



Microbiology of Post-Cardiac Surgery Infections in Children with Congenital Heart Diseases, A Single-Center Experience, Mashhad, Iran

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Abstract

Background: Children who have undergone cardiac surgeries due to congenital heart disease are prone to various kinds of infections.

Objectives: This study was done to investigate the prevalence of nosocomial infections and microbiology of post-cardiac surgery infections in pediatric patients with congenital heart disease (CHD).

Methods: In this cross-sectional study, the epidemiology and microbiology of post-cardiac surgery for pediatric patients with CHD at Imam Reza Hospital of Mashhad University of Medical Sciences between 2014 and 2017 were investigated. Demographic and clinical information was recorded, and the findings were analyzed using SPSS 16.

Results: Out of 1128 patients with open heart surgery during the four years of the study, 135 patients, including 80 males (60.1%) and 55 females (39.9%) with a mean age of 8.06 ± 3.86 months, were enrolled in the study. The prevalence of infection was 11.96%. The most common isolated bacteria were *Acinetobacter* (19/135, 14.1%), *Pseudomonas* spp. (13/135, 9.6%), and *Enterobacter* (13/135, 9.6%) as Gram-negative ones and *Corynebacterium diphtheria* (10/135, 7.4%) and *Staphylococcus epidermidis* (10/135, 7.4%) as Gram-positive types. *Candida albicans* (14/135, 10.4%) were also the most frequent fungi. The frequency of infection-causing masses did not differ significantly between different cardiac abnormalities ($P = 0.831$), sex ($P = 0.621$), age ($P = 0.571$), and weight ($P = 0.786$) groups. Also, the duration of hospitalization, intubation, bypass time, and urinary catheterization in positive culture cases were significantly longer than in negative cases.

Conclusions: In our study, the most common infections in children who underwent heart surgery were *Acinetobacter*, *C. albicans*, *Pseudomonas*, and *Enterobacter*. It is suggested to reduce the hospitalization, intubation, bypass, and urinary catheterization time to reduce nosocomial infections in these patients and decrease treatment costs.

Keywords: Cardiac Surgery, Children, Congenital Heart Disease, Nosocomial Infectious, Prevalence

1. Background

Congenital heart disease is the most common type of genetic disorder in children (1-3). The prevalence of congenital heart disease at birth has been reported to be between 6 and 13 subjects per 1000 live births (4-6), and more than half of these patients need surgery during the first year of life. These children are predisposed to more infections and more severe ones. In addition, some types of congenital heart disease are associated with genetic disorders that cause levels of immunodeficiency and increased susceptibility to infection (7); thus, these surgeries, in turn,

increase the risk of nosocomial infections due to ulcers (8). Although post-surgical infections are well-known complications, different causative germs depend on the type and site of cardiac surgery. Post-cardiac surgery infections are responsible for the high rate of morbidity, prolonged ICU stay, and even mortality; therefore, having a comprehensive epidemiological knowledge of responsible germs makes antibiotic prophylaxis more accurate and effective in these kinds of surgeries. In this study, we tried to find a microbiologic pattern on germs responsible for post-cardiac surgery infections.

Several studies have examined postoperative infec-

tions in children undergoing heart surgery as the most common postoperative infections in surgical sites, with a prevalence of 1.2 to 48% (9-12). The difference in the prevalence rate may happen because of the difference in factors affecting the infectious events, such as body mass index (BMI), cardiopulmonary bypass time, the surgery duration, the age of less than one month, blood transfusion, intravenous feeding, central venous catheters indwelling time, severe bleeding 24 hours after surgery, and misuse scheduling of antibiotics before surgery (7, 9, 13, 14). Common microbial agents that are found at the site of surgery-related infections are *Staphylococcus aureus* (15), *Coagulase-negative staphylococci* (16), *Escherichia coli* (17), *Enterococcus faecalis* (18), and *Pseudomonas* spp. (19), respectively. In the last decade, the number of Gram-negative bacilli has decreased, and the proportion of infections associated with *S. aureus* has increased (20).

In spite of the importance of congenital heart disease and the higher risk of complications and mortality rate, few studies have evaluated the prevalence and factors affecting the disease and its treatment methods and complications (21, 22). Although few studies on surgical infections in children with congenital heart disease in developed and developing countries, these studies have shown the high prevalence of infection in these surgical procedures (7). However, these studies have some limitations, like a small number of samples and errors in retrospective data collection (21). Therefore, identifying the most common causes of infections can help prevent nosocomial infectious diseases. Accordingly, the costs and the length of hospitalization can be decreased, and more importantly, the indiscriminate use of broad-spectrum antibiotics and, consequently, antibiotic resistance can be reduced. Furthermore, monitoring the disinfection of surfaces and equipment and checking the principles of infection control cause a significant reduction in the frequency of such infections.

2. Objectives

The higher susceptibility to severe infection causes the increased risk of nosocomial infection and consequent surgery complications (7). Therefore, additional studies are needed to identify pathogens and the risk factors affecting the disease to improve preventive procedures. In this regard, the present study was conducted to evaluate the type of microbial agents inducing post-cardiac surgery infection and also the possible differences between these infections in terms of age, sex, disease types, and hospitalization length.

3. Methods

3.1. Study Population

In this cross-sectional study, the records of all patients who underwent cardiac surgery for congenital and structural heart diseases during four years (2014 - 2017) in Imam Reza training hospital affiliated with the Mashhad University of Medical Sciences were reviewed. Demographic information included age (between less than one month and 36 months), sex, and weight. Medical data, including hospitalization time, underlying congenital heart disease, history of nosocomial infection, location of nosocomial infection, and the type of microbial infection in the laboratory samples (sputum, blood, urine, and surgical wound) were collected using medical documents.

The nosocomial infections included ventilator-related infections, pneumonia, sepsis, urinary tract infections, infections at the surgical wound site, and cardiovascular, skin, and soft tissue infections.

All the patients with congenital, structural, and valvular heart diseases who underwent palliative or total reconstructive surgical correction in eight of the pediatric cardiac surgery intensive care units in our center were included in this study.

The study was approved by the Mashhad University of Medical Sciences Ethics Committee with the ethical approval number: IR.MUMS.MEDICAL.REC.1397.356. All study procedures followed Helsinki ethics, and a coding system was used to keep the patients' information confidential.

3.2. Statistical Analysis

Data were analyzed using SPSS V16. A P-value less than 0.05 was considered significant. Descriptive indices of mean, standard deviation, frequency, and percentage were used to describe the variables. The chi-square test and *t*-test (or its non-parametric equivalent) were used to compare qualitative and quantitative data.

4. Results

A total of 135 patients were enrolled in the study (Figure 1), of whom 80 patients (60.1%) were male, and 55 patients (39.9%) were female. The mean age of patients was 8.06 ± 3.86 months. Most patients (90/135, 66.7%) were under one month old. Most patients had ventricular septal defect (VSD) (45/135, 32.9%), atrial septal defect (ASD) (32/135, 21.9%), and atrioventricular septal defect (AVSD) (22/135, 10.5%). Ventilator-related infections were among the most reported nosocomial infections (36%), followed by catheter-related urinary tract infections (29%), infections at the site of the surgical wound (16%), sepsis (8%), and skin and soft tissue infections (11%).

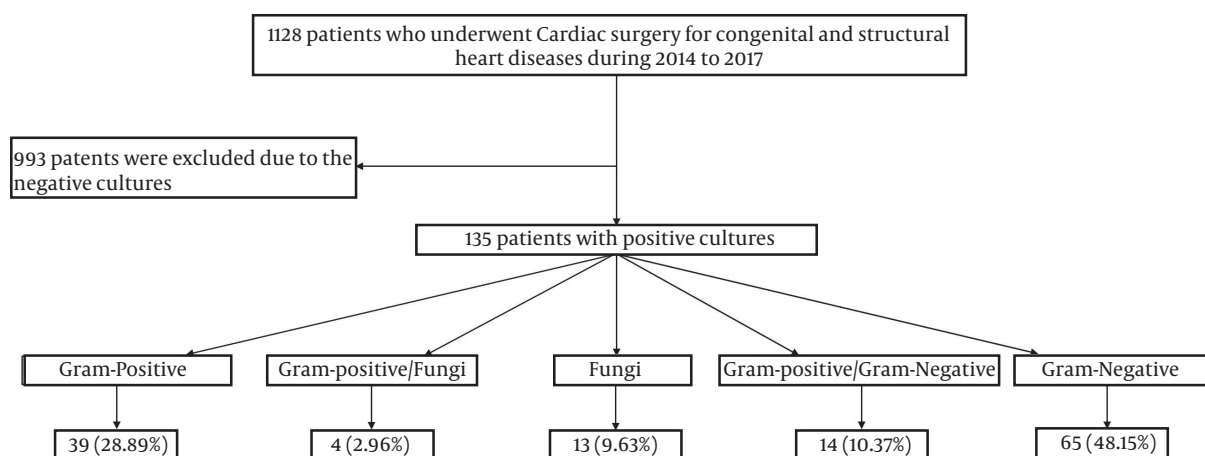


Figure 1. The study population characteristics

Regarding the isolated microorganisms' distribution, the most common Gram-negative types were *Acinetobacter* (14/135, 14.1%), followed by *Pseudomonas* spp. (13/135, 9.6%) and *Enterobacter* (13/135, 9.6%). Also, *Corynebacterium diphtheria* (10/135, 7.4%), *Staphylococcus epidermidis* (10/135, 7.4%), and *S. aureus* (10/135, 7.4%) were the most frequent Gram-positive types. Moreover, *Candida albicans* (14/135, 10.4%) was identified as the most common fungi. The type and frequency of all identified bacterial strains are reported in Table 1.

Comparing group differences based on age revealed that the most positive cultured subjects were under ten months of age, specifically one-month-old. Among 19 positive cultures of *Acinetobacter*, 15 cases were in subjects under one month of age, and four cases were in the second group (1 to 5 months). Moreover, the number of *C. albicans* was higher (10 out of 14) in the first group (one-month-old) than in others. However, there was no significant difference in infectious bacteria between different age groups ($P = 0.571$). The frequency of isolated infectious bacteria in age groups is reported in Table 2.

Analysis of the distribution of bacterial type between sex groups indicated that the prevalence of *Klebsiella*, vancomycin-resistant enterococci, *E. coli*, methicillin-resistant *S. aureus*, methicillin-resistant *S. aureus*, Coagulase-negative staphylococci, and Gram-positive bacilli was higher in males. Also, *Enterococcus*, *C. albicans*, Gram-negative bacilli, and *Strophomonas maltophilia* had equal prevalence in both groups. Although the frequency of other bacteria was higher in females, there was no significant difference between different sex groups regarding the bacteria causing the infection ($P = 0.621$) (Table 3).

Regarding the isolated bacteria in different diseases,

Candida was the most frequently isolated bacteria in ASD subjects. Moreover, *Acinetobacter* was the most abundant bacteria in VSD and AVSD patients. However, there was no significant difference in terms of the bacteria causing infection between different congenital diseases ($P = 0.831$) (Table 4).

Weight-based categorization indicated that 13 subjects with weight less than 5 kg, 45 patients with a weight of 5 to 10 kg, and 77 cases with more than 10 kg had positive cultures. The most abundant bacterium in the less than 5 kg group was *S. epidermidis*. The three most common bacteria in the 5 to 10 kg group were *Acinetobacter*, *S. aureus* (alone and with *Enterobacter*), and *Enterococcus*, respectively. The most common bacteria in the third group (> 10 kg) were *Acinetobacter*, *Candida*, *Pseudomonas*, and *Corynebacterium diphtheria*. There was no significant difference in the type of bacteria between the three weight groups ($P = 0.786$) (Table 5).

The duration of hospitalization ($P = 0.002$), intubation time ($P = 0.004$), bypass time ($P = 0.006$), and duration of urinary catheterization ($P > 0.0001$) were significantly longer in positive cultures than in negative ones (Table 6).

5. Discussion

This study was designed to identify microorganisms that caused post-cardiac surgery infection in these patients. In this regard, 135 patients who underwent heart surgery were enrolled the study. Most of the subjects were infants, and the most common abnormalities were VSD, followed by ASD and PDA. *Acinetobacter*, *C. albicans*, *Pseudomonas* spp., *Enterobacter*, *C. diphtheria*, and *S. epidermidis* were identified as the most common infectious bacteria.

Table 1. Study Population Characteristics and Identified Bacterial Types

Characteristic	No. (%)
Age	
< 1 month	90 (66.7)
2 - 5	26 (19.3)
6 - 10	7 (5.2)
11 - 15	3 (2.2)
16 - 20	0 (0)
21 - 25	3 (2.2)
26 - 30	4 (3.0)
≥ 31	2 (1.5)
Sex	
Male	80 (60.1)
Female	55 (39.9)
Hospitalization unit	
ICUA	3 (1.9)
ICUB	62 (48.6)
ICUC	2 (1.0)
ICUD	2 (1.0)
NICU	9 (6.7)
open-heart surgery ICU	44 (31.4)
Children ward	2 (1.0)
open-heart surgery ward	11 (8.6)
Final diagnosis	
VSD	45 (32.9)
ASD	32 (21.9)
AVSD	22 (10.5)
PDA	8 (7.7)
TOF	6 (5.6)
Cardiomyopathy	6 (5.6)
PS	4 (4.0)
PH	4 (4.0)
TGA	1 (1.0)
CAD	1 (1.0)
COA	1 (1.0)
TAPVC	1 (1.0)
Lupus+ cardiac tamponade	1 (1.0)
Heart failure	3 (2.8)
Bacteria type	
<i>Acinetobacter</i>	19 (14.1)
<i>Candida albicans</i>	14 (10.4)
<i>Pseudomonas spp.</i>	13 (9.6)
<i>Enterobacter</i>	13 (9.6)
<i>Corynebacterium diphtheriae</i>	10 (7.4)
<i>Staphylococcus epidermidis</i>	10 (7.4)
Methicillin-resistant <i>Staphylococcus epidermidis</i>	8 (5.9)
<i>Klebsiella</i>	7 (5.2)
Hemolytic <i>Staphylococcus</i>	7 (5.2)
<i>Enterococcus</i>	6 (4.4)
<i>Escherichia coli</i>	4 (3.0)
Multidrug resistant <i>Acinetobacter</i>	3 (2.2)
Gram-negative bacilli	2 (1.5)
<i>Stenotrophomonas maltophilia</i>	2 (1.5)
Vancomycin-resistant <i>Enterococcus</i>	1 (0.7)
Methicillin-resistant <i>Staphylococcus aureus</i>	1 (0.7)
<i>Micrococcus</i>	1 (0.7)
Coagulase-negative staphylococci	1 (0.7)
Gram-positive bacilli	1 (0.7)

Abbreviations: VSD, ventricular septal defect; ASD, atrial septal defect; AVSD, atrioventricular septal defect.

Our findings showed no significant difference between age, sex, weight, and disease types, and bacterial species. However, the duration of hospitalization, intubation, bypass time, and urinary catheterization was significantly different between positive and negative cultures.

The prevalence of post-cardiac surgery infection in children was reported at 11.96% in our study. Several examinations have studied the distribution of nosocomial infection in children. However, a vast variation was observed in prevalence reported by different studies ranging from 8.7% to 17.7% (23-26). In a new survey on a total of 11,651 subjects (0 - 10 years old), the rate of nosocomial infection was reported to be 10.8% (9). However, another study reported an extremely high frequency of the infection, almost 36% in 155 cases under consideration (13).

On the other hand, a study in the UK on 3090 consecutive pediatric cardiac surgical admissions revealed the post-surgical infection in 0.9 to 2.9% of subjects, independently and along with other morbidities, respectively, that was a lower frequency than previous studies (27). These variations can be explained by individual characteristics in considered populations, like genetic background and probable higher susceptibility to infectious diseases. Also, environmental factors, like infection control protocols and antiseptic materials used in hospitals, may be different. Therefore, evaluation of health issues, attention to disinfection of equipment, and the infection control disciplines lead to a significant reduction in the prevalence of the infection.

In the present study, the most common infectious bacteria in infants were *Acinetobacter*, *Candida albicans*, *Enterobacter*, *Pseudomonas spp.*, *C. diphtheria*, and *S. epidermidis*. Similarly, a previous study reported *S. epidermidis*, *S. aureus*, and *Klebsiella* as blood-borne infectious pathogens, and methicillin-resistant staphylococci, methicillin-susceptible staphylococci, *C. diphtheria*, *E. faecalis*, and *Escherichia coli* as infectious microorganisms at the surgical site (24). Other studies have also indicated the higher prevalence of *S. epidermidis*, *Klebsiella*, *Pseudomonas*, *Acinetobacter*, *S. pneumonia*, *Beta hemolytic streptococcus*, *Candida*, and *H. influenza* as main infectious post-cardiac surgery agents in children (23, 25, 28, 29). Although the identified infectious organisms were similar between studies, the rate and rank of pathogens were different.

Based on our results, bacterial type distribution was similar between different age, sex, and heart disease groups. Although the infection rate in patients under one month old was higher than in others, no difference was observed between age groups. Conversely, previous studies have reported age as an independent risk factor for surgical site infections after pediatric cardiovascular surgery (30, 31). Also, another study indicated that the younger age

Table 2. Distribution of Bacteria Based on Age

Bacteria Type	Age; No. (%)							Total
	< 1 Month	2-5	6-10	11-15	21-25	26-30	≥ 31	
<i>Corynebacterium diphtheriae</i>	7 (7.8)	1 (3.8)	2 (28.6)	0	0	0	0	10 (7.4)
<i>Staphylococcus epidermidis</i>	7 (7.8)	1 (3.8)	2 (28.6)	0	0	0	0	10 (7.4)
Methicillin-resistant <i>Staphylococcus epidermidis</i>	5 (5.6)	1 (3.8)	1 (14.3)	0	0	0	1 (50.0)	8 (5.9)
<i>Klebsiella</i>	4 (4.4)	2 (7.7)	0	0	0	1 (25.0)	0	7 (5.2)
<i>Acinetobacter</i>	15 (16.7)	4 (15.4)	0	0	0	0	0	19 (14.1)
<i>Enterococcus</i>	5 (5.6)	0	0	0	0	0	1 (50.0)	6 (4.4)
Vancomycin-resistant <i>Enterococcus</i>	1 (1.1)	0	0	0	0	0	0	1 (0.7)
<i>Pseudomonas</i> spp.	6 (6.7)	5 (19.2)	1 (14.3)	0	0	1 (25.0)	0	13 (9.6)
<i>Enterobacter</i>	9 (10.0)	2 (7.7)	0	2 (66.7)	0	0	0	13 (9.6)
Hemolytic <i>Staphylococcus</i>	4 (4.4)	1 (3.8)	0	1 (33.3)	1 (33.3)	0	0	7 (5.2)
<i>Candida albicans</i>	10 (11.1)	2 (7.7)	1 (14.3)	0	0	0	1 (50.0)	14 (10.4)
<i>Escherichia coli</i>	4 (4.4)	0	0	0	0	0	0	4 (3.0)
Methicillin-resistant <i>Staphylococcus aureus</i>	1 (1.1)	0	0	0	0	0	0	1 (0.7)
Gram-negative bacilli	1 (1.1)	1 (3.8)	0	0	0	0	0	2 (1.5)
Multidrug resistant <i>Acinetobacter</i>	1 (1.1)	0	0	0	1 (33.3)	0	0	3 (2.2)
<i>Stenotrophomonas maltophilia</i>	2 (2.2)	0	0	0	0	0	0	2 (1.5)
<i>Micrococcus</i>	1 (1.1)	0	0	0	0	0	0	1 (0.7)
Coagulase-negative staphylococci	0	1 (3.8)	0	0	0	0	0	1 (0.7)
Gram-positive bacilli	0	0	1 (14.3)	0	0	0	0	1 (0.7)

Table 3. Distribution of Bacteria Based on Sex

Bacteria Type	Sex; No. (%)	
	Male	Female
<i>Corynebacterium diphtheriae</i>	4 (7.3)	6 (7.5)
<i>Staphylococcus epidermidis</i>	4 (7.3)	6 (7.5)
Methicillin-resistant <i>Staphylococcus epidermidis</i>	1 (1.8)	7 (8.8)
<i>Klebsiella</i>	4 (7.3)	3 (3.8)
<i>Acinetobacter</i>	3 (5.5)	16 (20.0)
<i>Enterococcus</i>	3 (5.5)	3 (3.8)
Vancomycin-resistant <i>Enterococcus</i>	1 (1.8)	0
<i>Pseudomonas</i> spp.	6 (10.9)	7 (8.8)
<i>Enterobacter</i>	5 (9.1)	8 (10.0)
Hemolytic <i>Staphylococcus</i>	1 (1.8)	6 (7.5)
<i>Candida albicans</i>	7 (12.7)	7 (8.8)
<i>Escherichia coli</i>	3 (5.5)	1 (1.3)
Methicillin-resistant <i>Staphylococcus aureus</i>	1 (1.8)	0
Gram-negative bacilli	1 (1.8)	1 (1.3)
Multidrug resistant <i>Acinetobacter</i>	1 (1.8)	2 (2.5)
<i>Stenotrophomonas maltophilia</i>	1 (1.8)	1 (1.3)
<i>Micrococcus</i>	0	1 (1.3)
Coagulase-negative staphylococci	1 (1.8)	0
Gram-positive bacilli	1 (1.8)	0

Table 4. Distribution of Bacteria Based on Disease Type

Bacteria type	Disease Type; No. (%)														
	ASD	AVSD	CAD	COA	COM	HF	L+CT	PDA	PH	PS	TAPVC	TGA	TOF	VSD	
<i>Corynebacterium diphtheriae</i>	2(6.3)	1(4.5)	0	0	1(16.7)	0	0	0	1(25.0)	0	0	0	1(16.7)	3(6.7)	
<i>Staphylococcus epidermidis</i>	3(9.4)	1(4.5)	0	0	1(16.7)	0	0	0	2(50.0)	0	0	0	0	0	
<i>Staphylococcus epidermidis</i> + <i>Corynebacterium diphtheriae</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1(2.2)	
<i>Staphylococcus epidermidis</i> + <i>Candida albicans</i>	1(3.1)	0	0	0	0	0	0	0	0	0	0	0	0	0	
Methicillin-resistant <i>Staphylococcus epidermidis</i>	1(3.1)	1(4.5)	0	0	1(16.7)	0	0	0	0	0	0	0	0	3(6.7)	
Methicillin-resistant <i>Staphylococcus epidermidis</i> + <i>Enterobacter</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1(2.2)	
Methicillin-resistant <i>Staphylococcus epidermidis</i> + <i>Pseudomonas</i> spp.	0	1(4.5)	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Klebsiella</i>	2(6.3)	0	0	0	1(16.7)	0	0	0	0	1(25.0)	0	0	0	2(4.4)	
<i>Klebsiella</i> + <i>Enterobacter</i>	0	0	0	0	0	0	0	0	0	1(25.0)	0	0	0	0	
<i>Klebsiella</i> + <i>Pseudomonas</i> spp.	0	1(4.5)	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Acinetobacter</i>	3(9.4)	5(22.7)	0	0	0	0	0	0	1(25.0)	2(50.0)	0	0	1(16.7)	6(13.3)	
<i>Acinetobacter</i> + <i>Pseudomonas</i> spp.+ <i>Enterobacter</i>	0	0	0	0	0	0	0	1(12.5)	0	0	0	0	0	0	
<i>Enterococcus</i>	0	2(9.1)	0	1(100.0)	0	1(33.3)	0	0	0	0	0	0	0	1(2.2)	
<i>Enterococcus</i> + <i>Candida albicans</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1(2.2)	
Vancomycin-resistant <i>Enterococcus</i>	0	0	0	0	0	0	0	0	0	0	0	0	1(16.7)	0	
<i>Pseudomonas</i> spp.	2(6.3)	3(13.6)	0	0	0	0	0	3(37.5)	0	0	0	0	0	3(6.7)	
<i>Enterobacter</i>	2(6.3)	1(4.5)	1(100.0)	0	0	0	0	1(12.5)	0	0	0	0	0	4(8.9)	
<i>Hemolytic Staphylococcus</i>	0	2(9.1)	0	0	0	1(33.3)	0	1(12.5)	0	0	0	0	0	2(4.4)	
<i>Hemolytic Staphylococcus</i> + <i>Enterobacter</i>	0	1(4.5)	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Candida albicans</i>	8(25.0)	0	0	0	1(16.7)	0	0	0	0	0	0	0	0	3(6.7)	
<i>Escherichia coli</i>	0	1(4.5)	0	0	0	0	0	0	0	0	0	0	0	3(6.7)	
Methicillin-resistant <i>Staphylococcus aureus</i>	0	0	0	0	0	0	0	1(12.5)	0	0	0	0	0	0	
Gram-negative bacilli	0	0	0	0	0	0	0	1(12.5)	0	0	0	0	0	0	
Gram-negative bacilli+ <i>Staphylococcus epidermidis</i>	1(3.1)	0	0	0	0	0	0	0	0	0	0	0	0	0	
Multidrug resistant <i>Acinetobacter</i>	2(6.3)	1(4.5)	0	0	0	0	0	0	0	0	0	0	0	0	
<i>Stenotrophomonas maltophilia</i>	1(3.1)	0	0	0	1(16.7)	0	0	0	0	0	0	0	0	0	
<i>Micrococcus</i>	0	0	0	0	0	0	0	0	0	0	0	0	1(16.7)	0	
Coagulase-negative staphylococci	0	0	0	0	0	0	0	0	0	0	0	0	1(16.7)	0	
Gram-positive bacilli	0	0	0	0	0	0	0	0	0	0	0	0	0	1(2.2)	
Negative culture	4(12.5)	1(4.5)	0	0	0	1(33.3)	1(100.0)	0	0	1(25.0)	1(100.0)	1(100.0)	1(16.7)	11(24.4)	

Abbreviations: VSD, ventricular septal defect; ASD, atrial septal defect; AVSD, atrioventricular septal defect.

is related to more susceptibility to attract infections (32). However, sex was not a risk factor for infection (9). Furthermore, we found that prolonged hospitalization, intubation, bypass, and urinary catheterization time was more significant in patients with positive cultures. Consistent with these findings, it has previously been reported that a longer duration of surgery, prolonged ICU stay, catheter-in-dwelling time, and longer previous hospital stay were associated with nosocomial post-cardiac surgery infections (13, 30, 31).

As one of the limitations, this study was limited to one center, which reduced the sample size. Furthermore, the prevalence of post-cardiac surgery infection has not been

evaluated compared with other types of surgeries based on the age distribution. However, our study is one of the few studies in this area, especially to examine the age distribution and distribution of infectious masses among different types of congenital heart disease. The results of this study can be used in two parts. First, identifying the bacteria causing the infection in children who have had heart surgery can improve our knowledge about antibiotics that could be used for empirical treatment or as prophylaxis. Also, the distribution of infectious bacteria at different ages can help to prescribe these antibiotics more accurately. Second, this study can add more evidence to the literature about the distribution of pathogens in different

Table 5. Distribution of Bacteria Based on Weight

Bacteria Type	Weight (kg); No. (%)		
	< 5	5 - 10	> 10
<i>Corynebacterium diphtheriae</i>	1 (7.7)	1 (2.2)	8 (10.4)
<i>Staphylococcus epidermidis</i>	2 (15.4)	1 (2.2)	5 (6.5)
<i>Staphylococcus epidermidis</i> + <i>Corynebacterium diphtheriae</i>	0	0	1 (1.3)
<i>Staphylococcus epidermidis</i> + <i>Candida albicans</i>	0	0	1 (1.3)
Methicillin-resistant <i>Staphylococcus epidermidis</i>	1 (7.7)	0	5 (6.5)
Methicillin-resistant <i>Staphylococcus epidermidis</i> + <i>Enterobacter</i>	0	1 (2.2)	1 (2.2)
Methicillin-resistant <i>Staphylococcus epidermidis</i> + <i>Pseudomonas spp.</i>	0	1 (2.2)	1 (1.3)
<i>Klebsiella</i>	1 (7.7)	0	5 (6.5)
<i>Klebsiella</i> + <i>Enterobacter</i>	0	0	1 (1.3)
<i>Klebsiella</i> + <i>Pseudomonas spp.</i>	0	0	1 (1.3)
<i>Acinetobacter</i>	0	11 (24.4)	10 (12.9)
<i>Acinetobacter</i> + <i>Pseudomonas spp.</i> + <i>Enterobacter</i>	1 (7.7)	1 (2.2)	1 (1.3)
<i>Enterococcus</i>	1 (7.7)	6 (13.3)	0
<i>Enterococcus</i> + <i>Candida albicans</i>	1 (7.7)	1 (2.2)	1 (1.3)
Vancomycin-resistant <i>Enterococcus</i>	0	1 (2.2)	0
<i>Pseudomonas spp.</i>	1 (7.7)	3 (6.6)	8 (10.4)
<i>Enterobacter</i>	1 (7.7)	5 (11.1)	3 (3.9)
<i>Hemolytic Staphylococcus</i>	0	0	3 (3.9)
<i>Hemolytic Staphylococcus</i> + <i>Enterobacter</i>	1 (7.7)	5 (11.1)	1 (1.3)
<i>Candida albicans</i>	0	3 (6.7)	10 (12.9)
<i>Escherichia coli</i>	1 (7.7)	2 (4.4)	1 (1.3)
Methicillin-resistant <i>Staphylococcus aureus</i>	0	0	1 (1.3)
Gram-negative bacilli	0	0	1 (1.3)
Gram-negative bacilli+ <i>Staphylococcus epidermidis</i>	1 (7.7)	1 (2.2)	1 (1.3)
Multidrug resistant <i>Acinetobacter</i>	0	1 (2.2)	4 (5.2)
<i>Stenotrophomonas maltophilia</i>	0	1 (2.2)	1 (1.3)
<i>Micrococcus</i>	0	1 (2.2)	0
Coagulase-negative staphylococci	0	0	1 (1.3)
Gram-positive bacilli	0	0	1 (1.3)
Total	13 (100)	45 (100)	77 (100)

Table 6. Comparison of Hospitalization, Intubation, Bypass, and Urinary Catheterization Time Between Positive and Negative Cases

Characteristics	Culture		P-Value
	Positive	Negative	
Hospitalization time (day)	6.34 ± 3.21	4.05 ± 2.44	0.002
Intubation time (h)	14.28 ± 5.14	10.43 ± 4.31	0.004
Bypass time (min)	130.3 ± 10.4	91.5 ± 2.8	0.006
Urinary catheterization time (day)	13.16 ± 1.8	3.1 ± 0.3	< 0.0001

age categories and various heart diseases, as previous researches have paid less attention to these features.

5.1. Conclusion

The prevalence of nosocomial infection in open-heart surgery in our patients was 11.96%. The duration of hospitalization, intubation, bypass, and urinary catheterization was significantly higher in positive cultures than negative ones. Therefore, decreasing the time of these situations may reduce nosocomial infection. Consequently, treatment costs, as well as resistance to antibiotics, can be reduced. However, more studies should be conducted with a larger sample size focusing on age and microbial type distribution among different heart diseases to confirm these findings.

Footnotes

Authors' Contribution: Behzad Alizadeh, administrative; Ghasem Dolatkah, acquisition of data; Hossein Akhavan, statistical analysis; Hasan Birjandi, drafting of the manuscript; Mohammad Reza Naghibi Sistani, analysis and interpretation of data; Hasan Motaghi, study supervision.

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