

## Multiple colonization with highly resistant bacteria: carbapenemase-producing Enterobacteriaceae, carbapenemase-producing *Pseudomonas aeruginosa*, carbapenemase-producing *Acinetobacter baumannii*, and glycopeptide-resistant *Enterococcus faecium*

To the Editors,

The dissemination of carbapenemase-producing bacteria worldwide is an important source of concern because carbapenemase producers are multidrug resistant (Nordmann and Poirel, 2014). National guidelines increasingly recommend a systematic screening of at least carbapenemase-producing Enterobacteriaceae (CPE) and glycopeptide-resistant enterococci (GRE) in patients admitted to hospitals who have been hospitalized abroad during the preceding 12 months (Lepelletier et al., 2011). We have investigated the occurrence of colonization and infection with multiple highly resistant bacteria of more than 4 different genus in 2 patients directly transferred from a foreign country.

In June 2014, a 33-year-old French man (patient A) was admitted for a suicide attempt in a Vietnamese hospital where he was treated during 10 days for pneumonia with piperacillin + tazobactam before his transfer to Necker-Enfants Malades University Hospital in Paris, France. At the day of his hospitalization in France, distal protected pulmonary samples were collected, and imipenem was administered subsequently to a persistent fever. In addition, systematic screening to detect carbapenemase producers and GRE was also performed. Screening of extended spectrum  $\beta$ -lactamase (ESBL) producing Enterobacteriaceae, carbapenemase producers, and GRE was done on selective media (bioMérieux, La Balme-les-Grottes, France) ChromID ESBL, ChromID Carba Smart, and VRE medium, respectively. Carbapenemase production was identified using the Carba NP test for Enterobacteriaceae (Dortet

et al., 2014a) and *Pseudomonas aeruginosa* (Dortet et al., 2012) and CarbAcineto NP test for *Acinetobacter baumannii* (Dortet et al., 2014b). Definitive identifications of resistance determinant were done by PCR amplifications followed by sequencing. Pulmonary samples grew an OXA-23-producing *A. baumannii* isolate and an IMP-1-producing *P. aeruginosa* (Table 1). Screening identified also that the patient was colonized with a KPC-2-producing *Klebsiella pneumoniae*, a CTX-M-15-producing *K. pneumoniae*, and a VanA-positive glycopeptide-resistant *Enterococcus faecium* (Table 1).

Patient B was a 66-year-old French woman who was hospitalized in May 2014 in the intensive care unit of Paris suburb (Saint-Denis, France). She was directly transferred from Morocco, where she was admitted in the intensive care unit consecutively to a road accident. Seven days after her admission, she developed a ventilation-associated pulmonary infection due to a multidrug-resistant *P. aeruginosa* that was treated with imipenem, colistin, metronidazole, and fluconazole for 14 days. Twenty-one days after the admission, the patient developed another pulmonary infection due to a multidrug-resistant *A. baumannii* only susceptible to colistin, resulting in her transfer to the French hospital. At the admission, the systematic screening of CPE using rectal swab samples revealed the presence of an OXA-48-producing *K. pneumoniae* (Table 1). After 24 hours, the patient developed a pyelonephritis due to a VIM-4-producing *P. aeruginosa* and a bronchitis due to an OXA-23-producing *A. baumannii* (Table 1), which were treated with amikacin, fosfomycin, and colistin. Although apyrexia was

**Table 1**  
Clinical data on the 2 patients hospitalized in France after hospitalization in another country and were carrying multidrug-resistant bacteria.

Patient	Country of initial hospitalization	Species	Clinical sample	$\beta$ -Lactamase content <sup>a</sup>	Non- $\beta$ -lactam resistance determinants
A	Vietnam	<i>A. baumannii</i>	PDP	<b>OXA-23</b>	ArmA
		<i>P. aeruginosa</i>	PDP	<b>IMP-1</b>	
		<i>K. pneumoniae</i>	Rectal swab	<b>KPC-2</b> , <u>CTX-M-15</u> , TEM-1, SHV-1, OXA-1	AAC6'-Ib-cr, QnrB
		<i>K. pneumoniae</i>	Rectal swab	<u>CTX-M-15</u> , TEM-1, SHV-1, OXA-1	AAC6'-Ib-cr, QnrB
		<i>E. faecium</i>	Rectal swab	None	VanA
B	Morocco	<i>A. baumannii</i>	PDP	<b>OXA-23</b>	
		<i>P. aeruginosa</i>	Urine and blood culture	<b>VIM-4</b>	
		<i>P. rettgeri</i>	Urine	<b>NDM-1</b> , CTX-M-15, SHV-12, TEM-1,	AAC6'-Ib-cr, QnrB
		<i>K. pneumoniae</i>	Rectal swab	<b>OXA-48</b> , <u>CTX-M-15</u> , OXA-9	AAC6'-Ib-cr, QnrB

PDP = distal protected pulmonary sample.

<sup>a</sup> Carbapenemase are in boldface; ESBL are underlined.

obtained in 3 days, fever reappeared under colistin treatment. A naturally colistin-resistant *Providentia rettgeri* isolate producing the NDM-1 carbapenemase was isolated from 3 successive urine samples recovered by urinary catheter (Table 1). The urinary catheter was removed, and the patient received 2 doses of amikacin, leading to a rapid apyrexia.

Co-occurrence of multiple resistance determinants in the same species has become a common feature among carbapenemase producers (Compain et al., 2014). Nevertheless, co-occurrence in the same patient of a least 4 different highly resistant bacterial species is worrisome because it might potentially lead to a real therapeutic deadlock.

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