Letter to the Editor



Strategy to limit multidrug-resistant *Acinetobacter baumannii* transmission in a cohort coronavirus disease 2019 (COVID-19) critical care unit

Anucha Apisarnthanarak MD¹ ^(b) and David J. Weber MD, MPH²

¹Division of Infectious Diseases, Thammasat University Hospital, Pathum Thani, Thailand and ²Gillings School of Global Public Health, Chapel Hill, North Carolina, United States

To the Editor-Coinfection with multidrug-resistant organisms (MDROs) among coronavirus disease 2019 (COVID-19) patients is common in critical care patients with a prolonged length of stay in critical care units, likely due to the coadministration of high-dose steroids and the prolonged duration of mechanical ventilation.¹ The control of MDROs among COVID-19 patients is also difficult, given the requirement for airborne plus contact isolation among these patients and the difficulty in wearing and changing personal protective equipment (PPE) in a critical care unit.² The situation is much more challenging in middle- and lower-income countries where cohort areas in airborne-isolation critical care units are often designed, instead of single airborneisolation rooms in the critical care unit. We report the experience of controlling an MDR Acinetobacter baumannii outbreak in a COVID-19 critical care unit that featured airborne isolation cohorting areas together with limited standard single airborne isolation rooms in Thailand.

On March 1, 2021, at Thammasat University Hospital (Pratum Thani, Thailand), the first case of MDR A. baumannii was detected in a COVID-19 critical care unit, followed by 1 additional patient who was located next to the index patient 2 days later in the same cohorting area. In this 10-bed critical care unit, there were 2 sections of 4-bed airborne isolation cohort areas and 2 singleroom beds which were airborne isolation rooms. The nurseto-patient ratio in this unit was 2.5 to 1. After the detection of the first case, a root-cause analysis revealed the possibility of cross transmission because healthcare personal (HCP) are unable to change PPE between caring for patients in the cohort area as well as the possibility of an unrecognized case of MDR A. buamannii referred from another hospital. A policy to prevent transmission was initiated that included isolation of MDR A. baumannii patients in single-bed isolation rooms, assigning specific nurses to care for cases with MDR A. baumannii, changing gloves between cases, putting an extra sheet cover on the provider between care for MDR A. baumannii cases and daily environmental disinfection in cohort and single beds with a quaternary ammonium compound.

Feedback regarding compliance with infection prevention practices by HCP was given daily. After 2 weeks, 4 additional MDR *A. baumannii* cases occurred in the cohort area (the incidence rate, 16.9 cases per 1,000 patient days), despite full compliance with policies. A subsequent root-cause analysis suggested the possibility of widely disseminated environmental contamination with MDR *A. baumannii* together with the possibility of cross transmission of MDR *A. baumannii* by HCP unable to change gowns between patients in the cohort area.

Additional interventions at this stage included unit closure for hydrogen peroxide vapor disinfection, development of risk stratification criteria for housing high-risk patients with MDR A. baumannii in the 2 isolation rooms, development of an antibiotic stewardship program to limit broad-spectrum antibiotics and to de-escalate broad-spectrum antibiotics among COVID-19 patients, and development of a policy to discontinue isolation for COVID-19 patients.³ Continuous monitoring and feedback of MDR A. baumannii incidence and infection prevention compliance with such policies among HCP was performed daily. During the subsequent 8 weeks, 2 additional cases of MDR A. baumannii were detected. A significant reduction in MDR A. baumannii incidence occurred compared to the period before the intervention: 16.9 cases per 1,000 patient days versus 3.6 cases per 1,000 patient days (P < .001). Infection prevention compliance monitoring among HCP indicated full compliance with all components of infection prevention.

It is well recognized that MDR A. baumannii are selected by use of broad-spectrum antibiotics and often have an environmental reservoir that can facilitate rapid spread in critical care units if appropriate interventions are not introduced.^{4,5} Also, the constant use of gloves and gowns during a SARS outbreak led to an increase in transmission of MDROs, particularly methicillin-resistant Staphylococcus aureus.⁶ Outbreaks of MDR A. baumannii can be more difficult to control in middle- and lower-income countries where infrastructure may not be adequate (eg, suboptimal design of negative pressure airborne isolation units, inadequate nurseto-patient ratio). Such conditions require a practical strategy to control MDROs in the resource limited settings. Our experience suggests that policies that using an additional sheet to protect contamination of HCP gowns, frequent changes of gloves, assignment of specific nurses to care for MDR A. baumannii cases as well as basic environmental disinfection were not able to terminate an

© The Author(s), 2021. Published by Cambridge University Press on behalf of The Society for Healthcare Epidemiology of America.

Author for correspondence: Anucha Apisarnthanarak, Email: anapisarn@yahoo.com Cite this article: Apisarnthanarak A and Weber DJ. (2021). Strategy to limit multidrugresistant Acinetobacter baumannii transmission in a cohort coronavirus disease 2019 (COVID-19) critical care unit. Infection Control & Hospital Epidemiology, https:// doi.org/10.1017/ice.2021.289

outbreak of MDR *A. baumannii* if the HCP could not change PPE between patients in cohort areas. Additional strategies are needed in situations that do not allow changing PPE easily between cases in the cohort areas. These strategies must feature multimodal approaches that include risk stratification for index patients that may potentially harbor MDR *A. baumannii* with isolation in single rooms, an antibiotic stewardship program for COVID-19 patients, and policies to discontinue COVID-19 isolations as well as a policy to perform robust terminal environmental disinfection. Such strategies, together with fully compliance with infection prevention measures, will help limit the transmission of MDR *A. baumannii* in COVID-19 cohort areas in critical care units. Additional studies to evaluate practical strategies to help limit transmission of MDR-pathogens in cohort-type COVID-19 critical care units are needed.

Acknowledgments.

Financial support. No financial support was provided relevant to this article.

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

References

- Karruli A, Boccia F, Gagliardi M, et al. Multidrug-resistant infections and outcome of critically ill patients with coronavirus disease 2019: a single center experience. Microb Drug Resist 2021. doi: 10.1089/mdr.2020.0489.
- Lynch JB, Davitkov P, Anderson DJ, et al. Infectious Diseases Society of America guidelines on infection prevention for healthcare personnel caring for patients with suspected or known COVID-19. Clin Infect Dis 2020. doi: 10.1093/cid/ciaa1063.
- 3. Rhee C, Kanjilal S, Baker M, Klompas M. Duration of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infectivity: when is it safe to discontinue isolation? *Clin Infect Dis* 2021;72:1467–1474.
- Ng TM, Teng CB, Lye DB, Apisarnthanarak A. A multicenter case-case control study for risk factors and outcomes of extensively drug-resistant *Acinetobacter baumannii* bacteremia. *Infect Control Hosp Epidemiol* 2014;35: 49–55.
- Teerawattanapong N, Kengkla K, Dilokthornsakul P, Saokaew S, Apisarnthanarak A, Chaiyakunapruk N. Prevention and control of multidrugresistant gram-negative bacteria in adult intensive care units: a systematic review and network meta-analysis. *Clin Infect Dis* 2017;64 suppl 2:S51–S60.
- Yap FHY, Gomersall CD, Fang KSC, et al. Increase in methicillin-resistant Staphylococcus aureus acquisition rate and change in pathogen pattern associated with outbreak of severe acute respiratory syndrome. Clin Infect Dis 2004;39:511–516.