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EDITED AND REVIEWED BY
Jiun-Ling Wang,
National Cheng Kung University, Taiwan

*CORRESPONDENCE
Po-Yu Liu
pyliu@vghtc.gov.tw
Yao-Ting Huang
ythuang@cs.ccu.edu.tw

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Corrigendum: Comparative genomics revealed fluoroquinolone resistance determinants and OmpF deletion in carbapenem-resistant *Escherichia coli*

Wan-Ting Yang[1], I-Ju Chiu[2], Yao-Ting Huang[2]* and Po-Yu Liu[1,3,4]*

[1]Division of Infection, Department of Internal Medicine, Taichung Veterans General Hospital, Taichung, Taiwan, [2]Department of Computer Science and Information Engineering, National Chung Cheng University, Chia-Yi, Taiwan, [3]Ph.D. Program in Translational Medicine, National Chung Hsing University, Taichung, Taiwan, [4]Rong Hsing Research Center for Translational Medicine, National Chung Hsing University, Taichung, Taiwan

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A corrigendum on
Comparative genomics revealed fluoroquinolone resistance determinants and OmpF deletion in carbapenem-resistant *Escherichia coli*

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In the published article, there was an error in Table 2 as published. Column titles for TEM, CMY, and CTX were displayed as "AmpC," "ESBL," and "Broad-Spectrum Beta-Lactamase." The correct labels are "Broad-Spectrum Beta-Lactamase" for the TEM columns, "AmpC" for the CMY columns, and "ESBL" for the CTX columns. In addition,

the Elppa 2 CTX-M-5 block was displayed as "black." The correct indication is "white." The corrected Table 2 and its caption appear below.

The authors apologize for these errors and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Publisher's note

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TABLE 2 Genetic diversity, source of bacteremia, and antibiotic-resistant mechanisms identified in the *E. coli* strains.

Strain Elppa ID	ST	Phylogroup	*Fim*H Type	Serotypes		Source Of Bacteremia	Hospital Or Community-Acquired	Virulence Score	Plasmid	Carbapenemase		Broad-Spectrum Beta-Lactamase		AmpC		ESBL		Outer Membrane Porins Genes		FQ Resistant Gene
				O	H					NDM-1	OXA-48	TEM-1	TEM-135	CMY-2	CMY-29	CTX-M	CTX-M-5	*Omp*C D192G Mutation	*Omp*F 264-269 Deletion	Gyra (D87N)
7	44	A	54	O101	H4	IAI	HA	6	IncFIA,IncFIB, IncFII, IncI					■			■			■
6	58	B1	31	O9	H17			11	Col156, IncFIA, IncFIB IncFIC, IncI1-I			■		■		■		■		
10			32		H25	UTI	CA	12	IncFIB IncFIC			■						■		
1	95	B2	27	O25	H4			26	IncFIB			■						■		
3	131		30				HA	22	IncFIA, IncFIB IncFII IncI1-I			■		■	■			■		■
8	167	A		O101	H17		CA	13	IncC, IncFIB, IncFIC, IncI1-I, IncY	■										■
4	349	D	54	O86	H2		HA	22	IncFIA,IncFIB, IncFIC, IncI1-I, IncL		■									
5	405		27	O102	H6	IAI		16	IncFIB, IncFII, IncL, IncQ1		■								■	■
9							CA	15	IncFIA, IncHI1B, IncI1-I, IncY					■					■	■
2	1485*	F	58	O83	H42			21	IncFIA,IncFIB, IncFIC, IncHI2, IncHI2A				■						■	

Virulence score is the number of virulence genes that were detected in an isolate. CA, community-acquired; FQ, fluoroquinolone; HA, hospital-acquired; IAI, intraabdominal infection; ST, sequence type; UTI, urinary tract infection.