



Outbreak of neonatal infection by an endemic clone of *Serratia marcescens*

Surto de infecção neonatal a partir de clone endêmico de *Serratia marcescens*

Karla Valéria Batista Lima¹, Raimundo Gladson Corrêa Carvalho², Irna Carla do Rosário Souza Carneiro³, Josiane Lillian de Sousa Lima³, Cintya de Oliveira Sousa¹, Edvaldo Carlos Brito Loureiro¹, Lena Lillian Canto de Sá⁴ and Flávia Corrêa Bastos¹

ABSTRACT

Introduction: The outbreak occurred between February and June 2006 and included identification of the cases, analysis of medical records, cultures from environmental sources, resistance analyses and genotyping profile of *Serratia marcescens*. **Methods:** The cultures were composed of 13 blood isolates, 17 rectal and hand swabs and air sampling. **Results:** The data obtained by pulsed-field gel electrophoresis exhibited three strains that contaminated 24 patients. Systemic infection was the most common in neonates with lower weight, long periods of hospitalization, premature delivery and the use of mechanical ventilation. **Conclusions:** This investigation revealed the multifactorial nature of the outbreak. An endemic clone of *S. marcescens* was detected.

Keywords: *Serratia marcescens*. Outbreak. Nosocomial infection.

RESUMO

Introdução: O surto ocorreu entre fevereiro a junho de 2006 e incluiu identificação de casos, análise dos prontuários, culturas ambientais, análise de resistência e genotipagem dos isolados de *Serratia marcescens*. **Métodos:** Os cultivos foram compostos de 13 isolados de sangue e 17 swabs de reto e mãos e amostras do ar. **Resultados:** Os dados obtidos por eletroforese de campo pulsado evidenciaram três cepas que contaminaram 24 pacientes. Infecção sistêmica foi mais comum em neonatos com menor peso, longo tempo de internação, nascimento prematuro e uso de respiração mecânica. **Conclusões:** Foi evidenciada a natureza multifatorial do surto. Foi encontrado um clone endêmico de *S. marcescens*.

Palavras-chaves: *Serratia marcescens*. Surto. Infecção hospitalar.

Intensive care units are often involved in epidemics of *Serratia marcescens* colonization and infection. Important reservoirs in epidemics are the digestive, respiratory and urinary tracts and perineum of neonates and the artificial nails of adults and healthcare workers (HCWs)¹. Medical equipment, lotions, antiseptics, medications and sinks have also been described as sources of such epidemics^{2,3}. Transient transportation of *S. marcescens* on the hands of medical personnel is considered an important mode of transmission in epidemics².

The associations for neonatal infection by *S. marcescens* include low birth weight, long-term hospitalization, mechanical ventilation, invasive procedures and exposure to antibiotics and steroids².

1. Bacteriology and Mycology Section, Evandro Chagas Institute, Ananindeua, PA, Brazil. 2. Cytology Department, State University of Para, Belém, PA, Brazil. 3. Health Department, State University of Para, Belém, PA. 4. Environment Section, Evandro Chagas Institute, Ananindeua, PA, Brazil.

Address to: Dra. Karla Valéria Batista Lima. Instituto Evandro Chagas/ SABMI. Rodovia BR 316 Km 7, s/n, 67030-000 Ananindeua, PA.

Phone: 55 91 3214-2116

e-mail: karlalima@iec.pa.gov.br

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The occurrence of infections by *S. marcescens* in the neonatal intensive care unit (NICU) of reference in Belém motivated the development of this work, in order to establish the genotypic profile of this pathogen and contribute to the control of healthcare-associated infection (HCAI).

Characterization of the institution and of the samples

The study was developed in a neonatal unit (NU) of high complexity, located in the City of Belém, State of Pará. The unit is composed of an internal nursery (IN), for children born in the hospital, a neonatal intensive care unit (NICU) and an external nursery (EN). The IN is divided into five wards: special care (SC), intermediate care (IC), semi-intensive (SI), other disorders and the transition room.

Microbiological methods

From February to June 2006, 1,445 hemocultures were performed, as well as several other cultures: 375 rectal swabs from newborns, 127 from the hands of HCWs, 21 of water from the humidifier, 11 from drugs, alcohol and detergents, 10 involving air sampling in the environment, 5 from the humidifier filter, 5 of the water tap and 5 from the doorknob of the neonatal unit door. The hemocultures were performed in an automated system BACTEC 9,120 (Becton Dickinson) and the others were seeded on sheep blood agar (Difco), Cled agar (Difco) and Mac-Conckey agar (Difco) and incubated in a bacteriological incubator (Fanem) at 37°C for 24h.

Air was collected using AMS100 equipment (Merck), calibrated to collect 250L of air in 2 min 50 sec, which contained a 15/90mm-Petri dish with Cled agar (Difco).

The colonies originating from Gram negative bacilli, negative oxidase, were suspended in saline solution and incubated in a GNI+ card for bacterial identification and in a GNS 655 card (BioMerieux) for susceptibility assays, in accordance with the manufacturer's recommendations.

For evaluation of the sensibility to ertapenem, the methodology proposed by Clinical and Laboratory Standards Institute (CLSI) was used⁴.

Pulsed-field gel electrophoresis

Chromosomal DNA was extracted from stored *S. marcescens* isolates. DNA was digested with *XbaI* and PFGE was performed in electrophoresis system CHEF-DR III, Bio-Rad⁵.

Pulsed-field gel electrophoresis profiles were analyzed using BioNumerics software (*Applied Maths, Sint-Martens-Latem, Belgium*) and the similarity of fragment length patterns between two strains was scored by the Dice coefficient (4% of tolerance).

Case: control study

To determine associations with *S. marcescens* infection or colonization, a case-control study was conducted comparing case-patients (Table 1) with other neonates, who had been admitted to the NICU between February and June 2006. The medical records were reviewed to identify sex, birth weight, gestational age, delivery, length of hospitalization and reason for hospitalization. Based on these studies, each case was matched to a selected control. A case was defined as being a neonate with infection or colonization by *S. marcescens*. The controls were neonates who were hospitalized in the NICU during the same period but did not get infected or colonized.

Statistical analysis

Odds ratios, 95% confidence intervals and two-sided *p* values were calculated using the BioEstat software, version 5.0. The level of statistical significance adopted was $p < 0.05$.

Ethical considerations

The development of this work was approved by the Research Ethics Committee of the Evandro Chagas Institute under protocol CEP/ IEC no. 003/07, of 14 June 2007.

Results and discussion

The total number of positive cultures was composed of 13 isolates from hemocultures, 13 from rectal swabs, 2 from hand swabs, 2 from air sampling, 2 from blood collected some months after the outbreak and 3 from other NUs (PFGE profile 01, 15, 16). Cultures of water from the humidifier, from drugs, alcohol and detergents, the humidifier filter, water tap and doorknob of the neonatal unit door did not detect *S. marcescens*.

The grouping of the isolates identified by macro restriction fragment patterns resolved by PFGE exhibited 11 to 16 fragments ranging from 48.5 to 533.5kb. Typing of the outbreak-related isolates demonstrated that three epidemic strains with distinct fingerprints were associated with cross-infection (Figure 1). Eight patterns contained one isolate and the remaining 27 isolates were grouped into three clusters, containing 3 to 14 isolates. The first isolate of *S. marcescens* in group 1 (samples 4-13) was obtained on February 10th from EN14. Additional occurrences were observed in other wards (NICU and IC), 8 of them isolated from the NICU and two isolated in the EN and IC (Figure 1).

Group 2 was the largest; the first event was observed on March 7th from SC14, followed by dispersions to other wards. Samples 18 and 19 were isolated in the NU from blood of neonates attended some months after the outbreak. The samples of group 3 were dispersed over the months of February, March and June. The antibiotic susceptibility profiles were different for all three clinical isolates. They showed 82.78% similarity with group 2.

The case-control study showed that neonates with *S. marcescens* were more likely to have a lower birth weight, long-term hospitalization, premature delivery and exposure to continuous positive airway pressure (Table 1).

All the isolates of the *S. marcescens* outbreak were resistant to ampicillin and the association of ampicillin and sulbactam and 74.3% of the evaluated samples presented resistance to the β -lactams piperacillin associated with the inhibitor of β -lactamases tazobactam. The index of resistance to the aminoglycosides gentamicin and amikacin was also high in this work (Table 1). Both were susceptible to ertapenem.

TABLE 1 - Distribution of 26 patients according to clinical, demographic variables and the occurrence of neonatal infection.

	Cases		Controls		Odds ratio
	n	%	n	%	p
Gender					
male	14	53.9	15	57.7	0.8556
female	12	46.2	11	42.3	1.0000
Birth weight					
<2,500	24	92.3	13	50.0	12.0000
\geq 2,500	2	7.7	13	50.0	0.0020
Delivery					
cesarean	9	34.6	9	34.6	
vaginal	17	65.4	17	65.4	
Length of hospitalization (days)					
>11	3	11.5	17	65.4	0.0756
\geq 11	21	80.8	9	34.6	0.0004
Maternal age (years)					
<19	4	15.4	7	26.9	0.8352
>19	13	50.0	19	73.1	0.9139
Premature delivery					
yes	19	73.1	11	42.3	5.1818
no	5	19.2	15	57.7	0.0178
Respiratory distress syndrome					
yes	8	30.8	2	7.7	5.3333
no	18	69.2	24	92.3	0.0785
Surgical procedures					
yes	3	11.5	7	24.1	0.3540
no	23	89.7	19	75.9	0.2912
Orogastric tube					
yes	20	76.9	15	57.7	2.9333
no	5	19.2	11	42.3	0.1572
Continuous positive airway pressure					
yes	24	92.3	15	57.7	8.8000
no	2	7.7	11	42.3	0.0104

Internal nursery is recognized as one of the most significant causes of morbidity and mortality among hospitalized newborns, especially in NICUs⁶. However, the exact impact of this condition is difficult to determine, since a wide variation in infection rates is reported in the literature, possibly due to differences in surveillance or study methods. It is important to control the inherent aspects of each NICU and to make the data available to the local health authorities, healthcare workers and the scientific community interested in epidemiological data. Unfortunately, this practice is still not universal and there are few published studies that portray the epidemiology and risk factors for NI in Brazilian NICUs.

The overall NI incidence rates reported in Brazilian NICUs of 30.6%⁷ and 22%⁸ are much higher than those observed in most studies in the United States (11.4%)⁹ or Europe (2.5%)¹⁰.

The results obtained suggest cross-transmission between patients during the period of study. The colonized or infected neonates were the principal reservoir of *S. marcescens* in these outbreaks, as previously reported¹. The initial source and the mode of introduction to the NICUs remain unclear. For group 1, it is possible that the index case was the neonate admitted on February 10th and that later, other neonates in others wards were infected. Sample 10, which was isolated from the hands of a HCW, may have spread the microorganisms to other neonates in the NU.

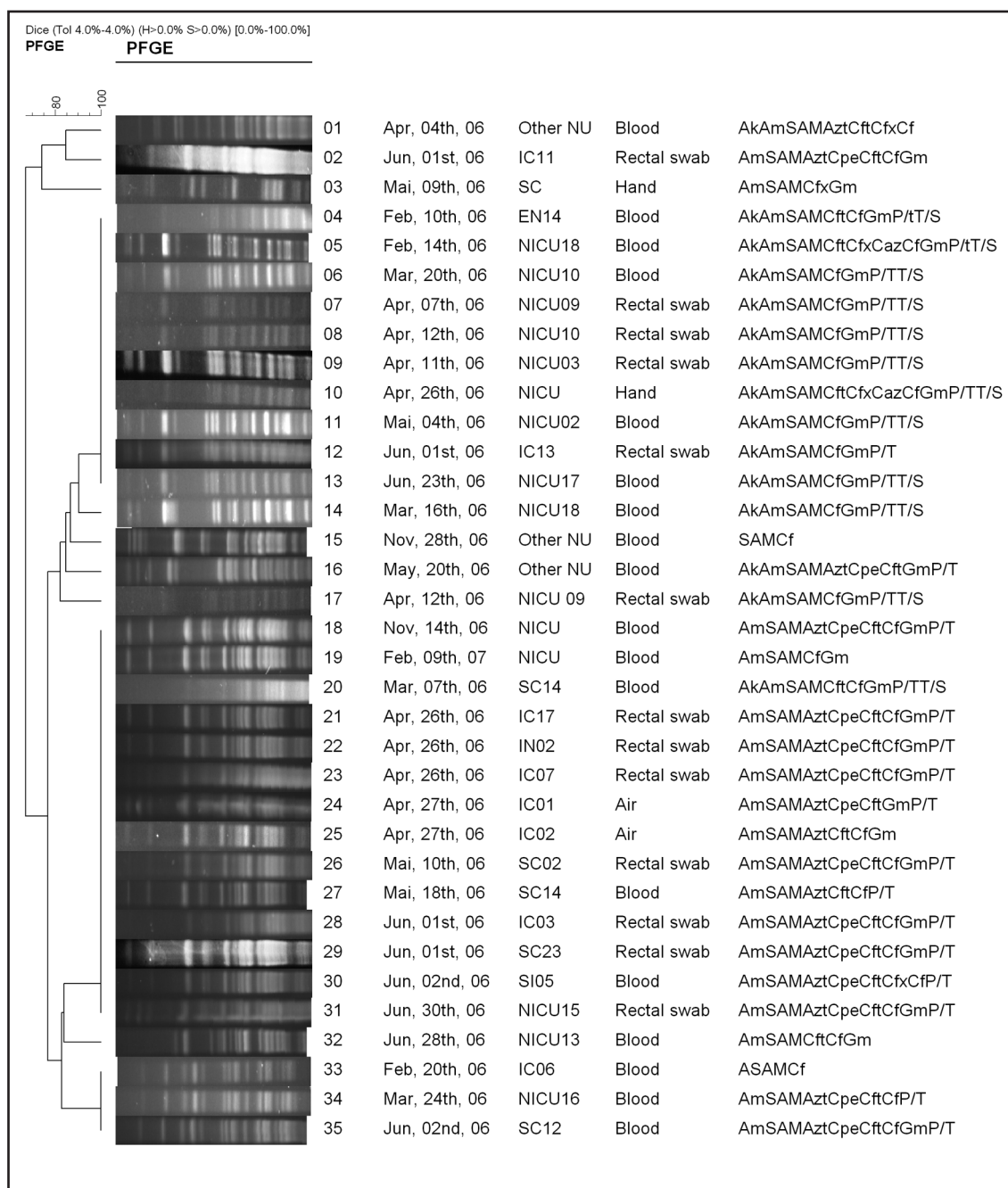


FIGURE 1 - Correlation of pulsed-field gel electrophoresis profile, isolation date, localization in the neonatal unit, site of isolation and resistance profile.

Ak: amikacin, Am: ampicillin, SAM: ampicillin-Subactan, Azt: aztreonam, Cpe: cefepime, Cft: cefotaxime, Cfx: cefoxitin, Caz: ceftazidime, Cf: cephalothin, Gm: gentamicin, P/Z: piperacillin-tazobactam, T/S: trimethoprim-sulfamethoxazole, EN: external nursery, IC: intermediate care, IN: internal nursery, NICU: neonatal intensive care unit, NU: neonatal unit, SC: special care, SI: semi-intensive.

The *S. marcescens*-positive HCW was known by colleagues to strictly adhere to hand hygiene procedures. Observational studies concerning hand hygiene during the epidemic were not performed; however, inspection of the hands of the HCW excluded the presence of rings, artificial nails and psoriasis. Rings, artificial fingernails and psoriasis can harbor gram-negative rods¹¹. Group 2 presented the more prevalent clone that affected ten neonates in 2006. The genotyping data also revealed that the *S. marcescens* strain (genotype 18 and 19) was identified in the same NICU some months after the outbreak, suggesting an endemic situation. The same clone was isolated from the air of the NICU. Subsequent isolates with similar patterns suggest that prolonged, persistent person-to-person

transmission of strains occurred. During the outbreak, patients were transferred from the NICU to surgery units and readmitted to the NICU in cases of patient deterioration. Another problem is related to the HCWs assisting in different wards in the NU. The actions outlined above facilitated the dissemination of the outbreak strains between different wards.

Low birth weight, longer periods of hospitalization, premature delivery and exposure to continuous positive airway pressure were risk factors for NICU acquired infections. In this study the majority of patients had low birth weight. Brito et al also reported that a large proportion (92.3%) of cases presented low birth weight and prematurity¹².

Waterless, alcohol-based hand antiseptic has been shown to be more effective than antimicrobial soap in eliminating pathogens from the hands. McNeil et al demonstrated that only 11% of HCWs with artificial nails who had pathogens at baseline effectively removed the pathogens using soap compared to 38% using alcohol-based gel¹³. Of those HCWs who had natural nails and pathogens at baseline, 14% removed the pathogens using soap compared to 80% of those using an alcohol-based antiseptic ($p = 0.09$). The introduction of alcohol-based hand antiseptic to the NICU effectively provided easy access to another method of hand hygiene, which may have contributed to more frequent hand washing between contacts with patient.

After the outbreak, several infection control measures were introduced at the same time, including the alcohol-based hand antiseptic; therefore, it is difficult to identify which measures were responsible for controlling the infection. Some months after the outbreak, two new NI (18 and 19 samples, **Figure 1**) were observed, but they were controlled. Future outbreak control measures should concentrate on the intensity of colonization, not only of the neonates, but also of the mothers and HWCs^{1,14,15}. The antibiotic use and prophylaxis in mothers should be evaluated on a case-by-case basis, thus limiting the potential for broad-spectrum antibiotics to select multiresistant Gram-negative bacteria.

The association tazobactam with piperacilin has been used as an alternative for treatment of HCAI caused by Gram-negative bacilli and anaerobes producers of β -lactamases. In the NICU studied, tazobactam with piperacilin should not have been used for *S. marcescens* infection.

Although genotyping has no influence on the therapy, clinicians should forward isolates for epidemiological research because genotyping can help trace infection sources and is useful in infection control and epidemiological surveys of hospital infections, particularly on the neonatal ward. The lack of cases in 2008, following the implementation of infection control measures to prevent spread, serves as additional evidence that this epidemic of 26 *S. marcescens* cases was indeed due to transmission.

Genotyping was available and showed identical strains in the NICU in the clusters, confirming that cross-infection occurred. However, *S. marcescens* most likely acted only as an indicator. It is possible that other pathogens were being horizontally transmitted at the same time, but they were not investigated and thus were not detected.

This investigation showed the multifactorial nature of the development of clusters caused by *S. marcescens*. These outbreaks seldom have only one cause or source, but are more likely due to several factors. Attention should be paid to healthcare workers, the mothers and the neonates, in addition to general risk factors and infection control policies, to ensure an adequate healthcare worker-to-patient ratio.

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CONFLICT OF INTEREST

The authors declare that there are no conflicts of interest.

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