### Letter to the Editor

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### First Detection of VIM-4 Metallo-β-Lactamase-Producing *Citrobacter freundii* in China

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#### Dear Sir,

Carbapenems are one of the most critically important antimicrobials and are considered as last-choice drugs in clinical settings. However, the emergence and dissemination of Enterobacteriaceae carrying acquired metallo- $\beta$ -lactamases (MBLs) is being increasingly reported throughout the world, presenting a serious threat to the effectiveness of carbapenem treatment [1]. Acguired MBLs, such as VIM, IMP, and NDM, are associated with different mobile elements, and the genes for VIM-type enzymes are carried as mobile gene cassettes inserted in class 1 integrons. In Citrobacter freundii, MBL-mediated resistance is rare and only scattered studies have reported the isolation of a few MBL-producing strains, particularly in the Far East [2]. To our knowledge, there is no report of VIM-producing C. freundii in China. In this study, we describe the first identification of the blavim-4 gene in a clinical C. freundii strain isolated from a patient in a hospital in Wuhu, China.

The multidrug-resistant *C. freundii* CF0638 isolate was recovered from the sputum of a 66-year-old man presenting with diabetic foot to a hospital in Wuhu, China, in September 2011. During the period of hospitalization, symptoms of lower respiratory tract infection emerged and this patient received treatment with cefotaxime and ciprofloxacin for 7 days. The effect of the treatment was not satisfactory. The patient continued to be febrile until he was treated with imipenem-cilastatin (3 g/day). Imipenem-cilastatin treatment was continued for a 2-week course, and the symptoms gradually disappeared during the patient's hospital stay.

Susceptibility testing was performed by an agar dilution method according to the CLSI guidelines (2012) [3]. C. freundii CF0638 exhibited high-level resistance to most β-lactam antibiotics tested. The minimum inhibitory concentrations (MICs) of imipenem and meropenem were 4 mg/L and 1 mg/L, respectively (Table 1). Positive results of the modified Hodge test with meropenem disc and double-disc synergy test between carbapenems and EDTA indicated the production of an MBL. CF0638 was subjected to PCR analysis using the respective primer pairs and amplifying conditions for *bla*<sub>IMP</sub> and *bla*<sub>VIM</sub>, as described previously [4], and followed by DNA sequencing. Nucleotide sequence analysis and homology searches were carried out using the BLAST database (http://www.ncbi.nlm.nih.gov). To determine whether the imipenem resistance was transferable, conjugation experiments were carried out in Luria-Bertani (LB) broth with sodium azide-resistant Escherichia coli J53 as the recipient. Transconjugants were selected on LB agar plates supplemented with sodium azide (100 mg/L) and cefotaxime (4 mg/L).

PCR amplification and nucleotide sequence analysis showed that CF0638 harbored  $bla_{VIM-4}$ . Conjugation experiments showed that imipenem resistance was successfully transferred to the recipient *E. coli* J53. PCR analysis of the transconjugant strain was positive for the  $bla_{VIM-4}$  gene. Susceptibility test results of *C. freundii* CF0638 and its transconjugant are presented in Table 1.

*C. freundii* has been recognized as an opportunistic pathogen that is rarely involved in nosocomial infections. In neonates and

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Strain	MIC (mg/L)												
	PIP	CTX	CAZ	FEP	FOX	ATM	IMP	MEM	CIP	LVX	GAT	GM	AMK
CF0638	512	128	64	8	>256	>256	4	1	>32	32	16	>256	>1,024
Transconjugant of CF0638	256	128	32	4	64	128	2	0.5	2	1	1	128	512
E. coli J53	< 0.5	< 0.5	< 0.5	< 0.25	< 0.5	< 0.5	< 0.25	< 0.25	< 0.06	< 0.125	< 0.125	< 0.25	< 0.25

Abbreviations: MIC, minimum inhibitory concentration; PIP, piperacillin; CTX, cefotaxime; CAZ, ceftazidime; FEP, cefepime; FOX, cefoxitin; ATM, aztreonam; IMP, imipenem; MEM, meropenem; CIP, ciprofloxacin; LVX, levofloxacin; GAT, gatifloxacin; GM, gentamicin; AMK, amikacin.

immunocompromised patients, however, invasive infections such as bacteremia, meningitis, brain abscesses, pneumonia, endocarditis, and intra-abdominal sepsis have also been reported [5]. VIM-4 is a single amino-acid variant of VIM-1  $\beta$ -lactamase. VIM-4 was first described in *Pseudomonas aeruginosa* in Greece [6], but was subsequently identified as the most common integronencoded MBL in Enterobacteriaceae species in several European countries. *C. freundii* isolates are usually susceptible to carbapenems and only a limited number of carbapenem-resistant strains have been identified. Prompt and accurate detection of MBL-producing *C. freundii* strains is necessary to prevent the dissemination of resistance vectors to more virulent pathogens.

# Authors' Disclosures of Potential Conflicts of Interest

No potential conflicts of interest relevant to this article were reported.

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