RESEARCH LETTERS

- Suthar MS, Pulendran B. Systems analysis of West Nile virus infection. Curr Opin Virol. 2014;6:70–5. http://dx.doi.org/10.1016/ j.coviro.2014.04.010
- Diuk-Wasser MA, Brown HE, Andreadis TG, Fish D. Modeling the spatial distribution of mosquito vectors for West Nile virus in Connecticut, USA. Vector Borne Zoonotic Dis. 2006;6:283–95. http://dx.doi.org/10.1089/vbz.2006.6.283
- Dunne DW, Shaw A, Bockenstedt LK, Allore HG, Chen S, Malawista SE, et al. Increased TLR4 expression and downstream cytokine production in immunosuppressed adults compared to non-immunosuppressed adults. PLoS One. 2010;5:e11343. http://dx.doi.org/10.1371/journal.pone.0011343
- Qian F, Goel G, Meng H, Wang X, You F, Devine L, et al. Systems immunology reveals markers of susceptibility to West Nile virus infection. Clin Vaccine Immunol. 2015;22:6–16. http://dx.doi.org/ 10.1128/CVI.00508-14
- Chabierski S, Barzon L, Papa A, Niedrig M, Bramson JL, Richner JM, et al. Distinguishing West Nile virus infection using a recombinant envelope protein with mutations in the conserved fusion-loop. BMC Infect Dis. 2014;14:246. http://dx.doi.org/ 10.1186/1471-2334-14-246
- Qian F, Thakar J, Yuan X, Nolan M, Murray KO, Lee WT, et al. Immune markers associated with host susceptibility to infection with West Nile virus. Viral Immunol. 2014;27:39–47. http://dx.doi.org/10.1089/vim.2013.0074
- Montgomery RR. Age-related alterations in immune responses to West Nile virus infection. Clin Exp Immunol. 2017;187:26–34.
- Shaw AC, Goldstein DR, Montgomery RR. Age-dependent dysregulation of innate immunity. Nat Rev Immunol. 2013;13: 875–87. http://dx.doi.org/10.1038/nri3547

Address for correspondence: Ruth R. Montgomery, Department of Internal Medicine, Yale University School of Medicine, 300 Cedar St/TAC S413, New Haven, CT 06520-8031, USA; email: ruth.montgomery@yale.edu

mcr-1 in *Enterobacteriaceae* from Companion Animals, Beijing, China, 2012–2016

Lei Lei,¹ Yang Wang,¹ Stefan Schwarz, Timothy R. Walsh, Yanran Ou, Yifan Wu, Mei Li, Zhangqi Shen

Author affiliations: China Agricultural University, Beijing, China (L. Lei, Y. Wang, Y. Ou, Y. Wu, M. Li, Z. Shen); Freie Universität Berlin, Berlin, Germany (S. Schwarz); Cardiff University, Cardiff, Wales, UK (T.R. Walsh, M. Li); Iowa State University, Ames, Iowa, USA (Z. Shen)

DOI: http://dx.doi.org/10.3201/eid2304.161732

To investigate the prevalence of the recently emerging colistin resistance gene *mcr-1* in *Enterobacteriaceae* among companion animals, we examined 566 isolates collected from cats and dogs in Beijing, China, during 2012–2016. Of these isolates, 49 (8.7%) were *mcr-1*–positive.

Multidrug-resistant and extensively drug-resistant gramnegative bacteria are a major threat to public health worldwide (1,2). The recent rapid dissemination of carbapenem-resistant *Enterobacteriaceae* has worsened this situation and further narrowed treatment options for infections caused by these bacteria (3). Colistin is a last-resort drug for treating carbapenem-resistant *Enterobacteriaceae* infections (4). In 2016, we identified the mobile colistin resistance gene *mcr-1* (1). Soon after its description, *mcr-1* was observed in *Enterobacteriaceae* from humans and food-producing animals in >30 countries on 5 continents (5).

A 2016 article reported that a 50-year-old man who worked in a pet store tested positive for mcr-1-harboring *E. coli* (6). Investigation identified 6 multidrug-resistant mcr-1-producing *E. coli* isolates in samples from 4 dogs and 2 cats in the pet store, indicating that the pathogens can be transmitted between humans and companion animals. So far, the prevalence of mcr-1-containing *Enterobacte-riaceae* in companion animals is largely unknown. In our study, we focused on estimating the prevalence of mcr-1 in *Enterobacteriaceae* of companion animal origin in Beijing, China, during 2012–2016, and investigated the presence of the mcr-1 gene in pet foods purchased there.

In Beijing, the total number of registered dogs and cats is ≈ 1.2 million. We collected samples from both healthy and sick dogs and cats in Veterinary Teaching Hospital of China Agricultural University.

A total of 566 nonduplicate *Enterobacteriaceae* strains were isolated from 1,439 nasal and rectal swab samples collected from 1,254 dogs and 185 cats during 2012–2016. We also isolated 25 *Enterobacteriaceae* from 32 nasal swab samples from the pet owners. Because the food chain is among the main routes for humans and companion animals to acquire foodborne pathogens, we collected a small sample of pet foods (dog food, n = 30; cat food, n = 5) containing chicken as the main ingredient in Beijing during June–August 2016.

The species of all *Enterobacteriaceae* were determined by 16S rDNA sequencing and matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of specimens cultured on brain-heart infusion agar plates containing 2 µg/mL colistin. A total of 79/566 (14.0%) of the *Enterobacteriaceae* isolates from companion animals were resistant to colistin: 56 *E. coli*, 16 *Klebsiella pneumoniae*, 5 *Enterobacter cloacae*, 1 *Enterobacter aerogenes*, and 1 *Shigella* spp. PCR amplification of *mcr-1* indicated that 8.7% (49/566) of *Enterobacteriaceae* and 62.0% (49/79)

¹These authors contributed equally to this article.

of colistin-resistant isolates harbored the *mcr-1* gene, 47 *E. coli* and 2 *K. pneumoniae*. Only 1 *E. coli* isolate from a pet owner was colistin-resistant and *mcr-1*–positive. The proportions of *mcr-1*–containing *E. coli* per year ranged from 6.1% to 14.3% (Figure).

We examined the susceptibility of colistin-resistant *E.* coli to 8 other antimicrobial agents by agar dilution, according to the recommendations of Clinical and Laboratory Standards Institute (7). The mcr-1–carrying *E.* coli exhibited high resistance rates to ampicillin (97.9%), cefotaxime (91.5%), chloramphenicol (89.4%), and gentamicin (85.1%) (online Technical Appendix Table 1, http://wwwnc.cdc.gov/EID/article/23/4/16-1732-Techapp1.pdf) but were susceptible to imipenem. The mcr-1–positive *E.* coli were more often resistant to amoxicillin/clavulanate, ampicillin, and chloramphenicol than were the mcr-1–negative *E.* coli (p<0.05) (online Technical Appendix Table 1).

All 57 colistin-resistant *E. coli* were subjected to XbaI pulsed-field gel electrophoresis (PFGE). The 55 colistin-resistant *E. coli* strains (2 nontypeable strains were excluded) were subdivided into 33 patterns and grouped into 31 clusters (A–Z, AB–AF) (online Technical Appendix Figure). The diversity and similarity of PFGE patterns of *E. coli* from different origins suggested that the dissemination of *mcr-1* was possibly related to both clonal expansion and horizontal transmission.

Of note, the 1 *E. coli* colistin-resistant, *mcr-1*-positive isolate from a pet owner had the same PFGE pattern as 5 isolates from dogs and cats. Multilocus sequence typing linked these 6 strains to sequence type 101. These results suggest that *E. coli* strains can be exchanged between companion animals and humans.

The PCR and sequence analysis of the pet food samples showed that 7 of 35 samples were positive for *mcr-1*. Companies in China produced 5 of these foods; the other 2 were from Italy and Belgium (online Technical Appendix Table 2). These results suggest that pet foods may be a source from which intestinal bacteria of companion animals can acquire the *mcr-1* gene.

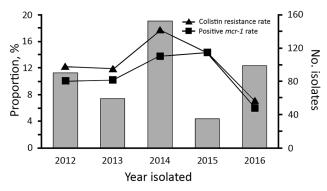


Figure. Proportion of colistin resistance and *mcr-1* in *Escherichia coli* of companion animal origin, Beijing, China, 2012–2016.

Currently, colistin is not used to treat companion animals in China. The companion animals included in this study were from an urban area of Beijing, so they had minimal or no contact with food-producing animals in which colistin may have been used. Because we found mcr-1 in pet foods, we speculate that the pet food industry may be a source of mcr-1 among companion animals. Because of frequent and close contact between humans and companion animals, our study proposes that opportunities exist to transmit colistin-resistant Enterobacteriaceae to and from both groups. Thus, colistin-resistant Enterobacteriaceae from companion animals may represent a potential risk to human health. Further surveillance and control efforts are needed to reduce colistin-resistant and mcr-1-containing Enterobacteriaceae in companion animals and food-producing animals.

This work was supported in part by National Natural Science Foundation of China (nos. 31672604, 31422055 and 81661138002). T.R. Walsh was also supported by MRC grant DETER-XDR-CHINA (MR/P007295/1).

Ms. Lei is a PhD student in the College of Veterinary Medicine, China Agricultural University. Her main interest is the prevalence of antibiotic resistance in enteric bacteria.

References

- 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. http://dx.doi.org/10.1016/S1473-3099(15)00424-7
- van der Bij AK, Pitout JD. The role of international travel in the worldwide spread of multiresistant Enterobacteriaceae. J Antimicrob Chemother. 2012;67:2090–100. http://dx.doi.org/10.1093/jac/dks214
- Temkin E, Adler A, Lerner A, Carmeli Y. Carbapenem-resistant Enterobacteriaceae: biology, epidemiology, and management. Ann N Y Acad Sci. 2014;1323:22–42. http://dx.doi.org/10.1111/nyas.12537
- Michalopoulos AS, Tsiodras S, Rellos K, Mentzelopoulos S, Falagas ME. Colistin treatment in patients with ICU-acquired infections caused by multiresistant Gram-negative bacteria: the renaissance of an old antibiotic. Clin Microbiol Infect. 2005;11: 115–21. http://dx.doi.org/10.1111/j.1469-0691.2004.01043.x
- Schwarz S, Johnson AP. Transferable resistance to colistin: a new but old threat. J Antimicrob Chemother. 2016;71:2066–70. http://dx.doi.org/10.1093/jac/dkw274
- Zhang XF, Doi Y, Huang X, Li HY, Zhong LL, Zeng KJ, et al. Possible transmission of *mcr-1*-harboring *Escherichia coli* between companion animals and human. Emerg Infect Dis. 2016;22: 1679–81. http://dx.doi.org/10.3201/eid2209.160464
- Clinical and Laboratory Standards Institute. Performance standards for antimicrobial disk and dilution susceptibility tests for bacteria isolated from animals; 3rd ed (VET01S). Wayne (PA): The Institute; 2005.

Address for correspondence: Zhangqi Shen, Beijing Advanced Innovation Center for Food Nutrition and Human Health, College of Veterinary Medicine, 2 Yuanmingyuan West Rd, Beijing 100193, China; email: szq@cau.edu.cn

mcr-1 in *Enterobacteriaceae* from Companion Animals, Beijing, China, 2012–2016

Technical Appendix

Technical Appendix Table 1. Antimicrobial drug MI	Cs of colistin-resistant Esch	nerichia coli from companion animals (r	า=56)*
mcr-1-posit	ive F coli (n = 47)	mcr-1-negative E coli (n = 9)	

	mcr - r - positive E. con (n = 47)		mcr - r - megative = 1 cont(n = 9)				
	MIC ₅₀ ,	MIC ₉₀ ,	Resistance,	MIC ₅₀ ,	MIC ₉₀ ,	Resistance,	
Antimicrobial agents†	µg/mL	µg/mL	%	µg/mL	µg/mL	%	p value‡
Colistin	8	256	100.0%	4	μg/mL	100.0%	NA
Polymyxin B	8	64	100.0%	4	16	100.0%	NA
Amoxicillin-clavulanic acid	64/32	>128/64	70.2%	16/8	16	33.3%	0.0296
Ampicillin	>256	>256	97.9%	128	64/32	66.7%	0.0111
Tigecycline	0.5	1	2.1%	0.25	>256	0.0%	NA
Enrofloxacin	32	64	74.5%	0.25	0.5	44.4%	0.1121
Cefotaxime	>256	>256	91.5%	128	128	66.7%	0.0740
Chloramphenicol	128	>256	89.4%	4	>256	22.2%	0.0002
Gentamicin	>256	>256	85.1%	64	>256	55.6%	0.0632
Imipenem	0.25	0.5	0.0%	0.25	>256	0.0%	NA

*Colistin-resistant *Escherichia coli* was isolated from 57 samples, including 1 from an owner of a companion animal that was excluded from this table; NA, not applicable.

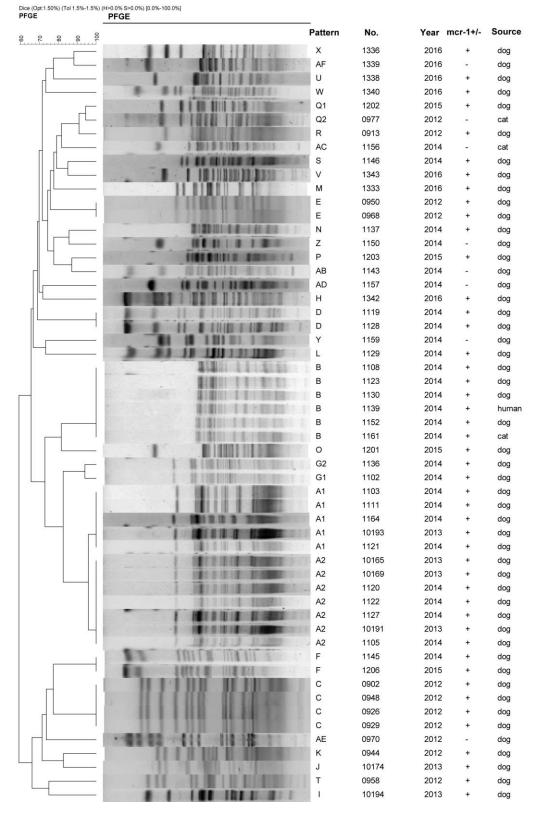
The MICs of amoxicillin-clavulanate, ampicillin, enrofloxacin, cefotaxime, chloramphenicol, gentamicin, and imipenum were interpreted according to the CLSI document VET01S 3rd ed. The MIC of colistin, polymixin B, and tigecycline were interpreted according to the European Committee on Antimicrobial Susceptibility Testing clinical breakpoints (version 6.0).

‡p values were determined by Fisher exact test.

Techncial Appendix Table 2. Characterization of pet foods and the presence of *mcr-1* gene

	Year of				
Type of pet food	production	Ingredients	Pet food sample numbers*	Origin	mcr-1-positive pet food
Dry foods	2015	Chicken meal	15–19, 22	China	16, 19
Dry foods	2015	Chicken meal	20	Belgium	20
Dry foods	2015	Chicken meal	27	Australia	0
Dry foods	2015	Chicken, fish/pork	21, 23, 27, 29, 30, 34, 35	China	21, 34, 35
Dry foods	2016	Chicken	24–26	China	0
Dry foods	2015	Chicken	31, 32	USA	0
Dry foods	2015	Fish/chicken	33	Germany	0
Semi-moist foods	2016	Chicken	1,2,5	China	0
Semi-moist foods	2016	Chicken, bovine spleen	6–8	China	0
Semi-moist foods	2016	Chicken, pork/fish	3, 4	China	0
Canned wet foods	2016	Chicken, beef, mutton	9–11	Thailand	0
Canned wet foods	2016	Chicken	12–14	Italy	14

*Total: 35 samples of pet foods. Numbers 2, 4, 5, 33, and 34 are cat food; remaining samples are dog food



Technical Appendix Figure. Xbal pulsed-field gel electrophoretic analysis of colistin-resistant *Escherichia coli.*