

R473

Publication Only

Infection Control: Clinical epidemiology of nosocomial infections

First description of KPC-3 producing *Klebsiella pneumoniae* in an acute care hospital in the north of Portugal

D. Gonçalves¹, P. Cecílio², C. Iglésias³, A. Faustino³, A. Estrada³, H. Ferreira¹

¹Biological Sciences - Microbiology, University of Porto - Faculty of Pharmacy - REQUIMTE, Porto, Portugal ; ²Biological Sciences - Microbiology, University of Porto - Faculty of Pharmacy, Porto, Portugal ; ³Clinical Pathology, Hospital de Braga, Braga, Portugal

Objectives:

Klebsiella pneumoniae and KPC-carbapenemases share a common history. *Klebsiella pneumoniae* isolates producing KPC-carbapenemases were described worldwide, and are today an endemic public health threatening problem, in some countries, with a relatively high hospital infection specific mortality. In Portugal, KPC-3 producing *Klebsiella pneumoniae* has been reported in multiple isolates in the center of the country, but in the North, KPC-3 has never been reported in *Klebsiella pneumoniae*, till now. A KPC-3 harboring *Escherichia coli* isolate has already been reported in our previous work. Here we describe the arrival of KPC-3 producing *Klebsiella pneumoniae* to an acute care hospital in the North of Portugal.

Methods:

Identification and susceptibility tests were performed by Vitek2 automated system. Carbapenem resistance mechanism was investigated by phenotypic methods by E-test IP/IP1. Carbapenemase coding genes (*bla*_{IMP}, *bla*_{VIM}, *bla*_{NDM}, *bla*_{KPC} and *bla*_{OXA-48}) were screened by PCR using specific primers. Amplified products were sequenced and compared to sequences available in the GeneBank database.

Results:

Three multidrug-resistant *Klebsiella pneumoniae* isolates were detected from November 2012 to September 2013, from different biological samples (urine n=1, bronchial aspirate n=1, expectoration n=1). Patient ages varied from 38 to 97 years old. One of the patients was admitted to the intensive care ward and two of them were in medicine. All the isolates showed resistance to at least one carbapenem (ertapenem n=3 MIC \geq 8, meropenem and imipenem n=2 MIC \geq 16). However, antimicrobial susceptibility phenotypes were found to be different. The only associate resistance common to all isolates was to trimethoprim/sulfamethoxazole. One isolate showed a resistance phenotype to aminoglycosides (resistant to gentamicin and tobramycin; intermediate to amikacin), a second one to nitrofurantoin, minocycline and fluoroquinolones (resistant to ciprofloxacin and levofloxacin), and the third one was also resistant to ciprofloxacin, and intermediate to levofloxacin and tobramycin. Metallo-beta-lactamase detection by Etest MIC determination for imipenem and imipenem+EDTA was negative. PCR amplification and sequence revealed the presence of *bla*_{KPC-3} in all isolates.

Conclusion:

As far as we know, this is the first description of KPC-3 producing *Klebsiella pneumoniae* clinical isolates, in the North of Portugal, although several isolates have already been described in the center of the country. Urgent application of strict infection control measures and active antibiotic resistance surveillance should be implemented to avoid further dissemination in the hospital. One of the patients from whom a KPC-3 producing *Klebsiella pneumoniae* was isolated, was transferred from a hospital in the center of Portugal. Importation by inter-hospital dissemination of KPC-3 producing *Klebsiella pneumoniae* is therefore something to consider. Nevertheless we cannot exclude horizontal gene transfer, once we have already described a KPC-3 producing *Escherichia coli* isolated in the same hospital. Clonal relationships and conjugation must be addressed in future work.