Title: Clonal spread of linezolid-resistant *Staphylococcus hominis* sups. *hominis* with rRNA gene C2190T and G2603T double mutations at hospitals in Brazilian northeast

Authors: Cidral, T. A¹; Carvalho, M.C.; Figueiredo, A.M.S.; Melo, M. C. N¹.

Institutions: ¹ - Universidade Federal do Rio Grande do Norte – Centro de Biociências – Departamento de Microbiologia e Parasitologia – Laboratório de Bacteriologia Médica. - Caixa Postal 1524 - Campus Universitário Lagoa Nova, CEP 59078-970 - Natal/RN – Brazil. - ²Instituto de Microbiologia Professor Paulo de Góes – Universidade Federal do Rio de Janeiro - Centro de Ciências da Saúde - Bloco I Cidade Universitária - Ilha do Fundão. Rio de Janeiro/RJ – Brasil

The first report of resistance to linezolid was for one MRSA isolate with a G2576T mutation in the V region of the 23S rRNA gene. The aim of this study were to determinate the mechanism of linezolid resistance in three Staphylococcus hominis subspecies hominis isolated from different patients from three public hospitals in the city of Natal-RN. The strains were identified using VITEK® 2 and MALDI-TOF. Susceptibility of the isolates to antibiotics was measured by disc diffusion method and Etest®. Extraction of whole genome DNA was performed and all strains were screened by Polymerase Chain Reaction (PCR) for the presence of mecA and cfr genes. The domain V region of 23S rRNA gene was sequenced and then aligned with the sequences from a linezolid-susceptible S. aureus reference strain. Pulsed-field gel electrophoresis (PFGE) analysis was performed with Xhol. The three strains were recovered from blood culture exhibited high levels of resistance to linezolid (MICs from 24 to 64 µg/mL) but a low MIC value for vancomycin (from 0.75 µg/mL to 1.5 µg/mL). In addition to methicillin resistance, presented a multidrug resistance phenotype involving different classes of antibiotic. Sequencing of the V region of 23S rRNA gene revealed the presence of C2190T and G2603T mutations in all strains. The cfr gene was not identified in any isolate. PFGE with Xhol for three Staphylococcus hominis sup. hominis isolates revealed the presence of a single pulsotype, suggesting an interhospital clonal spread. However, further studies are necessary to clarify this interhospital spread.

Key Words: Coagulase Negative staphylococci, Linezolid Resistance, C2190T and G2603T mutations, *Staphylococcus hominis*.

Grant: CNPQ, FAPERN