

## **POSTER PRESENTATION**

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# P249: Serratia marcescens among neonates

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From 2nd International Conference on Prevention and Infection Control (ICPIC 2013) Geneva, Switzerland. 25-28 June 2013

## Introduction

Healthcare-associated infections (HAIs) remain a permanent challenge among neonates. *Serratia marcescens* (SM) is a rare causative agent of HAIs, responsible for not more than 2% of HAI cases.

## **Objectives**

To analyze a colonization of SM in a neonatal intensive care unit (NICU).

#### Methods

We carried out an observational study among 489 neonates in a NICU, which included microbiological and epidemiological investigations during the period from March till October, 2012. We performed pulsed field gel electrophoresis (PFGE), random amplification of polymorphic DNA, and we also conducted a genotyping to reveal high pathogenicity island (HPI) and genes encoding haemolysin A (ShlA), haemolysin B (ShlB), and phospholipase A (PLA). In addition, we performed antimicrobial susceptibility testing and scanning electron microscopy (SEM).

## **Results**

We found that 72 neonates were colonized by SM without any cases of acute infection. PFGE revealed that only one SM strain was responsible for this colonization. SEM did not reveal any features of the pathogen, and inoculated SM strain was susceptible to imipenem and cefoperazone sulbactam only. Out of 41 cases investigated, 12 possessed ShlA gene, and no HPI, ShlB and PLA genes were determined. All bacteria belonged to RAPD type 1. Nebulizer was recognized as a source of SM.

# Conclusion

SM was revealed during the pre-epidemic situation and was successfully eliminated after additional chemical

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disinfection of nebulizers. Our data shed light on transmission of SM-caused HAIs.

### Disclosure of interest

None declared.

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Published: 20 June 2013

doi:10.1186/2047-2994-2-S1-P249

Cite this article as: Brusina et al.: P249: Serratia marcescens among neonates. Antimicrobial Resistance and Infection Control 2013 2(Suppl 1): P249.

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