

Bacteremia Due to Kosakonia cowanii in a Preterm Neonate

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1 **TITLE:** Bacteremia due to *Kosakonia cowanii* in a preterm neonate.

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22 ABSTRACT

Objective: Low-weight birth infants admitted to neonatal intensive care units are at high risk of hospital-acquired infections by opportunistic pathogens. The gut microbiota of preterm neonates lacks commensal bacteria providing a barrier against pathogens. We report a case of bacteremia due to *Kosakonia cowanii* in a preterm neonate.

Case report: Caesarian-section delivered a female baby of 680g at 28 weeks of 27 gestation due to intrauterine growth retardation and fetal rhythm abnormalities. On day 27, 28 two blood cultures grew Gram-negative bacilli in a context of functional ileus. No reliable 29 identification could be obtained using MALDI-TOF mass spectrometry, biochemical 30 reactions with the VITEK 2 GN ID card, or 16S rDNA sequencing. Kosakonia cowanii was 31 finally identified by gyrB sequencing. The source of infection may have been either the 32 central venous catheter or translocation from the gut microbiota. Evolution was favorable 33 after 14 days of cefepime (combined with amikacin for five days) and central venous catheter 34 removal. 35

36 **Conclusion:** *Kosakonia cowanii* is a member of the *Enterobacteriaceae* family that 37 was recently reclassified from the *Enterobacter* genus. Human infections due to *K. cowanii* 38 are scarce and have mainly been associated with traumatic inoculation from plants or transient 39 gut colonization. *K. cowanii* may be an underestimated opportunistic pathogen in susceptible 40 populations such as preterm neonates.

41 KEY-WORDS: Bacteriological Techniques; Kosakonia cowanii; Prematurity

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44 Introduction

Healthcare-associated infections frequently occur in neonatal intensive care units, with 45 a peak incidence reported between the 10th and 22nd day of life¹. Incidence of late-onset sepsis 46 is inversely correlated to birth weight: 36.3% of neonates with a gestational age under 28 47 weeks had at least one episode of sepsis compared to 29.6% for moderately preterm, 17.5% 48 for late preterm, and 16.5% for full term infants¹. Risk factors include the immature innate 49 immune system of preterm neonates, prolonged hospitalization, use of invasive medical 50 procedures, and failure of early enteral feeding with breast milk^{1,2}. Moreover, the gut 51 microbiota of preterm neonates lacks commensal bacteria providing resistance against 52 pathogens³. Thus, the risk of infection due to translocation from the intestinal tract may 53 increase. Indeed, Stewart et al. found that the the causative agent of late-onset sepsis in 54 preterm neonates was within the dominant genus of the gut microbiome⁴. Enterobacter 55 56 species are the most abundant bacteria present in preterm neonates' gut, especially after antimicrobial therapy⁵. Kosakonia cowanii is a member of the Enterobacteriaceae family that 57 was recently reclassified from the *Enterobacter* genus⁶. *K. cowanii* is frequently encountered 58 as a plant pathogen but rarely described in human infections. We present a case of bacteremia 59 due to K. cowanii in a preterm neonate. 60

61

62 **Case report**

Caesarian-section delivered a female baby of 680g at 28 weeks of gestation due to intrauterine growth retardation and fetal rhythm abnormalities. A central venous catheter was inserted, and enteral nutrition with pasteurized breast milk was initiated on day 1 of life. On day 10, the neonate presented bacterial sepsis due to *Staphylococcus epidermidis* : he had fever at 38°C, frequent apneas with bradycardia and desaturation. Continuous Positive Airway Pressure 68 (CPAP) and transfusion of one unit of packed red blood cells were required. The evolution69 was favorable after ten days of antimicrobial therapy with vancomycin.

On day 27, the baby presented abdominal distension with residual feeds attributed to a functional ileus, since she had no rectal bleeding nor vomiting. Hemodynamic and respiratory functions deteriorated again, requiring endotracheal intubation from day 28 to day 31, transfusion of five units of packed red blood cells and daily platelet transfusion from day 27 to day 34. Enteral feeding was temporarily stopped.

Blood cultures from day 27 yielded Gram-negative bacilli after 16 hours of incubation. No 75 reliable identification was assessed by MALDI-TOF spectrometry mass (Bruker Daltonics, 76 Wissembourg, France); the best score was 1.54 for K. cowanii. Biochemical reactions using 77 VITEK 2 GN ID card (BioMérieux, Marcy l'Etoile, France) identified the strain as Pantoea 78 spp., with a probability of 97%. However, the strain fermented sorbitol and did not use 79 80 malonate; those two characteristics distinguish K. cowanii from Pantoea agglomerans. Sequencing 16S rDNA performed universal of using primers 91E 81 was (TCAAAKGAATTGACGGGGGC-3) and 13BS (CCCGGGAACGTATTCAC) on the 3500 82 Dx Genetic Analyzer (Life Technologies, Cergy-Pontoise, France)⁷. Definitive identification 83 could not be obtained from the 16S partial sequence, because 99% identity out of 431 base 84 pairs was found for K. cowanii, Salmonella enterica, and Atlantibacter hermannii. Thus, gyrB 85 sequencing UP1 performed using the primers 86 was (GAAGTCATCATGACCGTTCTGCAYGCNGGNGGNAARTTYGA) 181r 87 and (CAGGAAACAGCTATGACCARRTGNGTNCCNCC) for amplification, followed by UP1-88 seq (GAAGTCATCATGACCGTTCTGCA) and 181r-seq (CAGGAAACAGCTATGACC) 89 for sequencing⁸. gyrB sequencing identified the isolate as K. cowanii, showing 99% identity 90 out of 521 base pairs with K. cowanii type strain JCM 10956 (Genebank accession number 91

JX425012.1). A phylogenetic tree based on *gyrB* sequences (Figure 1) was constructed in
MEGA X⁹, using the minimum evolution method.

In vitro antimicrobial susceptibility was determined by Mueller-Hinton agar disc diffusion 94 using the critical zone diameters recommended by the CASFM -EUCAST (version 2020) and 95 using an automated method (VITEK 2, BioMérieux, Marcy l'Etoile, France). Interestingly, 96 this strain did not produce an AmpC B-lactamase as expected from a former Enterobacter 97 species, but a penicillinase. The strain was resistant to amoxicillin (MIC \geq 32 mg/L), 98 99 ticarcillin (MIC \ge 128 mg/L), and piperacillin (MIC = 64 mg/L); all the other beta-lactams were active, including amoxicillin-clavulanate (MIC ≤ 2 mg/L) and cephalosporins (MIC ≤ 1 100 101 mg/L for cefotaxime, ceftriaxone, and cefepime).

102 After 14 days of cefepime, initiated on day 27 (combined with amikacin for five days) and 103 central venous catheter removal, evolution was favorable. Antimicrobial therapy and 104 laboratory values are represented as a timeline in Figure 2. Blood cultures remained sterile 105 after catheter change; *K. cowanii* also grew from the central venous catheter $(10^3 \text{ CFU}/\text{ mL})$, 106 indicating it could be the infection source. The neonate was discharged from the hospital after 107 three months of life, weighing 2615g for 48 cms.

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109 Discussion

Until 2013, the *Kosakonia* genus was part of the genus *Enterobacter*. Many species
from the *Kosakonia* genus (*K. radicincitans*, *K. sacchari*, *K. oryzae*, *K. cowanii*, *K. arachidis*)
have been isolated from plants, and only a few human infections have been described¹⁰. *K. cowanii* was first described from a patient sample but is mostly isolated from
environmental samples, including soil and trees. *K. cowanii* may promote plant growth and be
identified as an endophyte in trees with symptoms of blight⁶. Human infections due to *K. cowanii* are scarce. They have been associated with traumatic inoculation from plants. In a

117 case, rhabdomyolysis due to bacteremia was observed after rose thorn prick¹¹. More recently, 118 Berinson *et al.* reported a case of acute cholecystitis in an immunocompromised patient 119 without penetrating trauma¹², suggesting the origin of the infection may be food 120 contamination. The authors suggested that transient colonization of the gut by *K. cowanii* may 121 have caused acute cholecystitis in an immunocompromised patient.

Low-weight birth infants admitted in neonatal intensive care units are at high risk of 122 hospital-acquired infections by opportunistic pathogens². No case of K. cowanii opportunistic 123 infections in low-weight birth infants had been reported in the literature yet. Our case report 124 highlights that K. cowanii may be a relevant cause of infections in the population of preterm 125 neonates. Preterm neonates are vulnerable to infections caused by opportunistic foodborne 126 pathogens since their local natural barrier is immature, and secretory immunoglobulins A are 127 lacking during the first days of life. K. cowanii was previously isolated from powdered infant 128 milk formula¹³. However, food contamination may not be the origin of infection in our case 129 since the neonate was fed with pasteurized breast milk. In our patient, the source of infection 130 may have been bacterial translocation from the gut microbiota. Indeed, previous studies 131 showed that the pathogen identified in blood culture for late-onset sepsis in preterm infants 132 was usually abundant in the gut microbiome⁴. Additional risk factors of bacterial sepsis in 133 this population include a central venous catheter, parenteral nutrition, and steroids¹⁴. Another 134 hypothesis for the infection source may have been the central venous catheter, as suggested by 135 the positive bacteriological culture of the catheter. 136

Other environmental species, mostly former members of the *Enterobacter* group, were described as pathogens for neonates. The typical example is *Cronobacter sakazakii*, which has been isolated from environmental sources, milk, and infant formula, causing severe infections, especially bacteremia in preterm neonates^{6,15}. Powdered infant formula has been reported as the most commonly identified transmission vehicle of this bacterium¹⁵.

Berinson et al. suggested that K. cowanii might be an underestimated pathogen in 142 children¹². Indeed, in a retrospective study about pediatric infections due to 143 Pantoea agglomerans, seven patients out of 53 (13%) had a history of penetrating trauma 144 with a stick, plant thorn, or glass shard¹⁶. As shown in our case report, K. cowanii may be 145 falsely identified as *P. agglomerans* using biochemical reactions. It may be one of the reasons 146 for the few human infections described in the literature. The development of new 147 bacteriological techniques such as MALDI-TOF mass spectrometry and molecular biology 148 149 may improve our ability to identify this bacterium. K. cowanii is probably an underestimated opportunistic pathogen in susceptible populations such as preterm neonates. 150

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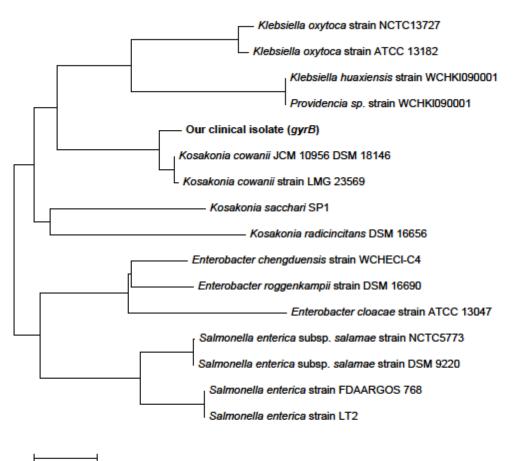
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Figure 1. Phylogenetic tree based on *gyrB* sequences.



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