Títle: Evaluation of resistance genes and genetic relationship in enterobacteria associated with meningitis

Authors: ¹Ballaben, A.S., ²Silva, P., ¹Ferreira, J. C., ¹Darini, A.L.C

Institution: ¹ School of Pharmaceutical Sciences of Ribeirão Preto – University of São Paulo (USP), Ribeirão Preto; ² Institute Adolfo Lutz – Ribeirão Preto

Abstract: Few reports describe enterobacteria causing meningitis, however it is still worrying. These microorganisms have been shown resistance to many beta-lactam antibiotics and guinolone, reducing the therapeutic options. This study aimed to investigate the presence of beta-lactamases genes (bla_{CTX-M} groups 1,2,8,9 and 25), bla_{TEM} and bla_{SHV}) and genes that confer resistance to quinolones (qnr A,B,S e acc(6)'-lb-cr) and to evaluate the genetic diversity among enterobacteria from cerebrospinal fluid and blood of patients with suspected meningitis in different hospitals from Ribeirão Preto region, from 2007 to 2014. Were studied 12 enterobacteria including 5 Klebsiella pneumoniae, 3 Enterobacter aerogenes, 3 Enterobacter cloacae and 1 Serratia marcescens. The genes conferring resistance to antibiotics were investigated by PCR and the genetic similarity was established by PFGE. The value used to determine the similarity was \geq 90%. The K. pneumoniae isolates were not considered genetically related. The gene blacTX-M group 2 was found in 2 isolates, bla_{TEM} in 1 and bla_{SHV} in 2. The gene aac(6)'-Ib-cr was also found in three K. pneumoniae isolates. All E. aerogenes isolates showed blaTEM, and in 2 also carried qnrS and aac(6)'-Ib-cr. Two isolates from the same hospital were considered genetically related while the other one from another hospital was not genetically related. All three E. cloacae isolates from the same hospital were not genetically related. One of the isolates harbored blaCTX-M group 1 gene, qnrB and aac(6)'-Ib-cr. The other two isolates carried aac(6)'-Ib-cr and one isolate showed blatem. The S. marcescens isolate showed *bla*_{TEM.} According to the evaluated data *bla*_{CTX-M} group 2 was most prevalent among K. pneumoniae isolates. Other studies have shown that this beta-lactamase gene is the most prevalent amongst Enterobacteriaceae in Brazil. The gene blactx-m group 1 was found in only one E. cloacae isolate.. This group is the most prevalent in Europe. In Brazil, it was recently identified for the first time, however with increasing prevalence. Studies showed that quinolone resistance genes have been detected in Enterobacter species, which corroborates with our data. The genetic diversity among isolates of each species even from the same hospitals, suggest a non-clonal dissemination. Resistant enterobacteria causing meningitis may be associated with complications in the development of the disease.

Key-words: Enterobacteriaceae, meningitis, ESBL, qnr

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