Emergence of NDM-producing *Pseudomonas aeruginosa* among hospitalized patients and impact on antimicrobial therapy during the coronavirus disease 2019 (COVID-19) pandemic

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To the Editor—*Pseudomonas aeruginosa* is a leading cause of nosocomial infection, mainly in intensive care patients. Notably, *P. aeruginosa* has a wide variety of resistance mechanisms that often restrict therapeutic options. Although carbapenem resistance among them has been observed worldwide in the past decade.1

Distinct mechanisms, such as *OprD* porin loss, overexpression of MexAB-OprM efflux pump, and carbapenemase production may confer a carbapenem-resistance phenotype in *P. aeruginosa*.1,2 Among the carbapenemase enzymes, metallo-β-lactamases (MBLs) have one of the most potent resistance mechanisms because they inactivate virtually all β-lactam agents, except aztreonam. Different types of MBLs have been described among *P. aeruginosa* isolates, particularly the SPM-1 type in Brazil, with varied degrees of endemicity.3,4 Nowadays, other MBLs have emerged around the world, mainly New Delhi metallo-β-lactamases (NDMs), which are present largely in Enterobacterales, but also in nonfermenters (*Acinetobacter* and *Pseudomonas* spp.).3

The current antimicrobial pipeline offers short-term treatment of infections due to MBL producers. Thus, our most important tool against the spread of MBL producers relies on intensified infection control, surveillance, and antimicrobial stewardship policies. We sought to determine the prevalence of carbapenem-resistance and MBL production and to assess the impact of them on resistance rates to other antimicrobial agents among *P. aeruginosa* nosocomial isolates.

From October 2020 to March 2021, *P. aeruginosa* isolates were recovered from hospitalized patients in Porto Alegre city and its metropolitan area as part of a survey study on antimicrobial resistance. Only the first isolate of each patient was included in the survey. Isolates were confirmed as *P. aeruginosa* based on oxidase and pyocyanin production or, for atypical isolates, by MALDI-TOF (bioMérieux, Marcy l’Etoile, France). Susceptibility was determined using the disk diffusion for amikacin, cefepime, cefazidime, ciprofloxacin, imipenem, meropenem and piperacillin-tazobactam, and the results were interpreted according to CLSI guidelines.6 Polymyxin B susceptibility was determined using broth microdilution. All isolates resistant to carbapenems were subjected to the NG-test CARBA 5 multiplex immunochromatographic assay for the detection of *bla*KPC, *bla*OXA-48-like, *bla*NDM, *bla*VIM, and *bla*IMP carbapenemases (NG Biotech, Gulpry, France) and were further confirmed by polymerase chain reaction assay (PCR). When an MBL enzyme was detected, an in vitro synergism between ceftazidime/avibactam and aztreonam was evaluated using a disk-diffusion approximation test, as published elsewhere, with modifications.7

Statistical analyses were carried out using SPSS version 13.0 software (IBM, Chicago, IL). Prevalence ratio (PR) and 95% confidence intervals (CIs) were calculated. *P* values were calculated using the $\chi^2$ or Fisher exact test.

In total, 278 *P. aeruginosa* isolates were recovered from 156 patients during the study period. Among the 156 nonduplicate isolates, 83 (53.2%) were recovered from intensive care patients. These were more prevalent in respiratory secretions (*n* = 88; 56.4%), urine (*n* = 34; 21.8%), blood (*n* = 16; 10.3%), and other secretions (*n* = 18; 11.5%).

Overall, higher resistance rates were noted for meropenem (47.4%), followed by imipenem (43.6%), cefepime (39.1%), piperacillin/tazobactam (37.2%), ciprofloxacin (33.3%), ceftazidime (32.7%), amikacin (28.2%), and gentamicin (22.4%).

An MBL-producing *P. aeruginosa* isolate was recovered at least once from 27 (17.3%: 95% CI, 12.2%–24.0%) of these 156 patients, and the *bla*NDM-1 was the sole MBL enzyme detected. All isolates were susceptible to polymyxin B. However, NDM-1 production was responsible for a high β-lactam resistance rate (Fig. 1). Additionally, resistances to amikacin (74.1% vs 17.0%; PR, 4.34; 95% CI, 2.12%–9.0%), gentamicin (70.4% vs 12.4%; PR, 5.67; 95% CI, 2.6%–12.4%; *P* < .001), and ciprofloxacin (100.0% vs 19.4%; PR, 5.16; 95% CI, 2.6%–10.2%; *P* < .001) were significantly higher among NDM-1 producers than nonproducing *P. aeruginosa* isolates. Table 1 illustrates the susceptibility profile of the 27 NDM-1–producing *P. aeruginosa* isolates. Ceftazidime/avibactam plus aztreonam showed an in vitro synergistic result in all NDM isolates, despite the resistance to both drugs when they were individually tested.

The relatively low prevalence of carbapenem-resistance driven by MBL enzyme revealed in this survey allowed us to discourage coverage with empirical therapy because there are therapeutic options for non–carbapenemase-producing *P. aeruginosa* isolates (eg, cefotaxime/tazobactam, data not shown). However, we emphasize the potential for dissemination of an NDM–1–producing *P. aeruginosa* among critically ill patients in Southern Brazil which possibly could displace other MBL enzyme (mostly SPM-1) or another resistance mechanism as driver for carbapenem resistance.
All NDM-1–producing *P. aeruginosa* isolates showed an in vitro synergism for the ceftazidime/avibactam plus aztreonam. This association has been described as an option to treat infections due to MBL-producing organisms.\(^8,9\) The rationale is based on the ability of avibactam to inhibit serine-\(\beta\)-lactamases and the lack of hydrolytic activity of MBLs against aztreonam.\(^10\) Despite the lack of knowledge about its effectiveness in MBL-related infections, this synergistic activity may be an important strategy because polymyxins can appear active in vitro but can be flawed in achieving a clinical response.

In conclusion, the emergence of NDM-1–producing *P. aeruginosa* isolates and its impact on a high resistance rate was observed. Because nosocomial antimicrobial resistance can result in higher cost of treatment and increased mortality, it becomes crucial to limit the risk of NDM-1–producing *P. aeruginosa* infection and its spreading. For this, proper use of antimicrobials and strict infection control measures avoiding coinfections in a pandemic scenario is mandatory.

**Acknowledgments.** The authors thank Pfizer for providing the ceftazidime/avibactam disks.

**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**


