

Rothia Bacteremia: a 10-Year Experience at Mayo Clinic, Rochester, Minnesota

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Rothia spp. are Gram-positive cocco-bacilli that cause a wide range of serious infections, especially in immunocompromised hosts. Risk factors for *Rothia mucilaginosa* (previously known as *Stomatococcus mucilaginosus*) bacteremia include prolonged and profound neutropenia, malignancy, and an indwelling vascular foreign body. Here, we describe 67 adults at the Mayo Clinic in Rochester, MN, from 2002 to 2012 with blood cultures positive for *Rothia*. Twenty-five of these patients had multiple positive blood cultures, indicating true clinical infection. Among these, 88% (22/25) were neutropenic, and 76% (19/25) had leukemia. Common sources of bacteremia were presumed gut translocation, mucositis, and catheter-related infection. One patient died with *Rothia* infection. Neutropenic patients were less likely to have a single positive blood culture than were nonneutropenic patients. Antimicrobial susceptibility testing was performed on 21% of the isolates. All of the tested isolates were susceptible to vancomycin and most beta-lactams; however, four of six tested isolates were resistant to oxacillin. There was no difference between the neutropenic and nonneutropenic patients in need of intensive care unit care, mortality, or attributable mortality.

Rothia mucilaginosa, previously known as *Stomatococcus mucilaginosus*, and other *Rothia* species (*R. dentocariosa*, *R. aeria*, *R. nasimurium*, and *R. amarae*) are part of the normal flora of the human oropharynx and upper respiratory tract (1). They are aerobic or facultatively anaerobic nonmotile non-spore-forming Gram-positive cocco-bacilli that can form filamentous branches (1). *Rothia* spp. are commonly associated with dental caries and periodontal disease (1). Invasive disease does occur, predominantly in immunocompromised hosts, but has rarely been reported in healthy hosts. The clinical syndromes associated with *Rothia* infection have included bacteremia (2), endocarditis (3), meningitis (4), peritonitis (5), bone and joint infections (1), pneumonia (6), skin and soft tissue infection, endophthalmitis (7), and prosthetic device infection (3). The main risk factors described for invasive disease have been hematological malignancy and severe neutropenia (8); other risk factors include diabetes mellitus, alcoholism, chronic liver disease, and infection with HIV (3). The clinical significance of *Rothia* spp. isolated from a blood culture is frequently unclear, especially in the setting of a single positive blood culture set with polymicrobial infection, suggesting contamination. To our knowledge, no large systematic evaluation of invasive *Rothia* infections has been published to date. We sought to describe the epidemiology and clinical significance of *Rothia* bacteremia in the past decade at our institution and to evaluate the differences in clinical outcomes between neutropenic and nonneutropenic patients. The other aim of our study was to describe the antimicrobial susceptibility pattern of *Rothia* isolates at our institution.

MATERIALS AND METHODS

We conducted a single-center retrospective cohort study of adult patients with blood cultures positive for *Rothia* between January 2002 and December 2012 at the Mayo Clinic in Rochester, MN. The protocol was approved by the Mayo Clinic Institutional Review Board. Cases were obtained by querying the microbiology records for *Rothia* spp. or *Stomatococcus* spp. grown from blood culture from 1 January 2002 to 31 December 2012 among patients presenting for medical care at the Mayo Clinic.

Patients were excluded from analysis if they were younger than 18 years of age or did not consent to the use of their medical records for research purposes.

The medical records of patients included for analysis were manually reviewed for patient demographics, medical comorbidities, antimicrobial exposure within the prior month, clinical outcomes, and microbiological data, including antimicrobial susceptibilities. The source of bloodstream infection was determined through a review of the medical records for a suspected source as documented by the attending infectious diseases physician (preferred) or by the attending physician of the primary service, if infectious diseases consultation was not involved in the patient's care. All documented potential sources were abstracted. If a suspected source was not documented, then the source was categorized as "no source identified." Attributable mortality was determined by two of the investigators (P.R. and P.K.T.), who reviewed the medical records of all patients who died with *Rothia* bacteremia. A case of mortality was considered nonattributable if a clear alternative cause of mortality was identified.

The Charlson comorbidity index score was used to assess the severity of underlying diseases (9). Analyses of microbiologic characteristics, including species identification and antimicrobial susceptibility, were performed on all the isolates recovered from blood culture. Analyses of clinical characteristics were performed on those with blood cultures positive for *Rothia* spp. from more than one set of blood cultures to minimize the inclusion of potential contaminants. A descriptive analysis was performed, as was an analysis to compare patients who were neutropenic (absolute neutrophil count, $\leq 1,000/\mu\text{l}$) at the time of bacteremia to those who were not neutropenic and to compare those with monomicrobial

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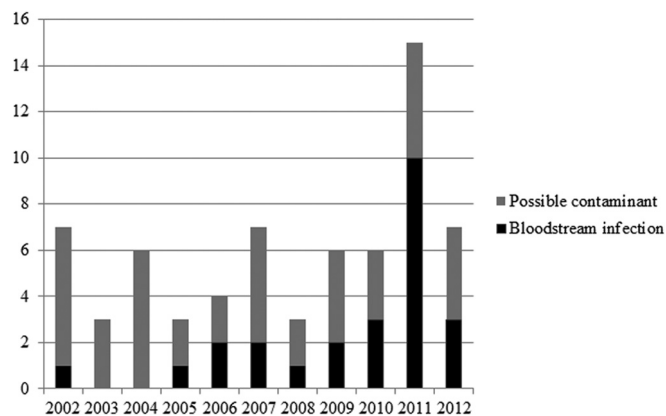


FIG 1 Distribution of patients with *Rothia* bacteremia per year at the Mayo Clinic in Rochester, MN (2002-2012).

infection to those with polymicrobial infection. Two-tailed Fisher’s exact and Kruskal-Wallis tests were used for comparisons of proportions and medians, respectively, with P values of <0.05 considered statistically significant. At our institution, blood samples are cultured using the Bactec instrumented blood culture system (Becton, Dickinson, and Company, Franklin Lakes, NJ, USA). Antimicrobial susceptibilities were determined using agar dilution with isolates from a pure subculture. The clinical microbiology laboratory at the Mayo Clinic uses CLSI breakpoints for *Corynebacterium* for reporting the susceptibilities of *Rothia* isolates. Most of the *Rothia* isolates were identified using morphological and biochemical properties; recent isolates were identified using matrix-assisted laser desorption ionization–time of flight mass spectrometry (MALDI-TOF MS).

RESULTS

After excluding 5 patients younger than 18 years of age, we identified 67 adults from 2002 to 2012 with blood cultures positive for *Rothia*; for 42 patients, *Rothia* grew from a single blood culture set and was considered potentially contaminated, and the remaining 25 patients were considered to have a true bloodstream infection (Fig. 1). All 67 patients identified with blood cultures that were positive for *Rothia* had at least two sets of blood cultures drawn at the same time, with at least one being from the periphery; for all of the cases in which only one set was positive for *Rothia*, there was a second blood culture set drawn at the same time from which *Rothia* did not grow. Neutropenic patients were significantly less likely to have a single positive blood culture set than were nonneutropenic patients (41% versus 90%, $P < 0.001$). The demographic, clinical, and microbiological characteristics and clinical outcomes of the patients considered to have had true bloodstream infection are summarized in Table 1. Twenty-two (88%) patients were neutropenic with a median of 9.5 days of neutropenia (interquartile range, 7 to 20 days) at the time of *Rothia* bloodstream infection, and 19 (76%) had an underlying diagnosis of leukemia.

The most common sources of *Rothia* bloodstream infection were presumed gut translocation ($n = 13$, 52%), catheter-related infection ($n = 8$, 25%), and mucositis ($n = 9$, 36%). No source of bacteremia was identified for 5 (20%) of the patients.

A comparison of the clinical characteristics and outcomes of patients with monomicrobial infection ($n = 16$) and those with polymicrobial infection ($n = 9$) identified no qualitative or statistically significant differences between the two groups for any of the variables collected, including median age (56 years versus 53 years, $P = 0.93$), recent corticosteroid use (6% versus 11%, $P = 1.0$),

presence of neutropenia (88% versus 89%, $P = 1.0$), ability to perform susceptibility testing (20% versus 33%, $P = 0.63$), median duration of hospital stay (20 days versus 21 days, $P = 0.32$), median duration of antimicrobial treatment (14 days versus 14 days, $P = 0.57$), need for intensive care unit (ICU) care (44% versus 44%, $P = 1.0$), or attributable mortality (0% versus 11%, $P = 0.36$). Of the nine polymicrobial *Rothia* bloodstream infections, five cases had additional growth of coagulase-negative staphylococci (three attributed to central line-related bloodstream infection, one attributed to presumed gut translocation, and one for which a source was not identified), one case had additional growth of viridans group streptococci attributed to presumed gut translocation, one case had additional growth of *Enterococcus faecium* for which a source was not identified, one case had additional growth of *Candida dubliniensis* attributed to central line-related bloodstream infection, and one case had additional growth of *Clostridium inoculum* attributed to presumed gut translocation.

We identified one patient with a *Rothia* bloodstream infection who died, potentially as a result of the infection. The patient was a 51-year-old man who developed acute abdominal pain, fever, and hypotension in the setting of profound neutropenia 4 days after allogeneic peripheral blood stem cell transplantation for multiple myeloma. Intravenous vancomycin, cefepime, metronidazole, fluconazole, and acyclovir were started empirically, and the patient was transferred to the ICU, where he required ventilatory support, vasopressor support, and dialysis. The blood culture bottles from all the sets were positive for vancomycin-susceptible *Enterococcus faecium* and *Rothia mucilaginosa* resistant to oxacillin (no other antimicrobial susceptibility tests were performed) without any significant differential time to positivity between peripherally drawn cultures and those drawn from the central venous catheter. He was felt to be too unstable to transport for computed tomography (CT) imaging of his abdomen. Despite maximal support, the patient developed cardiac arrest and died within 48 h of his initial presentation. An autopsy was not performed.

We also identified a 28-year-old woman with *Rothia* prosthetic aortic valve endocarditis (monomicrobial) who had a history of intravenous drug use and prior native tricuspid and aortic valve endocarditis with methicillin-susceptible *Staphylococcus aureus*. The diagnosis was confirmed with a transesophageal echocardiogram. There was not adequate growth to perform susceptibility testing, and the patient was treated with a 6-week course of intravenous ceftriaxone and vancomycin and with oral rifampin; 1 year after the completion of therapy, there was clinical resolution of the infection and no evidence of relapsed disease.

All of the 25 clinical bloodstream infections involved *Rothia mucilaginosa*, although one of the isolates from a patient with a single positive blood culture set (possible contaminant) involved *Rothia dentocariosa*. Due to poor growth of the organism on Mueller-Hinton agar (supplemented with 5% lysed horse blood), we were able to perform susceptibility testing on only 14 (21%) of the total 67 isolates and on 6 (24%) of the 25 isolates from true bloodstream infection. There was no significant association between the ability to grow the organism for susceptibilities and whether or not the patient received penicillin (40% versus 20%, $P = 0.29$), levofloxacin (25% versus 18%, $P = 0.55$), cefepime (9% versus 24%, $P = 0.43$), or vancomycin (14% versus 23%, $P = 0.72$) within 30 days prior to bacteremia. All tested isolates were susceptible to penicillin (9/9), ceftriaxone (8/8), ertapenem (2/2), meropenem (8/8), and vancomycin (13/13). Four isolates were resis-

TABLE 1 Demographic and clinical characteristics and clinical outcomes of patients with *Rothia* bloodstream infections at Mayo Clinic, Rochester, MN (2002–2012)

Variable	Data ^a for:			P value
	All patients (n = 25)	Neutropenic patients (n = 22)	Nonneutropenic patients (n = 3)	
Age (median [range]) (yr)	55 (22–78)	60 (22–78)	30 (28–48)	0.03
Male sex	19 (76)	18 (82)	1 (33)	0.2
Comorbidities				
Myocardial infarction	2 (8)	2 (9)	0 (0)	1.0
Congestive heart failure	1 (4)	0 (0)	1 (33)	0.12
Peripheral vascular disease	1 (4)	1 (5)	0 (0)	1.0
Chronic lung disease	1 (4)	1 (5)	0 (0)	1.0
Connective tissue disease	1 (4)	1 (5)	0 (0)	1.0
Diabetes mellitus	2 (8)	2 (9)	0 (0)	1.0
Solid tumor	3 (12)	3 (14)	0 (0)	1.0
Leukemia	19 (76)	19 (86)	0 (0)	0.009
Lymphoma/multiple myeloma	4 (16)	4 (18)	0 (0)	1.0
Moderate to severe liver disease	1 (4)	1 (5)	0 (0)	1.0
Moderate to severe kidney disease	2 (8)	2 (9)	0 (0)	1.0
Hemodialysis	1 (4)	1 (5)	0 (0)	1.0
Hematopoietic stem cell transplant				
Allogeneic	4 (16)	4 (18)	0 (0)	1.0
Autologous	2 (8)	2 (9)	0 (0)	1.0
Charlson comorbidity index				
0	1 (4)	0 (0)	1 (33)	0.002
1	1 (4)	0 (0)	1 (33)	
2	3 (12)	3 (14)	0 (0)	
3	3 (12)	3 (14)	0 (0)	
4	3 (12)	3 (14)	0 (0)	
5	8 (32)	8 (36)	0 (0)	
6	1 (4)	0 (0)	1 (33)	
7	5 (20)	5 (23)	0 (0)	
Medication use within 30 days prior to bacteremia				
Corticosteroid	2 (8)	2 (9)	0 (0)	1.0
Antimicrobial	25 (100)	22 (100)	3 (100)	1.0
Levofloxacin	21 (84)	20 (91)	1 (33)	0.06
Penicillin	3 (12)	3 (14)	0 (0)	1.0
Presence of central venous catheter at the time of bacteremia	24 (96)	22 (100)	2 (67)	0.12
Source of bloodstream infection				
Presumed gut translocation with negative CT of abdomen	12 (48)	9 (41)	3 (100)	0.28
Presumed gut translocation without a CT of abdomen	1 (4)	1 (5)	0 (0)	1.0
Neutropenic colitis with consistent CT of abdomen	1 (4)	1 (5)	0 (0)	1.0
Catheter-related bloodstream infection	8 (32)	6 (27)	2 (66)	0.23
Endocarditis	1 (4)	0 (0)	1 (33)	0.12
Dental abscess	2 (8)	2 (9)	0 (0)	1.0
Mucositis	9 (36)	9 (41)	0 (0)	0.28
No source identified	5 (20)	5 (23)	0 (0)	1.0
Presence of polymicrobial infection	9 (36)	8 (36)	1 (33)	1.0
Antimicrobial susceptibility test performed	6 (24)	6 (27)	0 (0)	1.0
Duration of hospitalization (mean [range]) (days)	21 (1–53)	24 (1–53)	12 (6–21)	0.32
Need for ICU care	11 (44)	10 (45)	1 (33)	1.0
Complications				
Septic shock	3 (12)	3 (14)	0 (0)	1.0
Respiratory failure	3 (12)	3 (14)	0 (0)	1.0
Liver failure	1 (4)	1 (3)	0 (0)	1.0
Renal failure	3 (12)	2 (9)	1 (33)	0.33
Altered mental status	2 (8)	2 (9)	0 (0)	1.0

(Continued on following page)

TABLE 1 (Continued)

Variable	Data ^a for:			P value
	All patients (n = 25)	Neutropenic patients (n = 22)	Nonneutropenic patients (n = 3)	
Attributable mortality	1 (4)	1 (3)	0 (0)	1.0
Antimicrobial treatment				
Cefepime	9 (36)	9 (41)	0 (0)	0.28
Ceftriaxone	1 (4)	1 (4)	0 (0)	1.0
Piperacillin-tazobactam	5 (20)	5 (23)	0 (0)	1.0
Ertapenem	3 (12)	3 (14)	0 (0)	1.0
Meropenem	4 (16)	4 (18)	0 (0)	1.0
Daptomycin	3 (12)	2 (9)	1 (33)	0.33
Linezolid	2 (8)	2 (9)	0 (0)	1.0
Vancomycin	20 (80)	18 (82)	2 (67)	0.50
Duration of antimicrobial treatment (median [range]) (days)	14 (2–42)	14 (2–28)	30 (10–42)	0.17

^a Data are no. (%) unless stated otherwise.

tant to oxacillin (4/6), but concomitant penicillin susceptibility testing was not performed on them.

DISCUSSION

We conducted a retrospective review of all adult patients with blood cultures positive for *Rothia* in a single academic institution during the past decade and have described the epidemiology and clinical characteristics of *Rothia* bacteremia. In addition, we performed a comparative analysis of clinical outcomes between neutropenic and nonneutropenic patients and between monomicrobial and polymicrobial infections. *Rothia* bloodstream infections occurred often in patients with significant medical comorbidities, most commonly hematologic malignancy. A majority of the patients were exposed to at least one antimicrobial agent (predominantly a fluoroquinolone) within the month preceding the infection. Most of these patients had an indwelling central venous catheter at the time of bacteremia, most likely related to the need for central venous access for the administration of chemotherapeutic agents (for patients with hematologic malignancy). There was no apparent temporal change in the incidence of bacteremia in the past decade, although a transient increase was noted in 2011; the reason for this transient increase is not clear, as there were no notable changes in patient management or concomitant increases in central line-related infections elsewhere in our institution. In this study, *Rothia mucilaginosa* caused all of the clinical bloodstream infections. Gut translocation was the most commonly identified source of *Rothia* bloodstream infection, although central line-related infections and mucositis were also common. There were no differences detected in clinical characteristics or clinical outcomes between those with monomicrobial and those with polymicrobial *Rothia* bloodstream infection.

When encountering *Rothia* bacteremia in clinical practice, many clinicians are faced with the challenge of deciding whether it represents a true bloodstream infection or contamination; 63% of the *Rothia* isolates in our study were potential contaminants since they grew from a single positive blood culture. Transient *Rothia* bacteremia has been reported in the literature, and its clinical significance remains unknown (10). In our study, neutropenic patients were less likely to have single blood culture set positivity than were nonneutropenic patients. This indicates that when

Rothia bacteremia is identified in neutropenic patients, it is likely to represent true infection. We did find that the vast majority of neutropenic patients had an underlying diagnosis of leukemia and had prolonged and profound neutropenia at the time they were diagnosed with *Rothia* bacteremia. Potential reasons for this predilection in patients with leukemia include a higher preponderance of mucositis due to the chemotherapeutic agents used to treat the underlying disease and prolonged duration of chemotherapy-induced neutropenia. Furthermore, the vast majority of patients undergoing chemotherapy for leukemia at our institution receive levofloxacin prophylaxis during neutropenia, so there may be a shift in the oral and gastrointestinal flora away from aerobic Gram-negative bacilli and toward other pathogens such as *Rothia*. The presence of polymicrobial *Rothia* bloodstream infection did not appear to be associated with higher-risk clinical characteristics or portend less favorable clinical outcomes than those with monomicrobial infection. We did not record whether the other organisms were identified in more than one blood culture set, raising the possibility that they were contaminants, but the *Rothia* species were identified in more than one blood culture set, which argues against its presence being a result of contamination.

Rothia (Stomatococcus mucilaginosa) bacteremia in neutropenic patients was first described in the 1990s (8, 11, 12). Ascher et al. described 10 patients with *Rothia mucilaginosa* bacteremia, 5 of whom had more than one positive blood culture. Among these 5 patients, 3 were neutropenic and had malignancies; all of them had an indwelling vascular foreign body. Most patients recovered with vancomycin (11). Henwick et al. characterized 8 cases of *Rothia mucilaginosa* bacteremia in children with cancer; 6 of them had leukemia, 7 had profound neutropenia, 4 had mucositis, and 5 had central venous catheters. Despite the prompt initiation of antibiotics, the rate of complications in this cohort were high (eg, septic shock [50%], pneumonia, altered mental status, meningitis, and acute respiratory distress syndrome). All the isolates were susceptible to vancomycin, but 50% were penicillin resistant and 29% were methicillin resistant (12). Fanourgiakis et al. described 8 patients with *Rothia mucilaginosa* bacteremia, the majority of whom (7/8) had hematological malignancies (6 leukemia); 1 had breast cancer. All these patients had profound neutropenia and

chemotherapy-induced disruption in the oral or gut mucosal barrier. All of the patients were on quinolone prophylaxis at the time of bacteremia; 5 of 6 tested isolates were quinolone resistant (13). One patient in our study expired potentially as a result of *Rothia* infection, despite organism identification, prompt appropriate antimicrobial initiation, and intense supportive management. He possessed many of the previously described risk factors for *Rothia* infection, demonstrating the pathogenicity of *Rothia* despite it normally being considered a benign and colonizing organism. However, the case description suggests that the patient may have died due to catastrophic gut wall breach with resultant polymicrobial bloodstream infection rather than from the pathogenicity of *Rothia* bloodstream infection itself. The mortality rates attributable to *Rothia* infections have varied in the literature according to age, immune status, and site of infection (14–16). Immunocompromised patients are more likely to develop severe complications, including death, from *Rothia* infections. To our knowledge, ours is the largest published cohort of patients with *Rothia* bacteremia. In prior publications, *Rothia mucilaginosa* isolates were generally susceptible to most beta-lactam antimicrobials (penicillin, ampicillin, imipenem, cefotaxime), rifampin, and vancomycin (17). However, isolates with partial resistance to penicillin have been described in the past (17). In one study, the incidences of penicillin and methicillin resistance among isolates were 50% and 29%, respectively (12). Antimicrobial susceptibility testing was not performed on most of the *Rothia* isolates in our study, owing to their poor growth *in vitro*, even when the growth medium was supplemented with 5% lysed horse blood. When testing was possible, all of the isolates from our study were susceptible to penicillin, ceftriaxone, meropenem, and vancomycin; however, four of six isolates were resistant to oxacillin. The reasons for this pattern of susceptibility are not clear.

The reason for the preponderance of cases in our study receiving vancomycin as part of dual therapy is likely because most of the patients had *Rothia* bloodstream infection in the setting of febrile neutropenia. Vancomycin is a recommended empirical antimicrobial agent (in combination with an antipseudomonal beta-lactam antimicrobial) for the treatment of neutropenic fever with Gram-positive bloodstream infection (18). Upon clearance of the bloodstream infection and identification of the causative organism, clinicians may have been inclined to continue with vancomycin, especially if susceptibility data were not available. The results of our study suggest that the addition of vancomycin to neutropenic fever therapy when Gram-positive bloodstream infection is identified would provide appropriate coverage for *Rothia* infections and that ceftriaxone is likely to be an effective definitive antimicrobial agent in a clinical setting when patients have clinically improved but antimicrobial susceptibility results are not available.

Further research is needed to develop microbiologic techniques to improve our ability to provide antimicrobial susceptibility results in cases of *Rothia* infection. Additionally, the potential role of fluoroquinolone prophylaxis in shifting oral and gastrointestinal flora in patients undergoing chemotherapy for hematologic malignancy needs further exploration.

In conclusion, members of the genus *Rothia*, despite their low virulence, have established themselves as significant pathogens, especially in patients with hematological malignancies and neutropenia. Mucositis and central venous catheters are common predisposing factors, and both of these factors are related to treat-

ment for hematologic malignancy. Neutropenic patients are more likely to have true bloodstream infection as evidenced by multiple positive blood culture sets and monomicrobial infection. There was no significant difference in clinical outcomes between the neutropenic and nonneutropenic patients. At this time, there are limited data available on the antimicrobial susceptibility patterns of *Rothia*; however, isolates are generally susceptible to vancomycin and beta-lactam antimicrobials, except oxacillin.

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