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Colonization profile and duration by multi-resistant organisms in a prospective cohort of newborns after hospital discharge

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ABSTRACT

The aim of this study was to determine the spontaneous decolonization period and characteristics in a prospective cohort of newborns colonized by multidrug-resistant organisms, after their discharge from the neonatal intensive care unit. Multidrug resistance is defined as bacterial non-susceptibility to ≥ 1 agent of ≥ 3 antimicrobial categories. In total, 618 newborns were included in the study, of which 173 (28.0%) presented a positive culture for multidrugresistant microorganisms, and of these, 52 (30.1%) were followed up in this study. The most frequent intrinsic factors were be born by cesarean section (86.5%), prematurity (84.6%), and very low birth weight (76.9%). The extrinsic factors were having remained hospitalized for an average of 27 days, during which 67.3% were submitted to invasive procedures and 88.5% received antimicrobials. The intrinsic and extrinsic factors of newborns were not associated to a decolonization period longer or shorter than 3 months, which was the average period of decolonization found in the present study. From the totality of colonization cultures sampled at hospital discharge, the Gram-negative Extended Spectrum β-lactamase producing bacteria were the most common, with 28.9% of babies colonized by Klebsiella spp. The median period of decolonization by multidrug-resistant microorganisms in the newborns population after hospital discharge was 3 months, but was highly dependent on the microbial species, and this period was not associated to any intrinsic and extrinsic factors of the newborn.

KEYWORDS: Premature. Neonatal intensive care. Antimicrobial resistance. Colonization. Decolonization. Very low birth weight.

INTRODUCTION

The human microbiota composition begins in the maternal uterus¹⁻³ and continues with the acquisition of microorganisms in the birth canal and later from maternal skin⁴. This natural process is modified in premature newborns who are usually delivered by cesarean sections⁵, and for whom the contact with the mothers is replaced by care in a neonatal intensive care unit (NICU), in which the babies are more exposed to microorganisms belonging to the hospital setting^{6,7}, as well as those of the health professionals⁸. Among the microorganisms in the hospital setting, antimicrobial resistant ones result from the selection pressure triggered by these drugs, widely used in intensive care units⁹, as well as by cross contamination¹⁰.

Multidrug-resistant (MDR) bacteria are considered a major threat to public health by the World Health Organization (WHO) due to the therapeutic limitations and high lethality of infections caused by these agents^{11,12}, especially in the neonatal population, whose immunological immaturity and fragility of skin barriers favor these infections⁸.

Colonization by MDR increases the risk of infection by these microorganisms¹³. Given this evidence, one can infer that this risk can extend to the domiciliary context when the newborn remains colonized by MDR after discharge. Colonization also poses a risk of MDR transmission to family members¹⁴ which remains until the moment of decolonization.

The process of decolonization has been studied in adults¹⁵⁻¹⁸ and is related to the topical use of antiseptics in aqueous formulations, and of antimicrobials in ointments. However, due to the immaturity of the skin barrier, these strategies are not applied to newborns due to the toxicity and systemic absorption potential of these antiseptic antimicrobial agents in this population¹⁹.

Considering the negative impact of MDR colonization and the knowledge gap regarding the period in which the newborn remains colonized after hospital discharge, the present study aimed to determine the period for spontaneous MDR decolonization in newborns after discharge from a neonatal intensive care unit. In addition, associations of the decolonization period with perinatal and clinical variables during hospitalization were investigated.

MATERIALS AND METHODS

Study design and setting

This is a prospective cohort study of newborns colonized by MDR during hospitalization in a neonatal intensive care unit of a university hospital. They were followed up after discharge until decolonization in a 4-year convenience period, from January 2014 to September 2018.

Inclusion and exclusion criteria

The study population consisted of 1,395newborns, of which 618 met the inclusion criteria as they remained hospitalized in the same unit for a period of more than 72 h and their mothers agreed to participate in the study. Of these, 28.0% (n=173) presented colonization by MDR as shown by positive cultures and were prospectively monitored. Every 30 days, swab samples collected during domiciliary visits or during an ambulatory consultation were cultured until a negative culture for MDR was obtained.

Babies who were readmitted to hospital or died during the period of domiciliary monitoring or outpatient followup were excluded. Losses were considered when swab collection at hospital discharge was not performed; the family moved to a municipality other than the study site; the mother withdrew consent to participate in the study and the contact with mothers was lost due to telephone number and address changes (Figure 1).

The study was carried out in a reference center for highrisk pregnancies and neonatal intensive care. The neonatal unit is composed of a Neonatal Intermediate Care Unit with 20 beds, and a Neonatal Intensive Care Unit (NICU) with ten beds.

Data collection

The clinical and epidemiological data regarding hospitalization were collected from the medical records of the newborns and analyzing the following variables: sex, type of delivery, birth weight, gestational age, invasive procedures, antimicrobial use, type of breastfeeding and period of hospitalization.

At hospital discharge, swabs were collected for culture (Transystem Stuart collection device, COPAN Diagnostic, Italy); the first set of swabs was collected from the oral, nasal, axillary and inguinal regions, simultaneously; the second set of swabs was collected from the inguinal and rectal regions, simultaneously. Then, swabs were sent to the microbiology laboratory and processed within 72 h.

Microbiology analysis

The first set of swabs (1) was inoculated into Tryptic Soy Broth (TSB, Oxoid, Cheshire, England) supplemented with 16 μg/mL cefotaxime (TSB-CTX) to select Gram-negative bacilli and TSB supplemented with 6.5% NaCl (TSB-NaCl) to select *Staphylococcus aureus*. The second set of swab (2) was inoculated in TSB-CTX and TSB supplemented with 6 μg/mL vancomycin (TSB-VRE) to select *Enterococcus* spp. The cultures were incubated at 35 °C for 24 h. Cultures that did not show growth were maintained for another 24 h. After growth, the bacteria were seeded in Petri dishes containing VRE Agar, Mannitol Salt Agar (MSA) and MacConkey Agar supplemented with 16 μg/L cefotaxime (CTX-MacConkey), that were incubated at 35 °C for 24 h. Plates that did not show growth in 48 h were discarded and the culture was considered negative.

The bacterial susceptibility profile to the antimicrobial agents was analyzed by the disc-diffusion technique, as recommended by the Centers for Disease Control and Prevention²⁰, considering the Clinical and Laboratory

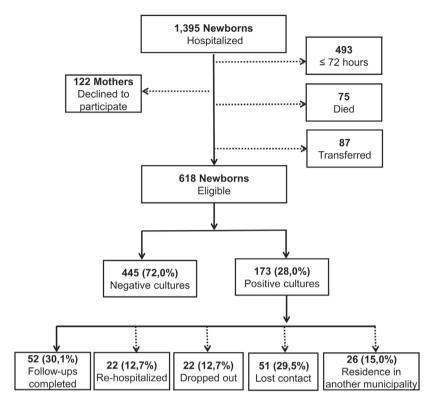


Figure 1 - Flowchart of the study sample design, Parana State, Brazil, 2014-2018.

Standard Institute²¹⁻²⁵ and European Committee on Antimicrobial Susceptibility Testing²⁶, to detect extended spectrum β -lactamase producing microorganisms (ESBL), performed using the double disc synergism test.

Molecular techniques were not performed as this study did not include the identification of clonal transmission of MDR as an objective²⁰.

MDR is defined as a bacterial non-susceptibility to ≥ 1 agent in ≥ 3 antimicrobial categories considering: Enterobacteria, *Acinetobacter baumannii* and *Pseudomonas aeruginosa* resistant to 3^{rd} or 4^{th} generation cephalosporins or monobactams (Extended Spectrum β -lactamase – ESBL) and carbapenems (Carbapenem resistance – CR); *Staphylococcus aureus* resistant to methicillin/oxacillin (MRSA) and *Enterococcus* spp. resistant to vancomycin (VRE)²⁷. The *Enterobacter* spp. and *Klebsiella* spp. were grouped for consistency in the frequency analysis. After the antimicrobial susceptibility test, the bacteria were stored in 30% TSB-glycerol at -20 °C.

The newborns who did not receive antimicrobial therapy for several weeks and who had two repeatedly negative cultures for MDR over two weeks were considered decolonized²⁰.

Statistical analysis

The data were tabulated in Microsoft Office Excel

97-2013® and analyzed in the Statistical Package for the Social Sciences (SPSS) version 20.0 (IBM, Armonk, New York, USA). The results of the continuous variables are described as mean and standard deviation (SD) or median and interquartile range (ITQ), depending on the distribution of the data. Categorical variables were analyzed using the Pearson *Chi-square* test. In order to investigate the magnitude of the association between variables, a binary logistic regression was performed to obtain the *odds ratio* (OR) and 95% confidence interval (95%CI).

Ethical considerations

This research was approved by the Ethics Committee on Research involving Humans (CAAE No. 15415413.4.0000.5231) of the Universidade Estadual de Londrina.

RESULTS

Among the newborns followed until decolonization (n=52), most of them were male (53.8%) and born by cesarean section (86.5%). Regarding the gestational age, 84.6% of the infants were premature, 34.6% were born before 30 weeks, 44.2% were between 31 and 34 weeks, and 21.2% were born after 34 weeks. In total, 76.9% of the newborns were born with low weight, while 23.1% were

within the normal range (> 2,500 g). At the time of hospital discharge, 48.1% of the infants were exclusively breastfed.

The hospitalization period presented a median of 27 days (8 \pm 124) and during this period, 88.5% of the infants received antimicrobial therapy, with a median of 28 days of use (16 \pm 40). All the infants treated with antimicrobials received β -lactam class antibiotics, with a median of 10 days (1 \pm 36), the most frequent of which were penicillin and oxacillin.

In relation to newborns who had an invasive procedure (n=35), 65.4% required central venous catheters (CVC), 46.2% mechanical ventilation (MPV) and 71.4% underwent more than one procedure.

The perinatal and clinical variables of the newborns during hospitalization were not associated to decolonization periods of more or less than 3 months, however, decolonization \geq 3 months was more frequent among infants born before 35 weeks (91.4%), with low birth weight (< 2000g, 87.0%), and hospitalized for more than 10 days (Table 1).

Gram-negative ESBL-producing bacteria were the most frequently isolated MDR in the colonization cultures of babies at discharge (Figure 2).

The MDR decolonization period after hospital discharge ranged from 1 to 9 months, with a median of 3 months (Figure 3).

The mean period of MDR decolonization corresponded to 5.2 months for *Escherichia coli* ESBL-producer, 4 months for ESBL-producing *Serratia marcescens*, 3 months for *Klebsiella* spp. ESBL/CR, ESBL-producing *Pseudomonas aeruginosa*, MRSA and *Serratia marcescens* CR, 2 months for *Acinetobacter baumannii* CR, 1.6 months for ESBL-producing *Enterobacter* spp., 1.5 months for ESBL-producing *Citrobacter freundii*, and 1 month for *Enterobacter* spp. CR.

Regarding the maximum period of decolonization related to bacterial species, carbapenem-resistant bacteria colonized infants up to 2 months after discharge, and ESBL-producing *E. coli* and ESBL-producing *Klebsiella* spp. continued to colonize infants up to 8 months in the domiciliary context (Figure 4).

DISCUSSION

Little is known about the period of MDR decolonization after discharge, both in infant and adult populations. The scarcity of studies can be justified by the methodological complexity of a follow-up of newborns after hospital discharge, as many factors hamper studies of this nature, such as the operational costs, deaths, rehospitalizations, follow-up in the domiciliary environment, data collection

and biological samples collection. These difficulties were corroborated in the present study, where the follow-up of 70% of the newborns colonized at discharge was interrupted. Even with these losses and exclusions, our study had an important neonatal population of 52 newborns followed-up to decolonization in a study lasting 4 years.

Among the results reported in the literature that address the theme of our study, two analyzed MDR colonization in feces collected from the children. Furthermore, these studies were limited to the monitoring of decolonization by ESBL-producing bacteria, excluding those resistant to carbapenems and MRSA^{14,28}, which were analyzed in the present study.

One of the studies cited above was conducted in Austria in a prospective study that followed-up 25 infants colonized by ESBL producing bacteria after discharge from an NICU. In that study Klebsiella spp. was the most prevalent and remained for up to 1 year, followed by Serratia marcescens, with colonization remaining until the 4th month after discharge. E. coli was isolated from a single newborn who remained colonized for 1 month²⁸. Our study has also shown that ESBL producing *Klebsiella* spp. was the most frequent MDR and continued colonizing the babies for an average of 3 months and a maximum period of 8 months. Although ESBL producing S. marcescens was less frequent in our study, the decolonization period was close to the aforementioned study. Finally, ESBL producing E. coli was more frequent and remained colonizing infants for longer periods after discharge when compared to the results of these authors.

Another research detected ESBL-producing *Klebsiella pneumoniae* in the feces of 51 infants who were domiciliary followed-up after hospital discharge from an NICU. These authors showed that the period of decolonization by this bacterium was on average 12.5 months after hospital discharge, with the maximum period for decolonization being 23.5 months. They also found that antimicrobial treatment and birth by cesarean section are risk factors for longer periods of colonization by ESBL-producing *K. pneumoniae* ¹⁴. In this study, the perinatal and clinical variables of infants were not related to the decolonization period of 3 months or less for any of the microorganisms studied. These disagreements can be explained by differences in sampling methods for culture.

Adult decolonization has been more frequently studied using different sampling sites¹⁵⁻¹⁸. A multicenter study performed in 13 European hospitals using nasal and perineal swab collection revealed that the period for MDR decolonization after discharge from the NICU was 4.8 months, and the study has also included patients

Table 1 - Association of perinatal and clinical variables of newborns (*n*=52) during hospitalization with the decolonization period after discharge from a neonatal unit, Parana State, Brazil, 2014-2018.

Neonatal Variables	Newborr	Newborns (N=52)				
	Decolonization ≥ 3 months 23 (44.2) n (%)	Decolonization < 3 months 29 (55.8) n (%)	OR	CI 95%		p-value
Gender						
Female	10 (43.5)	14 (48.3)	0.82	0.27	2.48	0.730
Male	13 (56.5)	15 (51.7)	1.00			
Type of Birth	,	,				
Normal	3 (13.0)	4 (13.8)	1.00			
Cesarean section	20 (87.0)	25 (86.2)	1.07	0.21	5.33	0.937
Gestational Age						
< 30 weeks	10 (43.6)	8 (27.6)	8.75	0.88	86.60	0.064
31-34 weeks	11 (47.8)	12 (41.4)	6.42	0.68	60.84	0.105
35-36 weeks	1 (4.3)	2 (6.9)	3.50	0.15	84.69	0.441
≥ 37 weeks	1 (4.3)	7 (24.1)	1.00			
Birth Weight	. ,	, ,				
< 1,000 g	3 (13.0)	6 (20.7)	1.50	0.22	10.08	0.677
1,000-1,499 g	7 (30.4)	4 (13.8)	5.25	0.87	31.55	0.070
1,500-1,999 g	10 (43.6)	7 (24.1)	4.29	0.84	21.76	0.079
2,000-2,499 g	Ô	3 (10.3)	NA			
>2,500g	3 (13.0)	9 (31.1)	1.00			
Discharge Weight	` '	, ,				
<1,000 g	0	0	NA			
1,000-1,499 g	0	1 (3.4)	NA			
1,500-1,999 g	14 (60.9)	12 (41.4)	3.21	0.81	12.75	0.098
2,000-2,499 g	5 (21.7)	5 (17.2)	2.75	0.51	14.86	0.240
> 2,500 g	4 (17.4)	11 (37.9)	1.00			
Breastfeeding on Discharge	,	,				
Exclusive	12 (42.9)	13 (54.2)	1.00			
Not exclusive	11 (57.1)	16 (45.8)	0.75	0.25	2.23	0.599
Hospitalization Period	,	,				
≤ 10 days	2 (8.7)	6 (20.7)	1.00			
> 10 days	21 (91.3)	23 (79.3)	2.74	0.49	15.09	0.247
Invasive Procedures	,	,				
Yes	14 (60.9)	21 (72.4)	0.59	0.18	1.91	0.380
No	9 (39.1)	8 (27.6)	1.00			
Number of Procedures		· (=:::)				
None	9 (39.1)	8 (27.6)	1.00			
1	5 (21.8)	5 (17.2)	0.88	0.19	4.24	0.883
2 or more	9 (39.1)	16 (55.2)	0.50	0.14	1.75	0.279
Antimicrobial Therapy	0 (00.1)	. 5 (55.2)	3.00	····	0	5.2.0
Yes	21 (91.3)	25 (86.2)	0.59	0.09	3.58	0.571
No	2 (8.7)	4 (13.8)	1.00	0.00	0.00	0.07
Number of Classes	_ (0)	. ()				
None	2 (8.7)	4 (13.8)	1.00			
1 class	0	0	NA			
2 classes	13 (56.5)	16 (55.2)	1.62	0.25	12.45	0.607
3 classes or more	8 (34.8)	9 (31.0)	1.78	0.25	12.45	0.562
Period of Use	5 (04.0)	0 (01.0)	0	0.20	12.40	0.002
< 15 days	9 (39.1)	12 (41.4)	1.00			
≥ 15 days	14 (60.9)	17 (58.6)	1.09	0.36	3.36	0.870

OR = odds ratio; CI = confidence interval, NA = not applicable

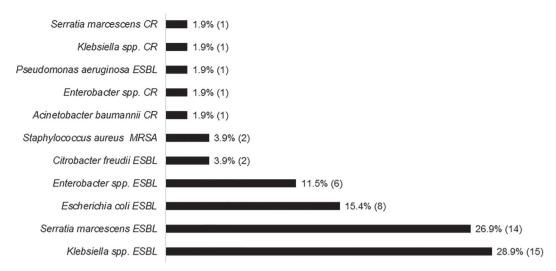


Figure 2 - Multidrug-resistant organisms isolated in colonization cultures collected at the time of infant discharge (*n*=52) from a neonatal unit, Parana State, Brazil, 2014-2018. CR: Carbapenems resistant; ESBL: Extended Spectrum β-lactamase; MRSA: methicillin/oxacillin resistant *Staphylococcus aureus*.

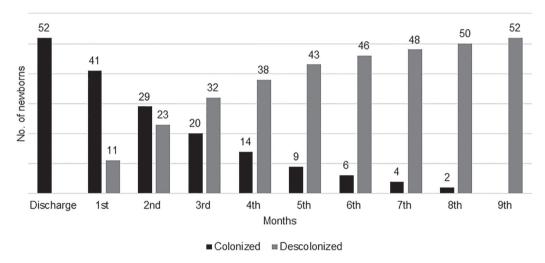


Figure 3 - Number of infants colonized and decolonized by multidrug-resistant organisms in the domiciliary context (*n*=52) according to the period in months after discharge, Parana State, Brazil, 2014-2018.

colonized by MDR readmitted to the NICU¹⁶. This was the only study that considered all the species of MDR, showing a mean period of 3.4 months for decolonization.

Regarding Gram-positive multidrug-resistant bacteria, only MRSA colonized infants in the current study, with an average time of 3 months for decolonization. This bacterial species is frequent in neonatal infections related to health care ^{19,29}. Given this threat, MRSA decolonization is recommended through the use of topical antimicrobial agents, however, as mentioned earlier, this practice is not recommended for infants due to the immaturity of their epithelial barrier, potential toxicity of the medication ¹⁹, as well as the risk of antimicrobial resistance and exposure to other pathogens ^{13,19}. In a cohort study conducted at a university medical center, after nasal decolonization with mupirocin and a chlorhexidine bath in infants colonized

with MRSA, 38% of infants were recolonized with the same microorganism and 16% developed MRSA infection, making this practice controversial for neonatal decolonization³⁰.

The limitations of the study were the failure to perform swab collection at hospital discharge, as well as the impossibility of investigating factors related to the use of antimicrobial therapy at home, considering the reports of the mothers and the absence of records on the medications used. However, this study presented as strengths its methodological rigor in the follow-up of infants after hospital discharge, as well as the investigation of different MDR species within a considerable neonatal population.

The novelty of the results of this research is in line with a relevant issue in global health, the antimicrobial resistance, contributing to fill some of the gaps in knowledge regarding

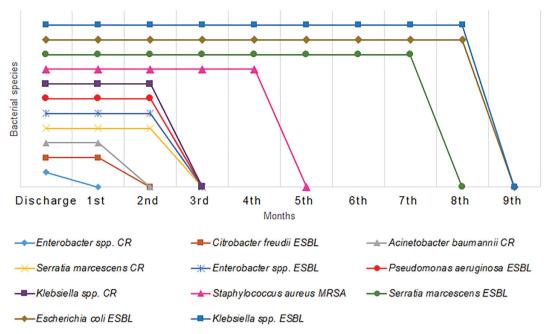


Figure 4 - Colonization period after discharge and decolonization according to multidrug-resistant organisms in infants in the domiciliary context, Parana State, Brazil, 2014-2018. CR: Carbapenems resistant; ESBL: Extended Spectrum β-lactamase; MRSA: methicillin/oxacillin resistant *Staphylococcus aureus*.

the period of colonization in infants by MDR after hospital discharge.

CONCLUSION

This research identified that the median period for spontaneous MDR decolonization in infants in the domiciliary setting was 3 months, considering all MDR species, and a maximum of 9 months, with a predominance of ESBL-producing Gram-negative bacteria.

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AUTHOR'S CONTRIBUTIONS

AMS wrote the manuscript and participated in data collection and analysis; GK, EGR, SFYO, MREP conceived, designed and supervised the study, wrote and performed data analysis; TNANI, KOP, NAAS, RLS and LGOP performed the laboratory procedures; RR gave substantial contributions for analysis and interpretation of data; JDC, MP, ECV, LMYL and MSAS participated in study supervision and data collection. All the authors read and approved the final manuscript.

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