

Incidence of Bloodstream Infections in a Speciality Hospital in Kuwait: 8-Year Experience

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Key Words

Bloodstream infections · Frequency of isolation · Antimicrobial susceptibility

Abstract

Objectives: To determine the frequency of isolation and antibiotic-susceptibility patterns of clinically significant bacterial pathogens isolated from blood. **Materials and Methods:** The study was conducted over a period of 8 years (1995–2002) at Infectious Diseases Hospital (IDH), Kuwait. Demographic and clinical data were obtained from medical records. 18,535 blood cultures were analyzed. Disk diffusion method was used to perform antibiotic-susceptibility testing. Minimum inhibitory concentrations of 9 antimicrobials were determined using E-test. Double disk (potentiation) test and E-test ESBL strips were used to detect the production of extended-spectrum beta-lactamases (ESBLs). **Results:** *Salmonella* spp. and *Brucella* spp. were predominant blood isolates, and represented 60.6 and 30.0% of all clinically significant episodes of bloodstream infections, respectively. Among the *Salmonella*, *Salmonella enterica* serotypes *typhi* and *paratyphi A* were most frequently isolated. The percentage of multidrug resistance (MDR) among them varied from 22 to 51%. A high percentage (40%) of MDR *S. enterica* serotypes *typhi* and *paratyphi A* also showed reduced susceptibility to ceftriaxone and ciprofloxacin.

Conclusion: During the study period, *Salmonella* spp. and *Brucella* spp. were predominant blood isolates. MDR *S. enterica* serotypes *typhi* and *paratyphi A*, with reduced susceptibility to ceftriaxone and ciprofloxacin, are among the most frequent causes of bloodstream infections in IDH, suggesting the need to monitor their susceptibility.

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Introduction

Bacterial bloodstream infections (BIs) are a leading cause of morbidity and mortality worldwide [1]. The detection of bacteremia and septicemia remains one of the most important services provided by clinical bacteriology laboratories [2, 3]. Management of infection in many cases is based on an empirical basis, especially in immunocompromised or elderly patients, who may not present with the usual clinical signs and symptoms. It is therefore important to be able to detect bacteremia/septicemia and to identify the causative pathogens and determine antibiotic susceptibility, so that optimal patient management can be instituted as early as possible. There are several studies available on BIs in large general and university hospitals [4–6], but fewer studies have been conducted in small and specialized hospitals. Several reports dedicated on *Salmonella enterica* serotypes *typhi*, *paratyphi A* and *Brucella* spp. (as predominant blood isolates in Infectious

Diseases Hospital) have been published [7–9]. However, no report is available presenting the total picture of BIs in Infectious Disease Hospital, Kuwait, hence this retrospective study was carried out to characterize the BIs to determine the frequency of bacterial isolation and their antimicrobial susceptibility pattern.

Materials and Methods

The study was carried out in the Infectious Diseases Hospital, Kuwait over a period of 8 years (1995–2002). The Infectious Diseases Hospital is a 151-bed specialized institution serving the entire population of Kuwait. The average number of admissions to the hospital is 4,000 per year. All the patients with bacteremia diagnosed by blood culture were included in the study. Data were collected from the microbiology and patient's medical records.

Blood Cultures

Blood was drawn aseptically, using Vacutainer system (Becton Dickinson) from all adult and pediatric inpatients and outpatients with suspected BIs and directly inoculated into BD Bactec Plus Aerobic/F, BD Bactec Plus Anaerobic/F and BD Bactec Peds/F bottles. The volume of blood obtained from adult was 5–10 ml and for the children 1–3 ml as previously recommended [10]. The Bactec 9120 (Becton Dickinson Diagnostic Instrument System, Sparks, Md., USA), a continuous-monitoring blood culture system, was used for further monitoring of blood cultures. The protocols of 7-day incubation for routine cultures and 21 days for fastidious microorganisms (including *Brucella*) were adopted.

All bottles positive in the system were gram-stained and subcultured on suitable media (blood agar, chocolate agar, MacConkey agar, *Brucella* blood agar, BBA). If no organisms were seen, the bottle was returned to the system to complete the testing cycle. All clinically significant bacterial isolates were identified according to standard methods [11].

Antimicrobial Susceptibility Testing

The isolated microorganisms were tested by Kirby-Bauer disk diffusion method on Mueller-Hinton agar and BBA, and the results were interpreted according to the current National Committee for Clinical Laboratory Standards (NCCLS) guidelines [12]. Antimicrobial agents (disks) were obtained from their respective manufacturers. *Escherichia coli* ATCC 25922, *Staphylococcus aureus* ATCC 29213, *Pseudomonas aeruginosa* ATCC 27853, *Staphylococcus epidermidis* ATCC12228, *Streptococcus pneumoniae* ATCC 6503 and *Enterococcus faecalis* ATCC 19212 were used as quality control strains. Minimum inhibitory concentrations (MICs) of ampicillin, amoxicillin, clavulanic acid, cephalothin, cefuroxime, ceftriaxone, piperacillin, Tazocin, gentamicin and ciprofloxacin were determined using E-test strips (AB Biodisk, Sweden) [13]. Antimicrobial susceptibility testing for *Brucella* isolates was performed on BBA. Since the disk diffusion method is not standardized for *Brucella*, the results were not categorized as susceptible, intermediate or resistant. Instead, the diameter of the zone of inhibition was used as indirect indicator to measure anti-*Brucella* activity. Inhibition zones of <16 mm and ≥16 mm were considered to represent low and good activity, respectively [9].

Detection of Extended-Spectrum Beta-Lactamase (ESBL) Production

Evaluation of isolates for ESBL production was done by double disk (potentiation) test and confirmed by E-test ESBL strip of Cormican et al. [13].

Data Interpretation

Episode of blood infection (BI) was defined by the isolation of one or more microorganisms from blood culture together with clinical evidence of systemic infection. A new episode was recorded for the same patient if the initial pathogen had previously been eradicated from the bloodstream or if an interval of at least 1 month had elapsed without signs of infection since the earlier episode.

The significance of isolates was judged according to the identity of the microorganism, the presence of more than one blood culture set for the same microorganism, the presence of the same microorganism as that found in the blood from another normally sterile site. The strains were regarded as multidrug resistant (MDR) if they showed resistance to three or more different classes of antibiotics.

Results

The distribution of patients by age was comparable during different years of the study (mean of 36.7 years and range of 0–65 years). Patients more than 60 years of age and pediatric age group accounted for about 8 and 9% of the total number of BIs, respectively. The male-to-female ratio was the same for all age groups (2.5:1).

During this period, 1,129 cases of bacteremia with 830 clinically significant episodes were diagnosed (table 1). Of the 18,535 blood cultures performed, 1,702 (9.2%) were positive and 1,129 species of microorganisms were detected. Of these, 299 (26.5%) were judged to be contaminants or to represent transient bacteremia without clinical significance. Coagulase-negative staphylococci (CNS), diphtheroids, *Propionibacterium* spp., *Bacillus* spp., *Acinetobacter* spp., *Pseudomonas* spp., representing 1.6% of total blood cultures, were considered contaminants. A total of 61.4% of the contaminants were CNS (*S. epidermidis*, *S. capitis*, *S. hominis*, *S. haemolyticus* and *S. auricularis*). The remaining blood cultures (n = 830), yielded clinically significant pathogens. The frequency distributions of these isolates are shown in table 2. Among all bacterial isolates, *Salmonella* spp. (503, 44.5%) and *Brucella* spp. (249, 22%) were predominant isolates. The other members of the family Enterobacteriaceae (*E. coli*, *Klebsiella* spp., *Enterobacter* spp.) accounted for only 7% of the total isolates. None of these produced ESBL. During the study period, we observed only a few cases of bacteremia caused by fastidious microorganisms such as nontypable serotypes of *Haemophilus influenzae* (n = 2),

Table 1. Details of bacterial isolates from blood cultures and their frequency of isolation

Year	Blood cultures			Cases of bacteremias	Bacterial isolates															
	total number performed	positive	sterile		<i>S. typhi</i>	<i>S. para. A</i>	<i>Salm. spp.</i>	<i>E. coli</i>	<i>Kleb. spp.</i>	<i>Ent. spp.</i>	<i>Brucella</i>	<i>H. infl.</i>	<i>N. mening.</i>	<i>S. aur.</i>	<i>S. pn.</i>	GN-NFB	CNS	<i>P. acnes</i>	Diphtheroids	<i>Bacillus</i>
1995	2,359	186	2,173	150	45	6	6	6	-	1	36	-	-	1	-	3	38	6	2	-
1996	2,943	274	2,669	199	95	18	3	5	5	2	16	-	-	-	-	2	37	4	2	10
1998	1,729	202	1,527	152	60	8	3	7	-	-	37	-	-	-	-	-	21	4	3	9
1999	1,594	180	1,414	134	62	7	-	4	3	-	24	2	-	2	4	1	18	2	-	5
2000	2,496	250	2,246	120	30	7	-	2	4	-	42	-	-	3	1	1	16	2	5	7
2001	2,789	202	2,587	115	20	3	1	4	-	-	46	-	-	5	-	2	22	-	4	8
2002	2,421	183	2,238	99	22	6	-	3	4	-	38	-	1	1	-	3	18	1	2	-
Total %	18,535	1,702	16,833	1,129	414	73	16	37	18	3	249	2	1	12	5	12	191	31	23	42
		9.2	90.8		36.7	6.5	1.4	3.3	1.6	0.3	22.0	0.2	0.1	1.1	0.4	1.1	16.7	2.7	2.0	3.7

S. para. A = *S. enterica* serotype *paratyphi A*; *Salm.* = *Salmonella*; *Kleb.* = *Klebsiella*; *Ent.* = *Enterobacter*; *H. infl.* = *H. influenzae*; *N. mening.* = *N. meningitidis*; *S. aur.* = *S. aureus*; *S. pn.* = *S. pneumoniae*; GN-NFB = gram-negative nonfermenting bacteria; *P.* = *Propionibacterium acnes*.

Table 2. Details of bacterial isolates from 830 clinically significant BIs (1995–2002)

Microorganisms	Number	%
<i>S. enterica</i> serotype <i>typhi</i>	414	49.9
<i>S. enterica</i> serotype <i>paratyphi A</i>	73	8.8
<i>S. enterica</i> serotypes (other than <i>typhi</i> and <i>paratyphi A</i>)	16	1.9
<i>E. coli</i>	37	4.5
<i>Klebsiella spp.</i>	18	2.2
<i>Enterobacter spp.</i>	3	0.4
<i>Brucella spp.</i>	249	30.0
<i>S. aureus</i>	12	1.4
<i>H. influenzae</i>	2	0.2
<i>N. meningitidis</i>	1	0.1
<i>S. pneumoniae</i>	5	0.6
Total species	830	

Neisseria meningitidis (n = 1), *S. pneumoniae* (n = 5). We also encountered a single episode of BI by gram-negative anaerobe, *Bacteroides spp.*

In less than 12 h 15% of all positive blood cultures were detected; within 24 h: 60%; 48 h: 70%; 72 h: 78% and within 7 days the remaining 22% were detected. At 12 h 45% of *S. aureus* infections and 55% of the facultative gram-negative enteric bacteria were detected. Fourteen percent of the CNS and 18% of the diphtheroids and *Propionibacterium spp.* were detected within 12 h. Seventy-three percent and 56% of significant and nonsignificant blood cultures, respectively, were detected during the first

Table 3. Antimicrobial susceptibility patterns (%) of *S. enterica* serotype *typhi* recovered from blood

		Antimicrobials					
		Am	Sxt	C	Ctx	Cro	Cip
1995 (n = 45)	S	38.5	65.9	65.8	100.0	100.0	100.0
	R	61.5	34.1	34.2	0	0	0
1996 (n = 95)	S	69.7	72.4	69.3	100.0	100.0	100.0
	R	30.3	27.6	30.7	0	0	0
1997 (n = 80)	S	47.0	56.3	48.2	100.0	100.0	100.0
	R	53.0	43.7	51.8	0	0	0
1998 (n = 60)	S	68.1	68.8	69.6	100.0	100.0	100.0
	R	31.9	31.2	30.4	0	0	0
1999 (n = 62)	S	68.1	63.2	65.8	100.0	100.0	100.0
	R	31.9	36.8	34.2	0	0	0
2000 (n = 30)	S	65.7	65.7	65.7	100.0	100.0	100.0
	R	34.3	34.3	34.3	0	0	0
2001 (n = 20)	S	77.3	77.3	77.3	100.0	100.0	100.0
	R	22.7	22.7	22.7	0	0	0
2002 (n = 22)	S	52.0	52.0	52.0	100.0	100.0	100.0
	R	48.0	48.0	48.0	0	0	0

S = Sensitive; R = resistant; Am = ampicillin; Sxt = trimethoprim-sulfamethoxazole; C = chloramphenicol; Ctx = cefotaxime; Cro = ceftriaxone; Cip = ciprofloxacin.

24 h of incubation. In all episodes of *Brucella* and gram-negative nonfermenting bacteria, the microorganisms were isolated only from the aerobic bottle.

The antimicrobial susceptibility results of clinically significant blood isolates are shown in tables 3 and 4. The

Table 4. Antimicrobial susceptibility results (%) of *E. coli* and *Klebsiella* spp. recovered from blood cultures

		Antimicrobial agents								
		Am	Amc	Cf	Cxm	Cro	Pip	Tzp	Gm	Cip
<i>E. coli</i> (n = 37)	S	30.6	64.0	74.5	97.9	98.0	50.4	87.8	92.8	95.0
	R	69.4	36.0	25.5	2.1	2.0	49.6	12.8	7.2	5.0
<i>Klebsiella</i> spp. (n = 18)	S	0	76.0	80.0	93.0	97.0	71.4	91.0	100.0	100.0
	R	100.0	24.0	20.0	7.0	3.0	28.6	9.0	0	0

S = Sensitive; R = resistant; Am = ampicillin; Amc = amoxicillin/clavulanic acid; Cf = cephalothin; Cxm = cefuroxime; Cro = ceftriaxone; Pip = piperacillin; Tzp = Tazocin; Gm = gentamicin; Cip = ciprofloxacin.

percentage of MDR strains of *Salmonella enterica* serotype *typhi* varied from approximately 31% (1995, 1996) to 44% in 1997 (table 3). During the second part of the period (1998–2001), the number of MDR *S. typhi* decreased to 23% in 2001. Antibacterial activity of ceftriaxone and ciprofloxacin against *S. typhi* (determined by disk diffusion method) remained stable during the study period but MICs for ceftriaxone increased from 0.19 to 0.25 µg/ml and ciprofloxacin from 0.35 to 1.0 µg/ml. Both *E. coli* and *Klebsiella* spp. showed a near-similar susceptibility pattern all through the study period. About 69.4, 49.6, 36 and 25.5% were resistant to ampicillin, piperacillin, amoxicillin/clavulanic acid and cephalothin, respectively (table 4).

All the *Brucella* isolates demonstrated good susceptibility to traditional anti-*Brucella* drugs such as trimethoprim, tetracycline, streptomycin, gentamicin, amikacin and ciprofloxacin. Rifampicin and sulfamethoxazole had variable in vitro anti-*Brucella* activity with 8 and 25% of the isolates, respectively, thereby showing low potency. Not a single isolate of methicillin-resistant *S. aureus* was isolated, 2 of the 5 isolates of *S. pneumoniae* expressed intermediate susceptibility to penicillin (MICs = 0.75 µg/ml).

Discussion

BIs have been reported as an important medical problem [4, 14–16], despite advances in antimicrobial therapy and supportive care. It is essential to evaluate prospectively the distribution of bacterial species isolated from blood and their susceptibility to the major antimicrobial agents and alternative drugs to adapt appropriate antibiotic therapy strategies. Ceftriaxone and ciprofloxacin are

drugs of choice for the treatment of MDR enteric fever. The clinically significant bloodstream isolates of this study [*Salmonella* spp. (*Salmonella enterica* serotypes *typhi* and *paratyphi A*) and *Brucella* spp.] were different Enterobacteriaceae, *S. aureus* and CNS isolated from those documented in large general and university teaching hospitals [4–6, 17–20]. The probable reasons for this difference may be due to the population mix, because our hospital is a specialized hospital dealing with only infectious disease patients. Our present observation is similar to our previous reports [7–9].

In our series, 26.5% (299/1,129) of total isolates were judged to be contaminants or from cases of transient bacteremia without clinical significance, a finding similar to previous studies [20, 21]. Of the contaminants, CNS were the predominant isolates. Several reports have documented the increasing importance of CNS as bloodstream pathogens [4, 5, 21–23], but in the present study, none of the CNS isolates met the clinical criteria to be included as cause of BIs.

The analysis of antimicrobial susceptibility patterns of *S. enterica* serotype *typhi* and *paratyphi A* indicated that a considerable number of enteric fever cases in our hospital are caused by MDR strains. The percentage of MDR strains during the study period remained between 23 and 48%. Panigrahi et al. [7] in an earlier study from our hospital had reported that 45% of the isolates of *S. typhi* between January 1993 and December 1994 were MDR but there were no MDR strains among *S. paratyphi A* [8]. The percentage of MDR *S. typhi* has decreased from 45 to 23% in 2001. Increased resistance of *S. enterica* serotype *typhi* and *paratyphi A* to quinolones, mainly ciprofloxacin, has been widely reported [24, 25]. In the earlier reports from our hospital no such observation was made [7, 8]. However, in the present study, we have noted that the percent-

age of reduced susceptibility of these two *Salmonella* species to ciprofloxacin (MIC > 0.125 µg/ml) [26] had increased up to 60% in 2002. All the *Brucella* isolates demonstrated susceptibility to tetracycline, gentamicin, amikacin, streptomycin and ciprofloxacin. The susceptibility pattern has remained the same as reported by us earlier [9]. In the present study other members of the family Enterobacteriaceae (*E. coli*, *Klebsiella* spp.) showed good in vitro susceptibility to imipenem, ceftriaxone, ceftaxime, ciprofloxacin and gentamicin. Resistance to aminopenicillins and first-generation cephalosporins was very high. This observation is similar to that of Decousser et al. [5].

Clinically significant gram-positive bacteremic episodes were very few in our present study. All the 12 isolates of *S. aureus* were methicillin-susceptible. This find-

ing differs from that of Decousser et al. [5] where 36% of the isolates of *S. aureus* were methicillin-resistant. In the present series, all the 12 cases were probably community-acquired infections, hence the difference. Two of the 5 isolates of *S. pneumoniae* (40%) showed intermediate susceptibility to penicillin, a trend which needs to be closely monitored.

Conclusion

Salmonella enterica serotypes *typhi* and *paratyphi A* were among the predominant clinically significant blood isolates in our hospital, and the number of these strains with reduced susceptibility to ciprofloxacin is increasing, thereby making the monitoring of these isolates essential.

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