linezolid, and tigecycline. Although the clinical significance of these findings is unknown, the decline in drug effectiveness against *S. aureus* infections represents a looming threat to patient health and highlights the possibility of a return to illness and death rates similar to those before antimicrobial drugs were available.

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# **Colistin-Resistant** *Enterobacteriaceae* Carrying the *mcr-1* Gene among Patients in Hong Kong

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To the Editor: Colistin belongs to the last line of bactericidal antimicrobial drugs active against multidrugresistant gram-negative bacteria such as carbapenemaseproducing *Enterobacteriaceae* and carbapenem-resistant *Acinetobacter baumannii*. Consequently, the discovery of the plasmid-mediated colistin-resistant gene *mcr-1* in *Escherichia coli* (1) raises concern in the medical community because colistin might be useless in treating infections caused by *mcr-1*–carrying *Enterobacteriaceae*.

During December 8, 2015–January 8, 2016, we conducted prospective laboratory surveillance of mcr-1-carrying Enterobacteriaceae and Acinetobacter species in a universityaffiliated tertiary hospital serving a population of  $\approx 0.53$  million in Hong Kong, China. Clinical specimens were processed by using standard operating procedures for different specimen types (2). All Enterobacteriaceae and Acinetobacter spp. isolates were plated onto MH1 agar, which is Mueller-Hinton agar (BD Diagnostics, Sparks, MD, USA), supplemented with 1 µg/mL colistin sulfate (Sigma-Aldrich, St. Louis, MO, USA) for overnight incubation at 37°C in air. Intrinsically colistin-resistant organisms, including Proteus spp., Providencia spp., Serratia spp., and Morganella morganii, were excluded. E. coli ATCC 25922 was used as a negative control. We screened bacteria that grew on MHC1 for *mcr-1* by real-time PCR that used specific primers MCR1 22697 F1 (5'-CACT-TATGGCACGGTCTATGA-3') and MCR1 22810 R1 (5'-CCCAAACCAATGATACGCAT-3') and the hydrolysis probe MCR1 22763 Pb1 (FAM-TGGTCTCGG/ ZEN/CTTGGTCGGTCTGTAGGGC-3IABkFQ) (Integrated DNA Technologies, Coralville, IA, USA). The complete mcr-1 gene found in PCR-positive isolates was amplified and sequenced by specific primers. The colistin MIC of positive isolates was measured by using Etest strips (Bio-Mérieux, Marcy l'Etoile, France). Susceptibility to other antimicrobial drugs was determined by using the Kirby-Bauer disk diffusion method, according to Clinical and Laboratory Standards Institute guidelines (3). We retrieved clinical details of patients whose sample had mcr-1-carrying Enterobacteriaceae from the hospital clinical management system.

## LETTERS

A total of 1,324 Enterobacteriaceae and 103 Acinetobacter spp. isolates were screened on MHC1 agar and isolated from blood, urine, stool, or respiratory samples; wound swab specimens; and other sterile and nonsterile body fluids, tissues, or swab specimens. Of the total 1,427 isolates, 62(4.3%)grew on MHC1: 26 E. coli, 24 Klebsiella spp., 7 Enterobacter spp., 4 Salmonella spp., and 1 Citrobacter sp. Among these 62 isolates, 1 Enterobacter cloaceae and 4 E. coli isolates were mcr-1 positive. All gene sequences were 100% identical to that of mcr-1 in E. coli strain SHP45 (GenBank accession no. KP347127), which was isolated from a pig farm specimen in China (1,3). Of the 5 mcr-1-positive isolates, 2 were from blood cultures from patients with biliary tract infection, 1 from a mid-stream urine specimen from a patient with symptomatic urinary tract infection, and 2 from stool specimens from asymptomatic patients. The range of colistin MICs of the 5 mcr-1-positive isolates was 3-64 µg/mL; all were susceptible to carbapenem. One E. coli isolate (from patient 4) exhibited extended-spectrum  $\beta$ -lactamase activity (Table). Patient 3 resided in mainland China before this admission; patient 2 received a liver transplant in China in 2004. None of the 5 patients had a history of colistin treatment.

Finding the *mcr-1* gene in 0.4% of *Enterobacteriace-ae* clinical isolates in Hong Kong is expected because of the high proportion of livestock and meat imported from China, where prevalence of colistin-resistant isolates is up to 25.4% and 28.0% in pigs and retail chicken meat,

respectively (5,6). Our findings highlight several issues. We noted a wide range (3–64  $\mu$ g/mL) of colistin MICs in the *mcr-1*–carrying *Enterobacteriaceae*; the *E. cloacae* isolate exhibited the highest MIC. This wide variation in MICs has been noted previously (1,7). Whether the variation results from the differential expression of the *mcr-1* gene or from potentially unidentified colistin resistance mechanisms co-existing in some isolates is unknown (8).

Our discovery of the *mcr-1* gene in an *E. cloacae* isolate adds diversity to the *Enterobacteriaceae* species known to be *mcr-1* carriers (e.g., *E. coli, Klebsiella pneumoniae,* and *Salmonella* sp.) (1,9). An in vitro study showed transfer of *mcr-1*–carrying pHNSHP45 (i.e., polymyxin-resistant plasmid) to *Pseudomonas aeruginosa* (1). Consequently, surveillance for the *mcr-1* gene should include all clinically relevant *Enterobacteriaceae* species, and screening for other gram-negative organisms (e.g., *P. aeruginosa*) infecting humans should be considered.

We show a potential workflow for screening *mcr-1* isolates by sequential use of MHC1 agar and real-time PCR. Clinical and Laboratory Standards Institute guidelines have no recommended colistin breakpoints for *Enterobacteriaceae* (3); however, the European Committee on Antimicrobial Susceptibility Testing recommends a breakpoint of  $\geq 4$ µg/mL to define colistin resistance in *Enterobacteriaceae* (10). Given that some *mcr-1*–positive isolates may have a colistin MIC of 2 µg/mL (1,7), which is lower than the

Table. Clinical details of 5 patients infected with mcr-1–carrying Enterobacteriaceae, Hong Kong*					
		Time from admission to	Antimicrobial drug		mcr-1-positive
Patient ID†		collection of specimen	use <1 mo before		species (colistin MIC,
(age, y)	Underlying conditions	(specimen type)	isolation	Outcome	µg/mL)
1 (55)	Acute myeloid leukemia 4 mo after bone marrow transplant	4 mo (stool sample‡)	Pipericillin/ tazobactam, meropenem	Asymptomatic colonization	Enterobacter cloacae complex (64)
2 (68)	Primary sclerosing cholangitis with liver transplant in China in 2004; currently on sirolimus and prednisolone; right hepatectomy in 2008 for right diffuse ischemic bile injury; history of recurrent cholangitis	On admission with sepsis workup for biliary sepsis resulting from biliary anastomotic stricture (blood culture)	None	Recovered	Escherichia coli (3)
3 (2)	Autologous bone marrow transplant for stage IV neuroblastoma	14 d (stool sample‡)	Pipericillin/ tazobactam	Asymptomatic colonization	<i>E. coli</i> (3)
4 (57)	Hepatitis B virus–related hepatocellular carcinoma; recurrent pyogenic cholangitis; recurrent biliary sepsis with portal vein thrombosis; cerebellar stroke in 2013	On admission with sepsis workup for biliary sepsis resulting from biliary stricture and recent transarterial chemoembolization (blood culture)	None	Recovered	ESBL-producing <i>E.</i> <i>coli</i> (4)
5 (80)	Duke's B carcinoma of rectum [lower anterior resection in 1996]; carcinoma of thyroid [post-thyroidectomy]; hypertension; diabetes mellitus	On admission with sepsis workup for symptomatic urinary tract infection (mid-stream urine sample)	None	Recovered	E. coli (4)

\*ID, identifier; ESBL, extended-spectrum β-lactamase.

†Patient 4 was male; others were female.

‡Routine surveillance of stool samples for multidrug-resistant organisms according to infection control protocol (4).

recommended breakpoint, we designed MHC1 with a colistin concentration of only 1  $\mu$ g/mL to minimize false-negative results. However, some colistin-susceptible organisms might grow on MHC1 (<5% in our study), resulting in the low PCR-positive rate for *mcr-1* among isolates.

Exact epidemiology of the *mcr-1* gene is unknown, indicating a need to conduct accurate surveillance of the gene's prevalence in humans. Additional mechanisms unique to the *mcr-1* gene may contribute to colistin resistance, suggested by the wide variation in colistin MICs among *mcr-1*-carrying *Enterobacteriaceae*.

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# *Cryptococcus gattii* Meningitis Complicated by *Listeria monocytogenes* Infection

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To the Editor: Among immunocompetent persons with cryptococcal disease, infection with a second organism is rare; all reported cases have involved concomitant mycobacterial infections (1). Immunosuppression is not a necessary precondition for infection with *Cryptococcus gattii* (2), and among immunocompetent persons, *C. gattii* infection confers high mortality rates: up to 24% according to a large case series (3). In addition, cryptococcomas are frequently observed in patients with *C. gattii*, as opposed to *C. neoformans*, infection, commonly necessitating longer courses of treatment. We report a fatal case of *C. gattii* and *Listeria monocytogenes* co-infection in an immunocompetent woman with cryptococcomas.

The patient was a previously healthy 23-year-old Hispanic woman who was hospitalized in 2009 after weeks of headache and recent-onset diplopia. Lumbar puncture revealed elevated opening pressure of 52 cm H<sub>2</sub>O; elevated leukocytes (1,030 cells/µL: 31% neutrophils, 55% lymphocytes, 14% monocytes); elevated protein concentration (117 g/L); and decreased glucose concentration (30 mg/dL). Cerebrospinal fluid (CSF) cryptococcal antigen (CrAg) titer was 1:64, and culture grew C. gattii. HIV antibody test result was negative. Magnetic resonance imaging of the brain demonstrated scattered enhancing round lesions within the cerebrum and cerebellum, consistent with cryptococcomas. The patient was prescribed intravenous amphotericin B (1 mg/kg/d) and intravenous flucytosine (2 g/6 h) (Table); after 5 days of therapy, culture of a repeat lumbar puncture sample was negative. The patient was then given intravenous liposomal amphotericin at 7 mg/kg, and after a 14-day induction period she was discharged with instructions to take fluconazole orally (400 mg  $2\times/d$ ) and to continue amphotericin B infusions  $(3\times/d)$ wk) (Table).