Difficulty in detecting low levels of polymyxin resistance in clinical Klebsiella pneumoniae isolates: evaluation of Rapid Polymyxin NP test, Colispot Test and SuperPolymyxin medium

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Abstract

Polymyxins are important therapeutic options for treating infections, mainly those caused by carbapenem-resistant *Klebsiella pneumoniae*. Specific chemical characteristics of polymyxins make it difficult to perform antimicrobial susceptibility testing, especially within the clinical laboratory. Here we aimed to evaluate the performance of three phenotypic methods: Rapid NP Polymyxin Test, ColiSpot test and the SuperPolymyxin medium. To accomplish this, 170 non-duplicate clinical *K. pneumoniae* isolates were analysed (123 colistin-resistant and 47 susceptible). The sensitivity and specificity obtained for Rapid Polymyxin NP Test, Colispot and SuperPolymyxin medium were, respectively, 90% and 94%, 74% and 100%, and 82% and 85%. Very major errors occurred more frequently in low-level colistin-resistant isolates (MICs 4 and 8 μg/mL). Rapid Polymyxin NP proved to be a method capable of identifying colistin-resistant strains in acceptable categorical agreement. However, major errors and very major errors of this method were considered unacceptable for colistin-resistance screening. Although the Colispot test is promising and easy to perform and interpret, the results did not reproduce well in the isolates tested. The colistin-containing selective medium (SuperPolymyxin) showed limitations, including quantification of mucoid colonies and poor stability. Nevertheless, Colispot and SuperPolymyxin medium methods did not present acceptable sensitivity, specificity and categorical agreement. It is essential to use analytical tools that faithfully reproduce bacterial resistance *in vitro*, especially in last-line drugs, such as polymyxins, when misinterpretation of a test can result in therapeutic ineffectiveness.

Keywords: Antibiotic resistance, Clinical isolates, Klebsiella pneumoniae, Phenotypic detection, Polymyxin resistance

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Introduction

Polymyxins (colistin and polymyxin B) are important therapeutic options for treating infections, mainly those caused by carbapenem-resistant *Klebsiella pneumoniae* [1]. Resistance has

been reported frequently, mainly mediated by chromosomal gene mutations [2–4]. In this context, the two-component systems PmrA/PmrB and PhoP/PhoQ and their negative regulatory gene *mgrB* are related to lipopolysaccharide modifications that decrease affinity for polymyxins, thereby reducing susceptibility [5]. More recently, mutations in the *crrB* gene, derived from a third two-component system, CrrAB, have been associated with changes in lipopolysaccharide, leading to resistance to polymyxins [6–8]. Resistance mediated by plasmid genes (*mcr*) encoding bacterial lipopolysaccharide-modifying enzymes has also been reported, starting with the characterization of the *mcr-1* gene in 2015 [9], which was followed by the description of numerous allelic variants (*mcr-2* to *mcr-9*) [10–17], including in carbapenem-resistant isolates and in high-risk clones [18,19].

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Whether caused by plasmid or chromosomal genes, the detection of polymyxin resistance faces technical difficulties, because of its cationic nature and its propensity to adsorb to plastic surfaces [20]. Therefore, methods widely used in clinical laboratories such as disc diffusion assays, and Etests are not suitable [21-23]. Furthermore, MICs obtained with gradient strips are markedly lower than those observed with the reference method, which could result in the misinterpretation of resistant strains as susceptible, especially in low-level colistinresistant bacteria [24]. Alternative phenotypic methods have been developed to enable the detection of polymyxin resistance in clinical microbiology laboratories. The Rapid NP Polymyxin Test, developed by Nordmann et al. in 2016, is based on verification of glucose metabolism associated with bacterial growth in the presence of polymyxin and phenol red as a pH indicator [25]. The Colispot Test is another phenotypic test published by Jouy et al. in 2017 that aims to determine susceptibility to polymyxins by analysing the diameter formed around a drop of a standardized antimicrobial solution on the agar surface [26]. Selective media containing polymyxin have also been developed for the detection of antimicrobial resistance [27-31].

Recently, broth microdilution (BMD), broth disc elution and agar dilution were recommended by the CLSI M100 30th edition guideline for determining colistin MICs. However, for polymyxin B, only BMD was approved [32]. When carrying out BMD, it is important to consider the use of polystyrene plates, as recommended by the joint CLSI-EUCAST Polymyxin Breakpoints Working Group [33]. It is known that most clinical microbiology laboratories lack the human or financial resources to perform the broth microdilution test recommended.

In this context, the Laboratório de Pesquisa em Infecção Hospitalar (LAPIH), located at Oswaldo Cruz Institute (Rio de Janeiro, Brazil), which is part of the Brazilian National Programme for Monitoring the Prevalence of Bacterial Resistance, conducted by the National Health Surveillance Agency, routinely receives clinical bacterial isolates from Brazilian health-care services to investigate resistance mechanisms. In this study, we aimed to evaluate the performance of these three alternative phenotypic methods: Rapid NP Polymyxin Test, ColiSpot test and the SuperPolymyxin medium (Plast Labor, Rio de Janeiro, Brazil), especially for clinical isolates with MICs near the breakpoint.

Material and methods

Clinical strains

We evaluated 170 non-duplicate *K. pneumoniae* clinical isolates (blood, urine, tracheal aspirate, rectal swab and catheter tip) received by LAPIH from January to December 2016 from eight

Brazilian states. The species identification was confirmed by conventional biochemical tests and the isolates were stored at -40° C in skimmed milk (DIFCO, Hants, UK) [34].

From these isolates, 123 were resistant to colistin (MIC 4–128 µg/mL) and 47 were susceptible (MIC \leq 2 µg/mL). From all resistant isolates, only three carried the *mcr-1* gene (screened by conventional PCR) [9]. Furthermore, several isolates showed mutations in chromosomal genes, including disruption of *mgrB* by insertion sequences such as ISKpn13 and IS903B (data not shown), detected by sequencing and manual curation with Geneious v.1.6.8 (Biomatters Ltd, Auckland, New Zealand) and BLAST tool (https://www.ncbi.nlm.nih.gov). For each isolate, the BMD reference method and the three phenotypic tests evaluated were performed on the same day, starting from the same primary culture.

Broth microdilution

The reference method was performed using colistin sulphate powder (Sigma, St Louis, MO, USA). No additives were included (no polysorbate-80 or other surfactants). Colistin stock solutions were prepared and kept at -40° C and diluted in fresh cation-adjusted Mueller–Hinton broth (Oxoid, Basingstoke, UK) during assay only. The susceptibility to colistin was evaluated in duplicate, according to the ISO 20776-1 methodology [35]. Hence, two-fold serial dilutions of colistin were performed in a 96-well polystyrene plate, obtaining a concentration range of 0.25–256 μ g/mL. The results were interpreted according to the EUCAST guideline (www.eucast.org) [36].

Rapid Polymyxin NP test

Fresh colonies grown on Mueller–Hinton agar (Oxoid) were used to perform the Rapid Polymyxin NP Test as described by Nordmann et al. [25]. A bacterial suspension with an optical density of 3.0–3.5 McFarland ($\approx 10^9$ CFU/mL) was prepared for each isolate and, after mixing with the Rapid NP colistin medium, resulted in a bacterial concentration of $\sim 10^8$ in each well of the plate. They were considered resistant when the colour of the developing solution changed from orange/red to yellow after incubation at 35°C \pm 2°C for 20 minutes to 4 hours, which indicated bacterial metabolism in the presence of a 3.75 µg/mL colistin concentration (Fig. 1).

Colispot test

ColiSpot was performed as described by Jouy et al. [26] and consisted of applying a single drop (10 μ L) of 8 μ g/mL colistin solution onto the surface of a previously inoculated Mueller–Hinton agar with a 0.5 McFarland bacterial suspension (1.5 × 10⁸ CFU/mL). Resistance was revealed when the inhibition zone did not exceed 5 mm (Fig. 2).



FIG. 1. Representative results of the Rapid Polymyxin NP test. Columns 1, 3 and 5 contain colistin-free solution and columns 2, 4 and 6 contain 5 μg/mL colistin-solution wells. A1-A2 wells were inoculated with the susceptible control strain *Escherichia coli* ATCC25922. Wells C1–C2 and D1-D2 were inoculated with the resistant strains *Klebsiella pneumoniae* CCBH22220 and *Proteus mirabilis* CCBH6854, respectively. G3-G4 wells were inoculated with NaCl solution only (no growth control). Wells D3-D4, F3–F4 and E5-E6 are examples of isolates considered resistant by the test. Wells A5-A6, B5–B6 and H5–H6 are examples of isolates considered susceptible by the Rapid Polymyxin NP test.

SuperPolymyxin medium

The SuperPolymyxin medium (PlastLabor, Rio de Janeiro, Brazil) was manufactured as described by Nordmann, Jayol and Poirel in 2016 [27]. Based on the eosin methylene blue agar with a final concentration of 3.5 µg/mL colistin, the medium also contained 10 µg/mL daptomycin for inhibition of Streptococcus and Staphylococcus species, and 5 µg/mL amphotericin B, an anti-fungal agent. An inoculum with an optical density of 0.5 McFarland standard (1.5 \times 10⁸ CFU/mL) was used. Then, 1 μ L of each bacterial suspension ($\sim 1.5 \times 10^5$ CFU/mL) was plated on the medium using calibrated bacteriological handles. The number of viable colonies was counted after 24 hours of incubation in an aerobic atmosphere at 35°C ± 2°C. To verify isolate viability, eosin methylene blue agar was inoculated concomitantly. Preliminary data from our laboratory show that the Super-Polymyxin medium tested in the present study allowed the growth of sensitive reference strains, and these data corroborate with others described in the literature [28]. Hence, we

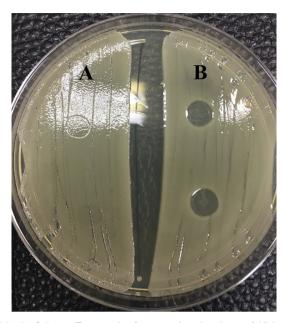


FIG. 2. Colispot Test results for two clinical isolates of *Klebsiella pneumoniae* evaluated in this study. In 'A', a resistant isolate (absence of zone of inhibition). In 'B', a susceptible isolate (7-mm inhibition zone). Two drops of colistin solution (8 μ g/mL) were added separately to each isolate for comparison.

proposed a cut-off point of five colonies, to increase specificity, above which the isolate was considered resistant to colistin.

Control strains, replicates and analysis of results

Escherichia coli ATCC25922 was included as a susceptible strain. Klebsiella pneumoniae CCBH22220 (colistin MIC 8 μ g/mL) carrying the mcr-1 gene (obtained from the CCBH collection, http://ccbh.fiocruz.br/) and Proteus mirabilis CCBH6854, an intrinsically polymyxin-resistant species, were used as resistant controls.

The BMD reference method was performed in duplicate, with concordant values for most isolates. In case of disagreement, a third determination was performed, and the MIC obtained twice was accepted. The phenotypic methods were evaluated twice when they diverged from the reference method, and in case of disagreement with the first run, a third analysis was performed. In case of inconsistent results, the final interpretation of the test was based on the result as suggested by the majority of the three test runs.

For the analysis of the results, the values of sensitivity and specificity of each phenotypic method were calculated. The Category Agreement (CA), which is considered the percentage of isolates that produce the same categorical result (susceptible, intermediate or resistant) when compared with the reference method, was obtained. Major errors (ME, resistant results by the new method and susceptible results by the reference standard

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method) and very major errors (VME, susceptible result by the new method and a resistant result by the reference standard method) were also calculated [36,37]. The analysis of the results followed the recommendations of CLSI guideline M23 [38] and the criteria required by the ISO 20776-I standard [35].

Results

Klebsiella pneumoniae isolates

The 123 clinical isolates of colistin-resistant K. pneumoniae had MICs ranging from 4 to >128 µg/mL (MIC₅₀ 32 µg/mL; MIC₉₀ >128 µg/mL). The 47 randomly selected susceptible isolates (MIC \leq 2 µg/mL) showed MICs that ranged from 0.25 to 2 µg/mL (MIC₅₀ 0.5 µg/mL; MIC₉₀ 2 µg/mL). Of all the isolates tested, 45 showed MICs close to the breakpoint (MICs I-8 µg/mL). Most of all resistant isolates were recovered from patients' blood, urine or tracheal aspirate, where the presence of K. pneumoniae may require the use of antimicrobial therapy (see Supplementary material, Table SI). In this context, we emphasize that I2I (98%) of the resistant isolates showed coresistance to at least one carbapenem (data not shown), which makes the use of polymyxin a probable alternative.

Performance evaluation of phenotypic tests

The Rapid Polymyxin NP Test showed 90% sensitivity and 94% specificity. Colispot Test showed 74% and 100% and Super-Polymyxin medium showed 82% and 85% sensitivity and specificity, respectively. Table 1 shows the results of the three phenotypic tests for the 170 isolates of *K. pneumoniae*. As required by CLSI and the ISO 20776-1 standard [35,38],

Categorical Agreement (CA) has been determined. Our study showed that for susceptible isolates, Rapid-NP and Colispot had CA >90%. The method using the polymyxin previously incorporated into the culture medium (SuperPolymyxin) presented low CA (70%) for the susceptible isolates with MICs close to the breakpoint (MIC I or 2 µg/mL). In the 123 resistant isolates, the accuracy of the tests increased with increasing MIC, >80% of CA is observed in high-level polymyxin-resistant isolates. It is important to highlight that VME occurred more frequently in low-level polymyxin-resistant isolates (MICs 4 and 8 µg/mL), meaning a false-susceptible result (Table 1). Especially for K. pneumoniae isolates, which characteristically have mucoid colonies in solid media [34], the quantification became difficult and lacked precision. The Rapid Polymyxin NP test had a CA >90% in isolates with MIC >16 µg/mL, which was considered acceptable (>90%) [35,38]. Despite this, the ME and VME calculated were 6.4% and 9.7%, respectively, which is not acceptable for colistin-resistance screening, required by the ISO 20776-1 standard (<3%). For resistant isolates, Colispot and SuperPolymyxin medium only showed CA >90% at MICs ≥128 μg/mL and 32 μg/mL, respectively. For low resistant MICs (4 and 8 µg/mL) (data underlined in Table I), CA was insufficient for acceptance (<90%) for all three methods. All phenotypic methods evaluated were able to detect resistance in the three mcr-1-positive isolates, as shown in Table 1.

Discussion

In 2017, the World Health Organization ranked carbapenemase-producing *Enterobacterales* as a priority I for

TABLE I. Performance evaluation of three phenotypic tests to determine resistance to polymyxins

(C ^a g/mL) 28 8	isolates (n = 170) 20 8 33 25	0 0 0 0 0	20 8 33 25	100% 100% 100% 100%	0 0 4	20 8 29	100% 100% 87.9%	3 0	R 17 8	CA 85% 100%	
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	13	0 I		100%	4			3	30	90.9%	
		1			4	21	84%	3	23	92%	
			12	92.3%	6	7	53.8%	3	10	76.9%	
	11 °	3	8 ^c	72.7%	6	5 °	<u>45.4%</u>	4	7 °	63.6%	
	13 ^d	<u>3</u> <u>8</u>	8 c 5 d	38.5%	<u>6</u> <u>12</u>	5 c 1 d	7.7%	<u>4</u> <u>6</u>	7 c d	53.8%	
	П	10	1	90,9%	П	0	100%	8	3	72.7%	
	10	10	0	100%	10	0	100%	7	3	70%	
;	22	20	2	90.9%	22	0	100%	21	1	95.4%	
.5	4	4	0	100%	4	0	100%	4	0	100%	
All isolates			CA = 91.2% No. of MF = 3 (6.4%)			CA = 81.2% No. of MF = 0 (0%)			CA = 82.9% No. of ME = 7 (14.9%)		
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Abbreviations: S, susceptible; R, resistant; CA, categorical agreement; ME, major errors; VME, very major errors. The underlined data represent low-level resistance isolates (MICs 4 or 8 µs/mL), where the lowest categorical agreement was observed.

⁴ or 8 µg/mL), where the lowest categorical agreement was observed.

aMICs were determined using the broth microdilution test according to European Committee on Antimicrobial Susceptibility Testing (EUCAST guidelines.

^bAntimicrobial susceptibility tests for colistin were interpreted according to EUCAST guidelines.

^cIncluding two *mcr-1*-positive isolates.
^dIncluding one *mcr-1*-positive isolate.

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research and development of new antibiotics—emphasizing the scarcity of therapeutic options [39]. In these scenarios, polymyxins are widely prescribed as a therapeutic option and then the clinical microbiology laboratory plays an important role in antimicrobial therapy. In Brazilian hospitals, carbapenem and polymyxin co-resistance has been described [18].

Considering the importance of early detection of polymyxin resistance for a positive patient outcome and the difficulty of performing the microdilution method in routine microbiology laboratories, phenotypic tests such as those evaluated in this study have been developed and evaluated worldwide [26,27,40-43]. All three methods evaluated proved to be easy to prepare and perform. The performance of the Rapid Polymyxin NP Test was comparable to that observed by other authors [40-43]. However, here we emphasize the difficulty of detecting resistance in isolates with MICs close to the breakpoint (4 and 8 µg/mL) for all evaluated methods. These findings have been described even for other methods, such as Colistin Broth Disk Elution [30,40,41]. Detection of polymyxin resistance in low MICs is important as the spread of resistance by plasmid mechanisms, such as the mcr gene, has been associated with bacteria that exhibit low resistance levels (i.e. 4-16 µg/ mL) [44-46].

Rapid Polymyxin NP proved to be a faster and less laborious technique than conventional BMD. Limitations for the use of this technique in some clinical laboratories are the preparation of the NP solution, which requires pH adjustment, and the standard polymyxin solution, which requires the use of analytical balances, which may be unavailable. For the Colispot Test, developed in 2016 [26], as far as we know, there are no published papers that aimed to evaluate its performance. Although the method is promising and easy to perform and interpret, the results did not reproduce well in the isolates tested, especially in strains with MICs close to the breakpoint. Regarding the SuperPolymyxin medium, other authors have observed satisfactory results, including the use of chromogenic media [29] and the Colistin Agar Test, which proposes to use Mueller-Hinton agar with colistin added as a screening [30]. However, in the present study, the categorical agreement observed was not satisfactory for SuperPolymyxin medium. Limitations included quantification of mucoid colonies and poor medium stability, evidenced by the growth of the susceptible control strain.

Although a recent study showed an adequate distribution of colistin in the agar, the authors reported that the culture medium could only be stored for 7 days [24]. This may have been one of the reasons for the low performance observed in the culture medium evaluated in our study. The storage conditions recommended by the manufacturer were strictly followed in the present study, but some tests were carried out after 7 days of storage.

In a screening for colistin-resistant bacteria in stool samples with SuperPolymyxin medium, 1851 phenotypically different colonies were recovered. After excluding intrinsically resistant species (n = 421), resistance to colistin was confirmed by BMD in 218 (15.3%) of the remaining 1430 isolates [28]. On the other hand, some studies have found a high sensitivity [31], and this variation may be related to the manufacturer or to the characteristics of the evaluated isolates.

Due to the wide variation in performance observed in these methods, the development of new tools, including phenotypic methods and genotyping, has been discussed [47], including selective culture media [25,27,29,48] and specific screening for detection of *mcr-1* [49]. Two phenotypic methods, colistin broth disc elution and colistin agar test, not evaluated in the present study, were described recently and included in the CLSI Guideline 2020 [32]. Although promising, they are still considered provisional, because they were established with limited discs and/or media manufacturers, endorsing the need for continuous evaluation of tests for the detection of polymyxin-resistant strains.

Conclusion

As noted by other authors, Rapid NP proved to be a method capable of identifying colistin-resistant strains in acceptable categorical agreement, but in clinical *K. pneumoniae* isolates evaluated in this work, ME and VME were considered unacceptable for colistin-resistance screening. Although very easy to perform, Colispot and SuperPolymyxin medium methods did not show acceptable sensitivity, specificity or categorical agreement. All methods have failed to detect resistance at low levels, which requires a more careful analysis in these cases, so avoiding the use of an ineffective therapeutic agent to treat an infection.

Conflicts of interest

The authors have stated that there are no conflicts of interest to disclose.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.nmni.2020.100722.

References

- [I] Potter RF, D'Souza AW, Dantas G. The rapid spread of carbapenemresistant Enterobacteriaceae. Drug Resist Updat 2016;29:30–46.
- [2] Poirel L, Jayol A, Bontron S, Villegas MV, Ozdamar M, Türkoglu S, et al. The mgrB gene as a key target for acquired resistance to colistin in Klebsiella pneumoniae. J Antimicrob Chemother 2015;70:75–80.
- [3] Cheong HS, Kim SY, Wi YM, Peck KR, Ko KS. Colistin heteroresistance in Klebsiella pneumoniae isolates and diverse mutations of PmrAB and PhoPQ in resistant subpopulations. J Clin Med 2019;8: 1444.
- [4] Aires CAM, Pereira PS, Asensi MD, Carvalho-Assef APDA. mgrB mutations mediating polymyxin B resistance in Klebsiella pneumoniae isolates from rectal surveillance swabs in Brazil. Antimicrob Agents Chemother 2016:60:6969–72.
- [5] Olaitan AO, Morand S, Rolain JM. Mechanisms of polymyxin resistance: acquired and intrinsic resistance in bacteria. Front Microbiol 2014;5: 1–18.
- [6] Cain AK, Boinett CJ, Barquist L, Dordel J, Fookes M, Mayho M, et al. Morphological, genomic and transcriptomic responses of Klebsiella pneumoniae to the last-line antibiotic colistin. Sci Rep 2018;8:1–11.
- [7] Cheng YH, Lin TL, Lin YT, Wang JT. Amino acid substitutions of Crrb responsible for resistance to colistin through CrrC in Klebsiella pneumoniae. Antimicrob Agents Chemother 2016;60:3709–16.
- [8] Wright MS, Suzuki Y, Jones MB, Marshall SH, Rudin SD, Van Duin D, et al. Genomic and transcriptomic analyses of colistin-resistant clinical isolates of Klebsiella pneumoniae reveal multiple pathways of resistance. Antimicrob Agents Chemother 2015;59:536–43.
- [9] Liu YY, Wang Y, Walsh TR, Yi LX, Zhang R, Spencer J, et al. Emergence of plasmid-mediated colistin resistance mechanism MCR-1 in animals and human beings in China: a microbiological and molecular biological study. Lancet Infect Dis 2016;16:161–8.
- [10] Xavier BB, Lammens C, Ruhal R, Kumar-Singh S, Butaye P, Goossens H, et al. Identification of a novel plasmid-mediated colistinresistance gene, mcr-2, in *Escherichia coli*, Belgium. Euro Surveill 2016;21:1–6.
- [11] Yin W, Li H, Shen Y, Liu Z, Wang S, Shen Z, et al. Novel plasmid-mediated colistin resistance gene mcr-3 in *Escherichia coli*. MBio 2017;8:1–6.
- [12] Carattoli A, Villa L, Feudi C, Curcio L, Orsini S, Luppi A, et al. Novel plasmid-mediated colistin resistance mcr-4 gene in Salmonella and Escherichia coli, Italy 2013, Spain and Belgium, 2015 to 2016. Eurosurveillance 2017;22:1–5.
- [13] Borowiak M, Fischer J, Hammerl JA, Hendriksen RS, Szabo I, Malorny B. Identification of a novel transposon-associated phosphoethanolamine transferase gene, mcr-5, conferring colistin resistance in d-tartrate fermenting Salmonella enterica subsp. enterica serovar Paratyphi B. J Antimicrob Chemother 2017;72:3317—24.
- [14] AbuOun M, Stubberfield EJ, Duggett NA, Kirchner M, Dormer L, Nunez- Garcia J, et al. mcr-1 and mcr-2 variant genes identified in Moraxella species isolated from pigs in Great Britain from 2014 to 2015. J Antimicrob Chemother 2017;72:2745–9.
- [15] Yang YQ, Li YX, Lei CW, Zhang AY, Wang HN. Novel plasmid-mediated colistin resistance gene mcr-7.1 in Klebsiella pneumoniae. J Antimicrob Chemother 2018;73:1791-5.

- [16] Wang X, Wang Y, Zhou Y, Li J, Yin W, Wang S, et al. Emergence of a novel mobile colistin resistance gene, mcr-8, in NDM-producing Klebsiella pneumoniae article. Emerg Microbe. Infect 2018;7:1–9.
- [17] Carroll LM, Gaballa A, Guldimann C, Sullivan G, Henderson LO, Wiedmanna M. Identification of novel mobilized colistin resistance gene mcr-9 in a multidrug-resistant, colistin-susceptible Salmonella enterica Serotype Typhimurium isolate. MBio 2019;10:1–6.
- [18] Aires AM, da Conceição-Neto OCC, Oliveira TRT, Dias CF, Montezzi LF, Picão RC, Albano RM, et al. Emergence of the plasmidmediated mcr-1 gene in clinical KPC-2-producing Klebsiella pneumoniae sequence type 392 in Brazil. Antimicrob Agents Chemother 2017;61:1–3.
- [19] Dalmolin TV, Martins AF, Zavascki AP, de Lima-Morales D, Barth AL. Acquisition of the mcr-I gene by a high-risk clone of KPC-2-producing Klebsiella pneumoniae ST437/CC258, Brazil. Diagn Microbiol Infect Dis 2018;90:132–3.
- [20] Turnidge J, Sei K, Mouton J. Polymyxin susceptibility testing and breakpoint setting. Adv Exp Med Biol 2019;117–32. https://doi.org/10. 1007/978-3-030-16373-0_9.
- [21] Humphries RM. Susceptibility testing of the polymyxins: where are we now? Pharmacotherapy 2015;35:22–7.
- [22] Simar S, Sibley D, Ashcraft D, Pankey G. Colistin and polymyxin B minimal inhibitory concentrations determined by etest found unreliable for Gram-negative bacilli. Ochsner J 2017;17:239–42.
- [23] Maalej SM, Meziou MR, Rhimi FM, Hammami A. Comparison of disc diffusion, Etest and agar dilution for susceptibility testing of colistin against Enterobacteriaceae. Lett Appl Microbiol 2011;53:546–51.
- [24] Turlej-Rogacka A, Xavier BB, Janssens L, Lammens C, Zarkotou O, Pournaras S, et al. Evaluation of colistin stability in agar and comparison of four methods for MIC testing of colistin. Eur J Clin Microbiol Infect Dis 2018;37:345–53.
- [25] Nordmann P, Jayol A, Poirel L. Rapid detection of polymyxin resistance in *Enterobacteriaceae*. Emerg Infect Dis 2016;22:1038–43.
- [26] Jouy E, Haenni M, Le Devendec L, Le Roux A, Châtre P, Madec JY, et al. Improvement in routine detection of colistin resistance in *E. coli* isolated in veterinary diagnostic laboratories. J Microbiol Methods 2017;132:125–7.
- [27] Nordmann P, Jayol A, Poirel L. A universal culture medium for screening polymyxin-resistant Gram-negative isolates. J Clin Microbiol 2016;54:1395–9.
- [28] Przybysz SM, Correa-Martinez C, Köck R, Becker K, Schaumburg F. SuperPolymyxinTM medium for the screening of colistin-resistant Gram-negative bacteria in stool samples. Front Microbiol 2018;9:1–7.
- [29] Abdul Momin MHF, Bean DC, Hendriksen RS, Haenni M, Phee LM, Wareham DW. CHROMagar COL-APSE: a selective bacterial culture medium for the isolation and differentiation of colistin-resistant Gramnegative pathogens. J Med Microbiol 2017;66:1554–61.
- [30] Humphries RM, Green DA, Schuetz AN, Bergman Y, Lewis S, Yee R, et al. Multicenter evaluation of colistin broth disk elution and colistin agar test: a report from the clinical and laboratory standards institute. I Clin Microbiol 2019:57:1–10.
- [31] Girlich D, Naas T, Dortet L. Comparison of the superpolymyxin and chromid colistin R screening media for the detection of colistinresistant *Enterobacteriaceae* from spiked rectal swabs. Antimicrob Agents Chemother 2019;63:1–8.
- [32] CLSI. Performance standards for antimicrobial susceptibility testing. 30th ed. Wayne, PA: Clinical and Laboratory Standards Institute; 2020. CLSI supplement M100.
- [33] European Committee on Antimicrobial Susceptibility Testing. Recommendations for MIC determination of colistin (polymyxin E) as recommended by the joint CLSI-EUCAST Polymyxin Breakpoints Working Group. EUCAST; 2016. Available from: http://www.eucast.org/fileadmin/src/media/PDFs/EUCAST_files/General_documents/Recommendations_for_MIC_determination_of_colistin_March_2016.pdf. [Accessed 3 May 2020].

- [34] Carrol KC, Pfaeller MA, Landry ML, McAdam AJ, Patel R, Richter SS, et al. Manual of clinical microbiology. 12th ed. Washington, DC: American Society of Microbiology; 2012.
- [35] International Organization for Standards. Susceptibility testing of infectious agents and evaluation of performance of antimicrobial susceptibility test devices. 2nd edition. Geneva, Switzerland: International Organization for Standards; 2019. ISO 20776-1.
- [36] The European Committee on Antimicrobial Susceptibility Testing. Breakpoint tables for interpretation of MICs and zone diameters. 2020. Available from:, version 10.0. http://www.eucast.org/clinical_breakpoints/.
- [37] Patel JB, Sharp S, Novak-Weekley S. Verification of antimicrobial susceptibility testing methods: a practical approach. Clin Microbiol Newsl 2013;35:103-9.
- [38] CLSI. Development of in vitro susceptibility testing criteria and quality control parameters. In: CLSI guideline M23. 5th ed. Wayne, PA: Clinical and Laboratory Standards Institute; 2018.
- [39] WHO. Critically important antimicrobials for human medicine. 5th revision. Geneva: WHO; 2017. Licence: CCBY-NC-SA 3.0 IGO. Available from: http://apps.who.int/iris/bitstream/10665/255027/1/9789241512220-eng.pdf?ua=1%0Ahttp://www.oalib.com/references/1801489%0Ahttp://apps.who.int/iris/bitstream/10665/77376/1/9789241504485_eng.pdf.
- [40] Dalmolin TV, Dias GÁ, de Castro LP, Ávila H, Magagnin CM, Zavascki AP, et al. Detection of Enterobacterales resistant to polymyxins using Rapid Polymyxins NP test. Braz J Microbiol 2019;50: 425–8.
- [41] Malli E, Florou Z, Tsilipounidaki K, Voulgaridi I, Stefos A, Xitsas S, et al. Evaluation of rapid polymyxin NP test to detect colistin-resistant Klebsiella pneumoniae isolated in a tertiary Greek hospital. I Microbiol Methods 2018;153:35–9.

- [42] Jayol A, Kieffer N, Poirel L, Guérin F, Güneser D, Cattoir V, et al. Evaluation of the Rapid Polymyxin NP test and its industrial version for the detection of polymyxin-resistant Enterobacteriaceae. Diagn Microbiol Infect Dis 2018:92:90—4.
- [43] Yainoy S, Hiranphan M, Phuadraksa T, Eiamphungporn W, Tiengrim S, Thamlikitkul V. Evaluation of the Rapid Polymyxin NP test for detection of colistin susceptibility in *Enterobacteriaceae* isolated from Thai patients. Diagn Microbiol Infect Dis 2018;92:102–6.
- [44] Lai CC, Lin YT, Lin YT, Lu MC, Shi ZY, Chen YS, et al. Clinical characteristics of patients with bacteraemia due to the emergence of mcr-1-harbouring *Enterobacteriaceae* in humans and pigs in Taiwan. Int | Antimicrob Agents 2018;52:651–7.
- [45] Eiamphungporn W, Yainoy S, Jumderm C, Tan-arsuwongkul R, Tiengrim S, Thamlikitkul V. Prevalence of the colistin resistance gene mcr-1 in colistin-resistant *Escherichia coli* and *Klebsiella pneumoniae* isolated from humans in Thailand. J Glob Antimicrob Resist 2018;15: 32–5
- [46] Liu X, Wang Y, Cui L, Li Y, Xue F, Liu J, et al. A retrospective study on mcr-1 in clinical Escherichia coli and Klebsiella pneumoniae isolates in China from 2007 to 2016. J Antimicrob Chemother 2018;73:1786–90.
- [47] Bardet L, Rolain JM. Development of new tools to detect colistinresistance among Enterobacteriaceae strains. Can J Infect Dis Med Microbiol 2018;2018. https://doi.org/10.1155/2018/3095249.
- [48] Bardet L, Le Page S, Leangapichart T, Rolain JM. LBJMR medium: a new polyvalent culture medium for isolating and selecting vancomycin and colistin-resistant bacteria. BMC Microbiol 2017;17:1–10.
- [49] Jayol A, Nordmann P, Lehours P, Poirel L, Dubois V. Comparison of methods for detection of plasmid-mediated and chromosomally encoded colistin resistance in *Enterobacteriaceae*. Clin Microbiol Infect 2018;24:175–9.