

RAPID COMMUNICATION

Detection of *Citrobacter koseri* carrying beta-lactamase KPC-2 in a hospitalised patient, Greece, July 2011

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This report describes the detection of *Citrobacter koseri* carrying *K. pneumoniae* carbapenemase (KPC-2) isolated in July 2011 from a Greek patient, who was also colonised by a *Klebsiella pneumoniae* strain co-producing KPC-2 and Verona integron-encoded metallo-beta-lactamase (VIM)-1.

Case description

On July 2011, a Greek male in his early 80s was admitted to the University Hospital of Larissa, Greece, due to an acute myocardial infarction. He had chronic hypertension due to diabetes and had been hospitalised in the same hospital two months earlier, due to suspicion of tuberculosis, which was later ruled out (sputum and bronchoalveolar lavage cultures were negative for *Mycobacterium tuberculosis*). His respiratory tract symptoms soon remitted and he was discharged.

On admission for the myocardial infarction in July, he was given an empirical prophylactic antibiotic regimen of amoxicillin and clindamycin. On the fourth day of hospitalisation, he became febrile (40 °C) and cultures taken from various samples (blood, vein catheter, urine, sputum and bronchoalveolar lavage) were examined. Gram staining of the bronchoalveolar lavage showed the presence of Gram-negative rods along with leucocyte infiltration, indicating a respiratory tract infection. The presence of the Gram-negative bacterium was confirmed by culture, while the rest of the clinical samples were negative.

Using the VITEK 2 system (bioMérieux, France), the isolate was found to be *Citrobacter koseri* with reduced susceptibility or resistance to beta-lactams, including carbapenems (cefotaxime, ceftazidime, cefepime, aztreonam and ertapenem), but susceptible to fluoroquinolones (ciprofloxacin) and aminoglycosides (amikacin and gentamicin) (Table). Identification of *C. koseri* was further confirmed by PCR (amplifying the 16S ribosomal (r) RNA) and sequencing [1]. Non-beta-lactam antibiotics, such as quinolones and aminoglycosides,

were active against the isolate (Table). The antimicrobial chemotherapy was altered to ciprofloxacin and the fever abated within days. Screening of faecal samples for carbapenem-resistant *Enterobacteriaceae* performed with McConkey agar supplemented with 1mg/L imipenem showed that the patient's gastrointestinal tract was colonised by an imipenem-resistant *Klebsiella pneumoniae* (Table).

Ten days later and while still on treatment with ciprofloxacin, the patient had a second febrile episode and imipenem-non-susceptible *K. pneumoniae* isolates were recovered from three cultures (two from blood and one from a vein catheter). The antibiotic resistance profiles of these three isolates were similar to that of the *K. pneumoniae* isolate from the faecal sample (Table). The patient was then successfully treated with a combination of tigecycline and gentamicin.

Background

The emergence of *Enterobacteriaceae* producing class A beta-lactamases of *K. pneumoniae* carbapenemase (KPC) type is a major clinical and public health concern as these enzymes have the potential to compromise treatment with all beta-lactams, including carbapenems. They are typically transposon-encoded determinants and therefore have the ability to move between plasmids and across bacterial species.

C. koseri, an environmental, Gram-negative bacterium, is occasionally found as a coloniser of the human gastrointestinal tract.* Although the potential virulence of the species is considered low, it is sporadically implicated in serious nosocomial infections [2]. Furthermore, *C. koseri* has the ability to easily incorporate antibiotic-resistance determinants [3]. Although there are several reports of *C. koseri* isolates bearing extended-spectrum beta-lactamase (ESBL) genes, the detection of *C. koseri* isolates producing a carbapenemase is rare [4-6]. To our knowledge, there is only one report of a *C. koseri* isolate producing Verona integron-encoded

metallo-beta-lactamase VIM-(1) [7]. Here, we present the first report of a *C. koseri* clinical isolate producing KPC-2.

Laboratory investigations

All five bacterial strains isolated from the patient (one *C. koseri* and four *K. pneumoniae*) were studied further. The minimum inhibitory concentration of various antibiotics were determined by Etest (bioMérieux, France) and the results were interpreted according to the criteria of the European Committee on Antimicrobial Susceptibility Testing [8]. The modified Hodge test was used for phenotypic detection of carbapenemase production of either KPC or VIM type: all five isolates were positive. Meropenem-boronate and meropenem-ethylenediaminetetraacetic acid (EDTA) combined disk tests, performed as described elsewhere [9], indicated the presence of a KPC in the *C. koseri* isolate and the co-production of KPC and metallo-beta-lactamase (MBL) in the *K. pneumoniae* isolates.

PCR assays were used to screen all five isolates for *bla* genes encoding KPC, VIM, SHV, TEM, OXA-1, CMY and CTX-M [10-12]. Sequencing of the entire amplified genes revealed the carriage of *bla*_{TEM-1} and *bla*_{KPC-2} by *C. koseri* and *bla*_{TEM-1}, *bla*_{KPC-2}, and *bla*_{VIM-1} by *K. pneumoniae*. No nucleotide differences were observed in the *bla*_{TEM-1} and *bla*_{KPC-2} genes (which are known to be highly conserved) in the *C. koseri* and *K. pneumoniae* isolates.

The genetic relatedness of the *K. pneumoniae* isolates was investigated by multilocus sequence typing based on seven housekeeping genes (*rpoB*, *gapA*, *mdh*, *pgi*, *phoE*, *infB* and *tonB*) [13]. All four isolates were found to belong to ST147.

It was not possible to determine whether *C. koseri* could be found in the patient's gastrointestinal tract. Detection of *C. koseri* in faecal samples relies on

amplification of 16S rRNA by PCR; however, this would amplify almost all bacteria that colonise the gastrointestinal tract.

Control measures taken

As recommended by the Greek health authorities, patients are not screened for carbapenem-resistant bacteria on admission or during hospitalisation. Weekly carrier screening is performed only in wards where a carbapenemase-producing bacterium is isolated.

When *C. koseri* was isolated from the patient, faecal samples were taken from the other six patients who were being treated in the same unit. None were found to be colonised by a KPC-positive *Enterobacteriaceae*. The patient was isolated in a single-bed room, under strict infection control measures.

Discussion

Previous studies have presented evidence for horizontal dissemination of *bla*_{KPC} genes among enterobacterial species worldwide [14-17]. However, to our knowledge, this is the first report of *C. koseri* producing KPC-2.

Enterobacteriaceae producing KPC were introduced in Greece in 2007 [18]. Since then, the prevalence of these microorganisms has risen to epidemic proportions, especially in teaching hospitals in the main urban areas [19]. This can be attributed to the overcrowded patient population in these hospitals, serious shortage of specialised personnel and consequently inadequate infection control. In the hospital in Larissa, where the prevalence of KPC-positive *K. pneumoniae* is approximately 27%, similar to that in other teaching hospitals in Greece [20], isolates of various species, such as *Serratia marcescens*, *Escherichia coli* and *Enterobacter aerogenes* carrying *bla*_{KPC} genes have been occasionally isolated from infected or colonised patients.

TABLE

Minimum inhibitory concentrations of various antimicrobial agents against isolated bacteria from a hospitalised patient, Greece, July 2011

Isolate (type of sample)	Etest results for the antimicrobial agents (mg/L and interpretation ^a)											
	IMP	MEM	ERT	CTX	CAZ	FEP	ATM	COL	TIG	CIP	GM	AN
<i>Citrobacter koseri</i> (bronchoalveolar lavage) n=1	0.75 S	0.19 S	8 R	4 R	2 NS	1.5 NS	4 R	0.25 S	0.25 S	0.25 S	0.19 S	2 S
<i>Klebsiella pneumoniae</i> (blood) n=2	32 R	16 R	32 R	64 R	256 R	32 R	64 R	16 R	0.5 S	16 R	1 S	12 NS
<i>K. pneumoniae</i> (vein catheter) n=1	32 R	16 R	32 R	64 R	256 R	32 R	64 R	16 R	0.5 S	16 R	1 S	12 NS
<i>K. pneumoniae</i> (faecal) n=1	32 R	16 R	32 R	64 R	256 R	32 R	64 R	16 R	0.5 S	16 R	1 S	12 NS

AN: amikacin; ATM: aztreonam; CAZ: ceftazidime; CIP: ciprofloxacin; COL: colistin; CTX: cefotaxime; ERT: ertapenem; FEP: cefepime; GM: gentamicin; IMP: imipenem; MEM: meropenem; TIG: tigecycline.

^a R: resistant; NS: non-susceptible; S: sensitive.

The fact that the patient in this report was colonised by a KPC-positive *K. pneumoniae* in his gastrointestinal tract suggests the in vivo transfer of *bla*_{KPC-2} to *C. koseri*. Given that *K. pneumoniae* was not isolated from bronchoalveolar lavage cultures, it is most likely that the transmission could have occurred in the gastrointestinal tract. Bronchoalveolar lavage and sputum cultures during the patient's first hospitalisation did not reveal the presence of *C. koseri* or *K. pneumoniae* or other bacteria. It could be hypothesised that, during his first hospitalisation, he was colonised in his gastrointestinal tract by a *K. pneumoniae* of ST147, which predominates in this setting (unpublished data). A probable scenario is that the *bla*_{KPC-2} gene was transmitted from *K. pneumoniae* to *C. koseri*, since both are part of the gastrointestinal flora. Identification of a KPC-encoding *C. koseri* strain from faecal samples would provide support for this hypothesis. No such strain was isolated, but given the low minimum inhibitory concentration of imipenem against the *C. koseri* isolate, a likely explanation is that the imipenem concentration used for screening of faecal samples (1 mg/L) suppressed the growth of *C. koseri*.

Rapid spread of carbapenemase-producing *Enterobacteriaceae* is a serious concern in clinical patient care in Greece. The finding of an additional enterobacterial species, *C. koseri*, producing KPC underscores the increasing clinical importance of carbapenemase-positive microorganisms in this country.

*Authors' correction:

At the request of the authors, the word 'anaerobic' was deleted from the sentence '*C. koseri*, an environmental, anaerobic, Gram-negative bacterium, is occasionally found as a coloniser of the human gastrointestinal tract'. This change was made on 24 October 2011.

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