



Comparison of the Superpolymyxin and ChromID Colistin R Screening Media for the Detection of Colistin-Resistant *Enterobacteriaceae* from Spiked Rectal Swabs

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ABSTRACT The dissemination of carbapenemase-producing *Enterobacteriaceae* (CPE) has led to the increased use of colistin, which has resulted in the emergence of colistin-resistant *Enterobacteriaceae* worldwide. One of the most threatening scenarios is the dissemination of colistin resistance in CPE, particularly the plasmid-encoded resistance element MCR. Thus, it has now become mandatory to possess reliable media to screen for colistin-resistant Gram-negative bacterial isolates, especially *Enterobacteriaceae*. In this study, we evaluated the performances of the Superpolymyxin medium (ELITechGroup) and the ChromID Colistin R medium (bioMérieux) to screen for colistin-resistant *Enterobacteriaceae* from spiked rectal swabs. Stool samples were spiked with a total of 94 enterobacterial isolates (*Escherichia coli*, *Klebsiella pneumoniae*, *Salmonella enterica*, *Enterobacter cloacae*), including 53 colistin-resistant isolates. ESwabs (Copan Diagnostics) were then inoculated with those spiked fecal suspensions, and culture proceeded as recommended by both manufacturers. The sensitivity of detection of colistin-resistant *Enterobacteriaceae* was 86.8% (95% confidence interval [95% CI] = 74.0% to 94.0%) using both the Superpolymyxin medium and the ChromID Colistin R plates. Surprisingly, the isolates that were not detected were not the same for both media. The specificities were high for both media, at 97.9% (95% CI = 87.3% to 99.9%) for the Superpolymyxin medium and 100% (95% CI = 90.4% to 100%) for the ChromID Colistin R medium. Both commercially available media, ChromID Colistin R and Superpolymyxin, provide useful tools to screen for colistin-resistant *Enterobacteriaceae* from patient samples (rectal swabs) regardless of the level and mechanism of colistin resistance.

KEYWORDS MCR, polymyxin, sensitivity, specificity

Colistin and polymyxin B represent some of the few remaining options for the treatment of infections caused by multidrug-resistant and extremely drug-resistant Gram-negative bacteria, especially carbapenemase-producing *Enterobacteriaceae* (CPE) (1). Uncertainty remains over the best treatment option that should be used to manage infections caused by CPE. Treatment with carbapenem in combination with amikacin and treatment with colistin have achieved therapeutic results in some cases (2). Unfortunately, due to the dissemination of CPE, the increased use of colistin has led to the emergence of colistin-resistant *Enterobacteriaceae* worldwide (3). Colistin is a cationic antimicrobial peptide that interacts with the lipid A moiety of the lipopolysaccharide (LPS), disrupting the negatively charged outer membrane of Gram-negative bacteria. In Gram-negative bacteria, the main resistance mechanisms consist of LPS modification through the addition of positively charged 4-amino-4-deoxy-L-arabinose or phosphoethanolamine. In *Enterobacteriaceae*, the operons encoding enzymes in-

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volved in these modifications are *arnBCADTEF* and *pmrCAB*, respectively (4–6). Activation of the LPS-modifying genes is associated with chromosome-encoded resistance mechanisms, such as mutations in the PmrA/PmrB or PhoP/PhoQ two-component system, or through alterations to the master regulator MgrB (5, 6). In 2016, the expression of a plasmid-encoded phosphoethanolamine transferase, named MCR-1, was described as being involved in colistin resistance in *Enterobacteriaceae* (7). Since then, eight families of *mcr* genes (*mcr-1* to *mcr-8*) have been assigned, and descriptions of seven were published previously (7–13). One of the most threatening scenarios is the wide dissemination of *mcr* in CPE isolates, again limiting therapeutic options. In addition, with (i) the rapid rise of *mcr* variants and (ii) the probability that an unknown number of polymyxin resistance mechanisms are as yet unidentified, the use of molecular techniques for the identification and the screening of colistin-resistant isolates is not universally possible. Accordingly, it has now become mandatory to possess reliable media to screen for colistin-resistant isolates (3).

Superpolymyxin and ChromID Colistin R are ready-to-use selective agar media designed for the screening for colistin resistance in Gram-negative bacteria. The target microorganisms are *Enterobacteriaceae* (mostly *Escherichia coli*, *Klebsiella pneumoniae*, *Salmonella* spp., and *Enterobacter* spp.) for both media and *Acinetobacter* spp. and *Pseudomonas aeruginosa* for the Superpolymyxin medium only. ChromID Colistin R is a chromogenic medium that distinguishes *E. coli* (pink), *Klebsiella* spp., *Enterobacter* spp., *Serratia* spp. (blue), and *Salmonella* spp. (colorless), while Superpolymyxin contains eosin Y and methylene blue dyes, which help to distinguish lactose-positive organisms (purple) from lactose nonfermenters (colorless). Both media are claimed to work on bacterial cultures, stool samples, rectal swabs (cecal samples from poultry, pigs, and calves might also be used). The present study aimed to compare the performance of these media with a collection of well-characterized colistin-resistant *Enterobacteriaceae* spiked into stool samples at different concentrations and inoculated onto swabs mimicking rectal swab samples.

RESULTS

The sensitivities for the detection of colistin-resistant *Enterobacteriaceae* were 86.8% (50% confidence interval [CI] = 74.0% to 94.0%) and 84.9% (95% CI = 71.8% to 92.8%) using the Superpolymyxin medium and a ChromID Colistin R plate, respectively, after 24 h of incubation. The sensitivity of both media was the same after 48 h of incubation (86.8% [95% CI = 74.0% to 94.0%]). Surprisingly, the isolates that were not detected were not the same for both media (Table 1). The specificities were high for both media, at 97.5% (95% CI = 85.6% to 99.9%) and 100% (95% CI = 89.3% to 100%) for the Superpolymyxin medium and the ChromID Colistin R medium, respectively. Overall, the ChromID Colistin R medium performed slightly better with *K. pneumoniae* and *Salmonella enterica* than the Superpolymyxin medium, with sensitivities of 100% (50% CI = 85.0% to 100%) and 96.2% (50% CI = 78.4% to 99.8%), respectively, and specificities of 100% (50% CI = 80.8% to 100%) and 87.0% (50% CI = 65.3% to 96.6%), respectively. Conversely, ChromID Colistin R did not detect 7/25 colistin-resistant *E. coli* isolates, while only 4 strains did not grow on Superpolymyxin (Table 1). The lack of detection was not correlated with the colistin MICs or the presence or absence of *mcr*-like genes (Table 1). For colistin-resistant isolates detected on both media (14 *E. coli* isolates, 24 *K. pneumoniae* isolates, and 1 *S. enterica* isolate), the limit of detection (LOD) was at least 1 log lower for ChromID Colistin R for 69.2% (27/39) of the isolates, equivalent for both media for 20.5% (8/39) of the isolates, and at least 1 log better for the Superpolymyxin medium for 7.7% (3/39) of the tested isolates (all *E. coli*). This lower LOD of the ChromID Colistin R protocol might be the result of the 4-h enrichment step in colistin-supplemented broth. In order to decipher whether such an enrichment step might increase the performance of the Superpolymyxin medium, the seven colistin-resistant isolates which did not grow on the Superpolymyxin medium were subjected to an enrichment step similar to that performed for the ChromID Colistin R protocol. This additional step did not allow them to grow on the Superpolymyxin medium,

TABLE 1 Limit of detection of colistin-resistant *Enterobacteriaceae* on ChromID Colistin R and Superpolymyxin media

Colistin susceptibility and species	Strain name	Colistin MIC (mg/liter)	Plasmid or chromosome encoded ^b	Mechanism	Colistin resistance		Lowest LOD (CFU/ml) in ^c :		Spiked stools	Reference or source		
					ChromID Colistin R	Superpolymyxin	ESwab Amies buffer				ChromID Colistin R	Superpolymyxin
							ChromID Colistin R	Superpolymyxin				
<i>Escherichia coli</i>	CNR 111 J7	16	Chr	PmrB mutations (D14N, S71C, V83A)	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁴	21		
	CNR 20160039	4	Chr	Unknown	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	21		
	CNR 20160235	8	Chr	MgrB mutation (V8A)	1 × 10 ⁵	>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	21		
	CNR 1728	8	Chr	PmrB mutation (G160E)	1 × 10 ⁶	1 × 10 ⁴	1 × 10 ⁸	1 × 10 ⁶	1 × 10 ⁶	21		
	41489	4	P	<i>mcr-1</i>	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁷	1 × 10 ⁷	21		
	J53 + <i>mcr-1f</i>	8	P	<i>mcr-1</i>	1 × 10 ⁶	1 × 10 ⁴	1 × 10 ⁸	1 × 10 ⁶	1 × 10 ⁶	21		
	CNR20140385	4	P	<i>mcr-1</i>	>1 × 10 ⁶	1 × 10 ⁴	>1 × 10 ⁸	1 × 10 ⁶	1 × 10 ⁶	21		
	S08-056	4	P	<i>mcr-1</i>	1 × 10 ⁴	1 × 10 ³	1 × 10 ⁶	1 × 10 ⁵	1 × 10 ⁵	21		
	CNR 117 G7	4	P	<i>mcr-1</i>	>1 × 10 ⁶	1 × 10 ⁴	>1 × 10 ⁸	1 × 10 ⁶	1 × 10 ⁶	22		
	CNR 121 G9	4	P	<i>mcr-1</i>	1 × 10 ⁶	1 × 10 ⁵	1 × 10 ⁸	1 × 10 ⁷	1 × 10 ⁷	23		
	R12 F5	4	P	<i>mcr-2</i>	1 × 10 ³	>1 × 10 ⁶	1 × 10 ⁵	>1 × 10 ⁸	>1 × 10 ⁸	11		
	CNR 1745	4	P	<i>mcr-1</i>	>1 × 10 ⁶	1 × 10 ⁴	>1 × 10 ⁸	1 × 10 ⁶	1 × 10 ⁶	21		
	CNR 1604	4	P	<i>mcr-1</i>	1 × 10 ⁶	1 × 10 ⁴	1 × 10 ⁸	1 × 10 ⁶	1 × 10 ⁶	21		
	CNR 1790	4	P	<i>mcr-1</i>	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁵	21		
	CNR 1859	4	P	<i>mcr-1</i>	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁵	21		
	CNR 1886	4	P	<i>mcr-1</i>	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁴	21		
	TOP10 + <i>mcr-5f</i>	8	P	<i>mcr-5</i>	1 × 10 ²	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁷	1 × 10 ⁷	21		
	4222	4	P	<i>mcr-1</i>	1 × 10 ²	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	21		
	4070	4	P	<i>mcr-1</i>	1 × 10 ³	1 × 10 ³	1 × 10 ³	1 × 10 ⁶	1 × 10 ⁶	21		
	979	4	P	<i>mcr-1</i>	1 × 10 ³	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	21		
	6383	4	P	<i>mcr-1.5</i>	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	21		
	1724	4	P	<i>mcr-1</i>	1 × 10 ⁴	1 × 10 ³	1 × 10 ³	1 × 10 ⁶	1 × 10 ⁶	21		
	1670	4	P	<i>mcr-1.5</i>	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁷	1 × 10 ⁷	21		
36070	8	P	<i>mcr-3.2</i>	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁷	1 × 10 ⁷	24			
CNR 164 A5	4	P	<i>mcr-1</i>	1 × 10 ^{5c}	>1 × 10 ⁶	1 × 10 ⁷	>1 × 10 ⁸	>1 × 10 ⁸	This study			
CNR 20140042	16	Chr	MgrB N42Y and K43I	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 20140661	64	Chr	MgrB Q30 stop	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20151119	64	Chr	MgrB L4 stop	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20150622	64	Chr	MgrB Y41 stop	1 × 10 ²	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 20150777	128	Chr	MgrB Y41 stop	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 20150944	64	Chr	MgrB modified sequence starting at aa ^d	1 × 10 ²	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 20150309	64	Chr	MgrB modified sequence starting at aa ^d	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20150675	64	Chr	<i>mgrB</i> truncated in ORF ^e by IS10	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20140483	32	Chr	<i>mgrB</i> truncated in ORF by IS1F-like	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20140563	64	Chr	<i>mgrB</i> truncated in ORF by IS1R	1 × 10 ²	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 20150050	32	Chr	<i>mgrB</i> truncated in promoter by IS1R	1 × 10 ³	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 20140591	64	Chr	<i>mgrB</i> truncated in ORF by IS5-like	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20140550	32	Chr	<i>mgrB</i> truncated in promoter by IS903D	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20151285	32	Chr	<i>mgrB</i> truncated in ORF by IS903-like	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
S14-002	64	Chr	<i>mgrB</i> truncated in promoter by IS <i>kpn14</i>	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20140101	32	Chr	<i>ΔmgrB</i>	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 2015007	32	Chr	<i>ΔmgrB</i>	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20150066	16	Chr	<i>ΔmgrB</i>	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
CNR 20151223	32	Chr	<i>ΔmgrB</i>	1 × 10 ³	1 × 10 ³	1 × 10 ³	>1 × 10 ⁶	>1 × 10 ⁸	This study			
S15	64	Chr	<i>ΔmgrB</i>	1 × 10 ²	1 × 10 ³	1 × 10 ³	1 × 10 ⁵	1 × 10 ⁵	This study			
CNR 1630	64/32	Chr	<i>mgrB</i> truncated in ORF by IS <i>kpn25</i>	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	25			
CNR 1861	16	Chr	<i>mgrB</i> truncated in ORF by IS5	1 × 10 ²	1 × 10 ⁵	1 × 10 ⁵	1 × 10 ⁷	1 × 10 ⁷	This study			
CNR 1601	32	Chr + P	PmrB mutation (T157P)	1 × 10 ³	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			
			<i>mcr-1</i> + <i>mgrB</i> truncated in ORF by IS5	1 × 10 ²	1 × 10 ⁴	1 × 10 ⁴	1 × 10 ⁶	1 × 10 ⁶	This study			

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TABLE 1 (Continued)

Colistin susceptibility and species	Strain name	Colistin MIC (mg/liter)	Plasmid or chromosome encoded ^b	Mechanism	Colistin resistance		Lowest LOD (CFU/ml) in ^c :		Reference or source		
					Colistin MIC (mg/liter)	ESwab Amies buffer	Spiked stools				
							ChromID Colistin R	Superpolymyxin		ChromID Colistin R	Superpolymyxin
<i>Salmonella enterica</i> Serovar Paratyphi B D-tartrate + S. <i>enterica</i> biotype Java Serovar Typhimurium Serovar Paratyphi B D-tartrate + S. <i>enterica</i> biotype Java	CNR 1732	4	P	<i>mcr-1</i>	1×10^3	1×10^3	1×10^3	1×10^5	1×10^5	This study	
	CNR 1853	4	P	<i>mcr-1</i>	1×10^3	1×10^3	1×10^3	1×10^5	1×10^5	This study	
	201610686	8	P	<i>mcr-1</i>	1×10^3	1×10^3	1×10^5	1×10^5	1×10^7	This study	
	CNR 1776	8	P	<i>mcr-1</i>	1×10^3	1×10^3	$>1 \times 10^6$	1×10^5	$>1 \times 10^8$	This study	
	13-SA01718	8	P	<i>mcr-5</i>	1×10^3	1×10^3	$>1 \times 10^6$	1×10^5	$>1 \times 10^8$	8	
	Colistin-susceptible <i>Enterobacteriaceae</i> (n = 41) <i>Escherichia coli</i>	TOP10	0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21
		1608071881	0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21
		1608072264	0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21
1608073733		0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1608073228		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1608078635		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1608078858		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1608062671		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1608064819		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
2H6		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
LAN 10.48		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
VER 9.39		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1F1		0.25			$>1 \times 10^6$	$>1 \times 10^6$	1×10^6	$>1 \times 10^8$	$>1 \times 10^8$	21	
1A6		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1A8		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
2A1		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
2D9		0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
2C4		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
2D5		0.25			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	21	
1609056413		0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study	
1609061149		1			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study	
2 E8		0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study	
2 I4	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
2 F1	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
2 I5	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
3 B4	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
3 B7	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
1 B6	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
CNR 173 F9	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
1 C9	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
1 E3	1			$>1 \times 10^6$	$>1 \times 10^6$	1×10^4	$>1 \times 10^8$	1×10^6	This study		
2 B1	1			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
CNR 173 E3	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
2 C6	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		
2 D2	0.5			$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^6$	$>1 \times 10^8$	$>1 \times 10^8$	This study		

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TABLE 1 (Continued)

Colistin susceptibility and species	Strain name	Colistin MIC (mg/liter)	Colistin resistance		Lowest LOD (CFU/ml) in ^a :				Reference or source
			Plasmid or chromosome encoded ^b	Mechanism	ESwab Amies buffer		Spiked stools		
					ChromID Colistin R	Superpolymyxin	ChromID Colistin R	Superpolymyxin	
<i>Salmonella enterica</i> 4,12i:—	201604739	1			>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	This study
Serovar Enteritidis	201608919	1			>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	This study
Serovar Typhimurium	201606509	1			>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	This study
Serovar Enteritidis	201607559	0.5			>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	This study
Serovar Venezuela	201610299	0.5			>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	This study
<i>Enterobacter cloacae</i>	CNR 131 G4	0.5	P	<i>mcr-4.2</i>	>1 × 10 ⁶	>1 × 10 ⁶	>1 × 10 ⁸	>1 × 10 ⁸	This study

^aUnderlined CFU counts are considered negative results. The sensitivity was 86.8% (95% CI = 74.0% to 94.0%) for both media after 48 h of incubation. The specificity was 100% (95% CI = 89.3% to 100%) and 97.5 (95% CI = 85.6% to 99.9%), respectively, for ChromID Colistin R and Superpolymyxin.

^bP, plasmid; Chr, chromosome.

^cAfter 48 h of incubation (no colony at 24 h).

^daa, amino acid.

^eORF, open reading frame.

^f*E. coli* J53 was a recombinant strain harboring the *mcr-1* gene on plasmid pDM1 (41522) (21). *E. coli* TOP10 was a recombinant strain harboring the *mcr-5* gene on plasmid pSE13-SA01718 (21).

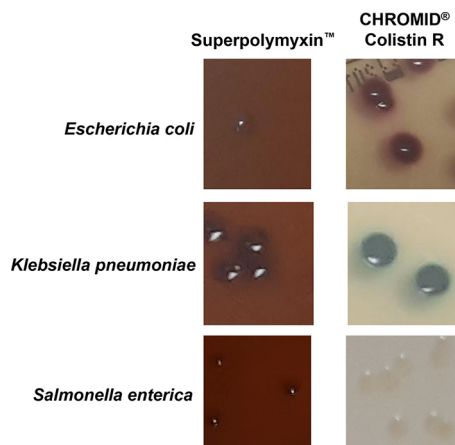


FIG 1 Morphological aspect of colonies of *E. coli*, *K. pneumoniae*, and *Salmonella enterica* grown on Superpolymyxin and ChromID Colistin R media.

suggesting that this enrichment should not be recommended for use with this selective medium. As previously reported by Jayol et al. for the Superpolymyxin medium, prolongation of the incubation from 24 to 48 h did not modify the performance of the Superpolymyxin medium (14). Regarding ChromID Colistin R, prolongation of the incubation time to 48 h for one *Mcr-1*-producing *E. coli* isolate (strain CNR 164 A5) allowed us to identify typical pink colonies that were barely detectable at 24 h of incubation. Finally, one *Enterobacter cloacae* isolate positive for *mcr-4.2* was not detected by either medium. As previously described for *mcr-3* and *mcr-4* variants of CPE isolates (15), the presence of *mcr-4.2* does not confer phenotypic resistance to polymyxins in this *E. cloacae* isolate (colistin MIC, 0.5 mg/liter).

DISCUSSION

Based on this study performed with spiked rectal swabs, ChromID Colistin R and Superpolymyxin selective media showed very similar performances. The main advantage of the Superpolymyxin medium is that it could be directly inoculated with the rectal swabs without any enrichment step (4 h) in colistin-supplemented broth, whereas ChromID Colistin R requires an enrichment step. On the other hand, the main advantage of ChromID Colistin R lies in the use of chromogenic molecules enabling the rapid presumed identification of growing colonies (pink for *E. coli*, blue for *Klebsiella*, *Enterobacter*, and *Serratia*, and white for *Salmonella*). Indeed, the morphological aspect of the colonies on the Superpolymyxin medium was indistinguishable between *E. coli*, *K. pneumoniae*, and *Salmonella enterica* (Fig. 1). As species cannot easily be differentiated on Superpolymyxin, clinical labs must then identify the growing colonies before reporting results. In our study, the selectivity of both media was good, since no Gram-positive bacteria or fungi grew on them.

Of note, unlike the ChromID Colistin R medium, which is currently limited to use with *Enterobacteriaceae*, the Superpolymyxin medium is also claimed to be able to detect colistin resistance in all Gram-negative bacteria, including *Acinetobacter* spp. and *P. aeruginosa*. Accordingly, we tested the Superpolymyxin medium with three colistin-resistant isolates (all producing the OXA-23 carbapenemase) and four colistin-susceptible *Acinetobacter baumannii* isolates. In all three colistin-resistant isolates, a mutation of PmrB (A226T, A226V, and R263H) resulted in MICs ranging from 16 to 64 mg/liter. The Superpolymyxin medium fully detected all colistin-resistant isolates, while none of the four susceptible strains grew on the medium.

As the rate of colistin resistance is likely to increase in the near future, clinical microbiology laboratories will require rapid and reliable screening media to identify carriers in hospital settings. Here, we have shown that both commercially available media, ChromID Colistin R and Superpolymyxin, are useful tools to screen for colistin-

resistant *Enterobacteriaceae* from patient samples (rectal swabs) regardless of the level and mechanism of colistin resistance.

MATERIALS AND METHODS

Susceptibility testing. MICs were determined by broth microdilution according to the guidelines of a CLSI and EUCAST joint subcommittee (16). Results were interpreted using EUCAST breakpoints, as updated in 2018.

Bacterial isolates. Ninety-four enterobacterial isolates, including 53 isolates exhibiting resistance to colistin (MIC > 2 mg/liter), were tested. The colistin resistance mechanism of all these isolates has been characterized at the molecular level (Table 1). The tested isolates were as follows: colistin-resistant isolates with colistin MICs of ≥ 4 mg/liter, consisting of *Escherichia coli* ($n = 25$, including 20 isolates carrying *mcr* genes), *Klebsiella pneumoniae* ($n = 25$, including 3 isolates carrying *mcr* genes), and *Salmonella enterica* ($n = 3$ isolates carrying *mcr* genes); colistin-susceptible *E. coli* ($n = 19$), *K. pneumoniae* ($n = 16$), and *Salmonella enterica* ($n = 5$) isolates; and one *mcr-4.2*-positive *Enterobacter cloacae* isolate (Table 1). Chromosomally encoded mutations in genes responsible for colistin resistance (the *pmrA*, *pmrB*, *phoP*, *phoQ*, *mgrB*, and *crpB* genes) were also searched as described previously (17).

Spiked rectal swabs. Suspensions of bacterial strains with an optical density of a 0.5 McFarland standard (inoculum, $\sim 10^8$ CFU/ml) were serially diluted in water, and 10-fold dilutions of pure solution to 10^{-3} were used to spike liquid stools from healthy volunteers (1 g in 1 ml of sterile water), as previously described (18). The bacterial suspensions that were used to spike stools from healthy volunteers were verified by the concomitant inoculation of Mueller-Hinton agar with 10 μ l of the suspension diluted to 10^{-4} in water. Ten microliters of bacterial suspension was added to 90 μ l of stool. The totality (100 μ l) of this spiked stool was then absorbed on the ESwab and introduced into 1 ml Amies transport medium (Copan Diagnostics, Murrieta, CA, USA) to mimic true rectal swabs. Each ESwab containing stool with each dilution of bacteria was then cultured according to the recommendations of both manufacturers (see Fig. S1 in the supplemental material). Briefly, 10 microliters of the inoculated Amies medium was transferred to the Superpolymyxin agar (ELITechGroup, Puteaux, France) and spread with a plate spreader without an enrichment step. The ChromID Colistin R agar plates (bioMérieux, La Balmes-Les-Grottes, France) were inoculated after an enrichment step, as follows: 200 μ l of each inoculated Amies suspension was introduced into 10 ml of brain heart infusion (BHI) medium (bioMérieux) supplemented with one disc of colistin (10 μ g) and incubated for 4 h at 37°C before seeding of 50 μ l in dials.

Determination of LOD. The lowest limit of detection (LOD) corresponds to the minimum number of bacteria that must be present in the sample to obtain growth on selective medium. In contrast to other studies that evaluated the performance of selective media with cultured bacteria (14, 19, 20), our study was performed on inoculated rectal swabs. This involves further dilution of the spiked stool sample in the ESwab Amies buffer (Fig. S1). As indicated by the manufacturer of the Superpolymyxin medium (ELITechGroup), the threshold value for susceptible strains could not be greater than 5×10^6 CFU/ml (directly from a bacterial suspension) because susceptible bacteria could benefit from an inoculum artifact to grow on the selective medium. Accordingly, the threshold for the LOD value was set at $\geq 1 \times 10^6$ CFU/ml in ESwab Amies buffer, corresponding to an initial concentration of 1×10^8 CFU/ml in the spiked stool (Table 1; Fig. S1). A fecal suspension without addition of bacteria was used as a negative control. In addition, 10 randomly selected strains were tested by a second experimenter to assess reproducibility. In all cases the results were identical between all experimenters.

Statistical analysis. The sensitivity and specificity values with their respective 95% confidence intervals (CI) were calculated using the free software vassarStats (website for statistical computation, <http://vassarstats.net/>).

SUPPLEMENTAL MATERIAL

Supplemental material for this article may be found at <https://doi.org/10.1128/AAC.01618-18>.

SUPPLEMENTAL FILE 1, PDF file, 0.3 MB.

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We have no conflicts of interest to declare.

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