

Title: DIVERSITY OF BETA-LACTAMASE GENES IN *ACINETOBACTER* SP. FROM AN ONCOLOGICAL HOSPITAL IN BRAZIL

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ABSTRACT

Acinetobacter calcoaceticus-*Acinetobacter baumannii* (ACB) complex isolates have been detected among the top five gram-negative bacteria isolated from patients in hospital infections. *Acinetobacter baumannii* is the most important species of the ACB complex and it was recognized as the priority pathogen (critical bacteria), according to World Health Organization (WHO), due to antibiotic multidrug-resistant (MDR) phenotype. The aim of this study was to investigate the diversity of beta-lactamase genes in *Acinetobacter* sp. isolates (n=85) from an Oncological Hospital in Brazil. In addition, *A. baumannii* species were also identified. Extended-spectrum beta-lactamases (ESBL) genes (*bla*_{CTX-M} groups 1, 2, 8, 9 e 25, *bla*_{GES}, *bla*_{VEB}, *bla*_{BEL} and *bla*_{PER}) and carbapenemases genes (*bla*_{KPC}, *bla*_{IMP}, *bla*_{VIM}, *bla*_{NDM} and *bla*_{OXA}) were searched by PCR. *bla*_{OXA-51} gene was used to confirm the *A. baumannii* specie. 75/85 (88.2%) of the isolates were identified as *A. baumannii* and the other ones (10/85, 11.76%) were hereafter denominate *Acinetobacter* sp. 56/75 (74.6%) *A. baumannii* also presented *bla*_{OXA-23-like} gene and one of them (1.17%) presented *bla*_{OXA-48-like} as well. Besides, one *Acinetobacter* sp. presented *bla*_{OXA-143-like} gene. In one hand, OXA-23 beta-lactamase is widely disseminated in *A. baumannii*, on the other hand, OXA-48 beta-lactamase is more common in *Enterobacteriaceae*. ESBL genes were also found in the isolates studied, *bla*_{CTX-M} group 2 (n = 2) and *bla*_{CTX-M} group 8 (n = 1) genes. CTX-M beta-lactamases are common in *Enterobacteriaceae* and rarely found in *A. baumannii*. The presence of *bla*_{CTX-M} and *bla*_{OXA-48-like} genes in the isolates studied demonstrates increased horizontal genes transfer, contributing to selection and persistence of MDR isolates, impacting on morbidity and mortality of patients.

Keywords: *bla*_{CTX-M}, *bla*_{OXA-48-like}, resistance genes, hospital pathogens, hospital infection

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