolfe Street, Halsted 831, Baltimore, MD 21287-5425 (Clare.Rock@jhmi.edu).

PREVIOUS PRESENTATION: Part of the data reported here were presented as a poster at the Society for Hospital Epidemiology of America (SHEA) Spring Meeting 2017 in March 2017, in St. Louis, Missouri.

Received March 28, 2017; accepted June 2, 2017.; electronically published July 11, 2017

Infect Control Hosp Epidemiol 2017;38:1117-1119

© 2017 by The Society for Healthcare Epidemiology of America. All rights reserved. 0899-823X/2017/3809-0020. DOI: 10.1017/ice.2017.136

## REFERENCES

- 1. Centers for Disease Control and Prevention. Vital signs: carbapenem-resistant Enterobacteriaceae. Morb Mortal Wkly Rep 2013;62:165-170.
- 2. Snitkin ES, Zelazny AM, Thomas PJ, et al. Tracking a hospital outbreak of carbapenem-resistant Klebsiella pneumoniae with whole-genome sequencing. Sci Transl Med 2012;4: 148ra116.
- 3. Kanamori H, Parobek CM, Juliano JJ, et al. A prolonged outbreak of KPC-producing Enterobacter cloacae and Klebsiella pneumoniae driven by multiple mechanisms of resistance transmission at a large academic burn center. Antimicrob Agents Chemother 2016;61: e01516-16
- 4. Leitner E, Zarfel G, Luxner J, et al. Contaminated handwashing sinks as the source of a clonal outbreak of KPC-2-producing Klebsiella oxytoca on a hematology ward. Antimicrob Agents Chemother 2015;59:714-716.
- 5. Ben-David D, Maor Y, Keller N, et al. Potential role of active surveillance in the control of a hospital-wide outbreak of carbapenem-resistant Klebsiella pneumoniae infection. Infect Control Hosp Epidemiol 2010;31:620-626.
- 6. Facility guidance for carbapenem-resistant Enterobacteriaceae. Centers for Disease Control and Prevention website. https:// www.cdc.gov/hai/organisms/cre/. Published 2015. Accessed March 21, 2017.
- 7. Kizny Gordon AE, Mathers AJ, Cheong EYL, et al. Is the hospital water environment a reservoir for carbapenem-resistant organisms causing hospital-acquired infections? A systematic review of the literature. Clin Infect Dis 2017:cix132.