ORIGINAL RESEARCH

Effectiveness of Pre-Transplant Screening for High-Priority Multidrug-Resistant Pathogens on Pre-Engraftment Infections After Hematopoietic Stem Cell Transplantation

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Objective: Owing to the rising incidence of multidrug-resistant organisms (MDRO) and the high mortality rates associated with such bacterial infections post-hematopoietic stem cell transplantation (HSCT), we investigated the MDRO colonization rate prior to transplantation using an active surveillance approach and determined its impact on subsequent infection during the pre-engraftment period.

Methods: A single-center observational study was conducted, and surveillance cultures from multiple body sites, including the rectum, nasal cavity, and groin, were performed at admission to determine MDRO colonization. Serological tests were used to detect certain viruses and toxoplasmosis before HSCT.

Results: In the pre-transplant setting, 59 MDRO were recovered from the 40 HSCT recipients. Of the 59 isolates recovered from one or more body sites, 29 were positive for methicillin-resistant *Staphylococcus aureus* (MRSA), 7 for carbapenem-resistant Enterobacterales (CRE), and 23 were positive for extended-spectrum β -lactamase (ESBLs). Serological assessment before HSCT revealed active or reactivation of latent infection with cytomegalovirus (7.5%), Epstein–Barr virus (EBV; 5%), and *Toxoplasma gondii* (2.5%) among HSCT patients. In terms of factors associated with pre-engraftment infections, the type of transplant (*p*=0.04) was statistically significant, whereas other factors, such as age, sex, and underlying conditions, were not. In post-transplant settings, bloodstream infections (BSIs) were documented in 2 allogeneic HSCT patients (5%), and the isolated microorganisms were ESBL-producing *E. coli* and non-MDR *Acinetobacter baumannii*.

Conclusion: Active screening cultures are a helpful tool for identifying patients colonized by MDRO or relevant viruses before HSCT, and for predicting those at risk of developing subsequent pre-engraftment infections. Additionally, active screening may aid in predicting those who are likely to develop subsequent pre-engraftment infections. Our findings highlight the importance of pre-transplant screening for high-priority multidrug-resistant pathogens and the application of infection control interventions after HSCT. **Keywords:** pre-transplant screening, MRSA, ESBLs, CRE, pre-engraftment infections

Introduction

To replace and repopulate the hematopoietic system, hematopoietic stem cell transplant (HSCT), previously known as bone marrow transplant, has been used as a potentially curative treatment for several life-threatening malignant diseases and immunological disorders over the last decades.¹ Before stem cell infusion, patients undergo a preparative regimen

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(myeloablative conditioning regimen) that involves high-dose chemotherapy and/or total body irradiation to eliminate the numerous malignant cells and to prevent rejection of the transplanted graft.² As a result of the interplay between weakened defense mechanisms induced by conditioning regimens and opportunistic pathogens, life-threatening infections with bacteria, especially multidrug-resistant organisms (MDRO), as well as the acquisition of respiratory viral infections, and/or reactivation of human herpes viruses remains a major determinant of morbidity and mortality after allogeneic or autologous HSCT.^{3–6} Additionally, reactivation of the Hepatitis B virus (HBV),⁷ Hepatitis C virus (HCV),⁸ and latent toxoplasmosis⁹ remain a risk exacerbated by the immunosuppressive status of HSCT recipients.

Carbapenem-resistant Enterobacterales (CRE), MDR-*Pseudomonas aeruginosa*, MDR-*Acinetobacter baumannii*, extended-spectrum β -lactamase producers (ESBLs), and methicillin-resistant *Staphylococcus aureus* (MRSA) currently account for the majority of hospital-acquired infections (HAIs) within 30 days after HSCT in the pre-engraftment phase,^{10,11} data on HSCT-related infections remain limited in developing countries. During the pre-engraftment neutropenic phase, the dysbiosis of endogenous gastrointestinal flora, mucosal damage with indwelling devices, and prior colonization with MDRO can contribute to different clinical manifestations, including bloodstream infections (BSIs), pneumonia and gastrointestinal infections.^{12,13} Despite major advances in the management of post-transplant outcomes, the incidence ranges from 13–46% and 5–10% among allogeneic HSCT, in which patients receive progenitor cells from the donor^{13,14} and autologous HSCT, in which the patient receives their own progenitor cells,¹⁵ respectively, with a mortality rate of 32.7%.¹⁶

Although active surveillance strategies to detect the colonization state of HSCT recipients are important tools for controlling and preventing the spread of MDRO within hospital settings and may be useful in guiding empirical therapy for post-transplantation neutropenia, their impact on patient outcome and cost-effectiveness remains controversial,¹⁷ particularly in developing countries where MDRO is endemic. Accordingly, this study aimed to evaluate the effect of pre-transplant screening for high-priority MDRO and relevant viruses on the outcome of pre-engraftment infections. We also studied the effect of infection control policies on the rate of HAIs during the pre-engraftment period after post-HSCT.

Materials and Methods

Study Design

This prospective observational cohort study was conducted on adult patients who underwent HSCT between January 2021 and December 2021 at the International Medical Center (IMC), Cairo, Egypt. The bone marrow transplant (BMT) unit at the IMC contains four highly isolated and super-equipped units for autologous and allogeneic transplantations of hematological cancers. Data of the enrolled patients, including demographics, underlying diseases, and microbiological data, both pre- and post-HSCT, were retrieved from the hospital's electronic medical record system. The study procedure was in accordance with the Declaration of Helsinki and approved by the Institutional Ethics Committee of the Faculty of Pharmacy, Ain Shams University (ACUC-FP-ASU RHDIRB2020110301 REC #72).

Pre-Transplant and Post-Transplant (Pre-Engraftment) Screening

Within 48 hours of admission, all patients who underwent BMT were subjected to routine CRE, ESBLs, and MRSA pretransplant screening. Rectal swabs were used for CRE/ESBL screening, whereas swabs from the anterior nares, groin, and axilla were used for MRSA screening. Additionally, extensive screening covering different body sites, including the eye, ear, sputum, urine, and blood, adhering to international guidelines <u>https://apps.who.int/iris/rest/bitstreams/911060/retrieve</u> (accessed on February 5, 2024) was performed.

HB&L-carbapenemase[®] and ESBL-producers-AmpC[®] kits (Alifax, Padua, Italy) were used for active screening of CRE and ESBLs producers from rectal swabs, respectively. The patented laser beam technology with dedicated reagents (selective supplement broth that contains carbapenem and certain third-generation cephalosporins), enables us to perform rapid bacterial culture and to rapidly screen CRE and ESBLs with high sensitivity and specificity.¹⁸ For MRSA screening/isolation, an oxacillin-resistant screening agar base medium was used (Oxoid, UK) and a cefoxitin disk (30 µg) was used to identify MRSA isolates as per clinical laboratory standard institute.¹⁹ The automated system Vitek-2 (BioMérieux, Marcy L'Etoile; Lyon, France) was used for the further confirmation and detection of antimicrobial

breakpoints, in accordance with the manufacturer's recommendations. Polymerase chain reaction (PCR) was used to determine the type of carbapenemase enzyme and ESBLs as previously reported.¹⁸

Serological tests were used to determine prior (IgG) or recent (IgM) exposure to a multitude of pathogens, including certain viruses (human herpes virus type 1 and 2, CMV, EBV, HCV, human immunodeficiency virus (HIV)), and parasites such as *Toxoplasma gondii* which are prevalent among HSCT recipients. HBsAg, anti-HBc IgM, and total anti-HBc serum markers were tested for HBV. An Architect Plus immunoassay analyzer (Abbott, Illinois, USA) was used to deliver reliable results without delay and to enhance the laboratory workflow, as previously mentioned. The Architect anti-HCV assay was used for qualitative detection of anti-HCV antibodies in human serum and plasma. The chemiluminescent signal in the reaction was compared to the cutoff signal obtained from Architect Anti-HCV calibration to determine the presence or absence of IgG/IgM anti-HCV in sample https://www.corelaboratory.abbott/int/en/offerings/segments/infectious disease/hepatitis.html (accessed on 5 February 2024).

The Architect HIV Ag/Ab combo assay was used for the detection of HIV, based on the simultaneous qualitative detection of HIV-p24-Ag and Ab to HIV-1/HIV-2. <u>https://www.ilexmedical.com/files/PDF/HIVAgAbCombo.pdf</u> (accessed on 5 February 2024).

To exclude primary infections with CMV, CMV IgG reactive samples (detected by Architect CMV IgG assay) should be tested for CMV-IgM and CMV-IgG Avidity as previously reported <u>https://www.ilexmedical.com/files/PDF/CMVIgM_ARC.pdf</u>; (accessed on 5 February 2024).

A rapid antigen test was used for coronavirus disease 2019 (COVID-19) screening and positive results were confirmed by PCR whenever indicated according to the Centers for Disease Control and Prevention (CDC) standard guidelines (<u>https://www.cdc.gov/coronavirus/2019-ncov/lab/resources/antigen-tests-guidelines.html</u>) (accessed on February 5 2024).²⁰ For monitoring post-transplant infections, patients with fever or with any other signs or symptoms of infections were monitored through the collection of two blood culture sets in addition to other diagnostic procedures based on the patient's clinical presentation, as a part of follow-up care during the pre-engraftment period (https://apps.who.int/iris/rest/bitstreams/911060/retrieve) (accessed on February 5, 2024).

Microbiological Definition

MDR phenotype is defined as acquired non-susceptibility to at least one agent in three or more antimicrobial categories.²¹ Colonization is defined as the presence of the respective organism in at least one surveillance culture. BSIs are defined as the isolation of the microorganism from at least one set of blood cultures associated with systemic signs of infection, such as fever, chills, and hypotension, when drawn from a transplantation recipient at any phase of transplantation.²² In cases of skin commensals, BSIs are defined by the identification of at least two consecutive blood cultures of the same species isolated within 24 hours, as reported by the Centers of Disease Control and Prevention (CDC) (https://www.cdc.gov/nhsn/pdfs/pscmanual/4psc_clabscurrent.pdf); (accessed on February 5, 2024).

Infection Prevention and Control

Cornerstones of infection control procedures, including routine hand hygiene, environmental cleaning, contact precautions, antimicrobial stewardship programs along with using intense aseptic/antiseptic regimens were routinely applied to stop transfer of MDRO within BMT unit (https://apps.who.int/iris/rest/bitstreams/911060/retrieve) (accessed on February 8, 2024). Except for those colonized with MDRO Gram-negative bacteria, all patients received levofloxacin (500 mg/day) as a prophylaxis from the beginning of the conditioning regimen until engraftment, while colonized individuals by MDRO received a regimen based on the isolate's susceptibility profile.²³ Acyclovir was used as a routine antiviral prophylactic agent for all HSCT recipients, while ganciclovir was only used for CMV-related indications among high-risk HSCT recipients.²⁴ Fluconazole was used as an antifungal prophylactic agent^{25,26} and posaconazole was used among allogeneic patients with acute myeloid leukemia or of high risk to develop graft versus host disease.^{27,28} Trimethoprim-sulfamethoxazole was used as a prophylactic agent to prevent both toxoplasmosis and *Pneumocystis jirovecii* pneumonia.²⁹ A decolonization protocol was used to reduce asymptomatic MRSA carriers, but it was not used in the case of ESBLs and CRE due to concerns about the emergence of resistant strains and negative effects on the gut microbiome.²³ For MRSA decolonization regimen, the body was washed with chlorhexidine gluconate 4% on a daily basis for 5 days. To determine the clearance of

MRSA, a post-decolonization screening step was conducted by reswabbing different multibody sites to ensure the effectiveness of decolonization. <u>https://www.cdc.gov/infectioncontrol/pdf/strive/MRSA202-508.pdf</u> (accessed on 5 February 2024).

Association Between Colonization State and Pre-Engraftment Infections

Before transplantation, we determined the colonization state of recipients by initial screening for MDRO including MRSA,ESBLs and CRE. After HSCT, we examined the association between prior colonization with MDRO and probability of developing subsequent infections with the same MDR pathogens at pre-engraftment period.

Statistical Data Analysis

All analyses were conducted using the GraphPad Prism 8.0 (GraphPad Software, San Diego, CA, USA). Chi-square or Fisher's exact test was used to compare the proportions of categorical variables, and p < 0.05 was considered statistically significant.

Results

Patient Demography and History

During the study period, forty patients received HSCT procedures, (15, 37.5%) allogeneic transplant and (25, 62.5%) autologous transplantation. The median patient age was 43 years (range: 34.5–55), 25 patients were males (62.5%) and 15 (37.5%) were females. The most common underlying condition was multiple myeloma (11, 27.5%), followed by Hodgkin Lymphoma (10, 25%) and acute myeloid leukemia (6, 15%).

Patient Surveillance Prior to HSCT

Among the 40 patients, 280 surveillance cultures (2 rectal swabs, n = 80; 1 swab from nostrils, axilla, and groin, n = 40 each) were performed per patient to detect the colonization state by CRE, ESBLs, and MRSA upon admission to the hospital prior to HSCT (Table 1). Other multi-body site colonization screening, including eye, ear, and urine, was performed. Of the 40 patients, 59 MDRO, including 29 MRSA, seven CRE, and 23 ESBLs, were isolated from one or more body sites. Most MRSA isolates were recovered from groins (n = 19), followed by axial (n = 16) and nasal swabs (n = 8). Of the 40 patients screened for MRSA, 4 had results assessable by all three sites (axilla, groin, and nostrils), 9 had results assessable by two combined sites (axilla and groin). On the other hand, 6, 4, and 3 patients were assessable using only a single site, including groin, nasal, and axillary swabs, respectively.

For viral/parasitic serological assessment prior to HSCT, CMV (3, 7.5%), EBV (2, 5%), and *Toxoplasma gondii* (1, 2.5%) were recorded, whereas none of the other tested viruses were detected.

Distribution of MDRO Pre- and Post-HSCT and Antibiotic Resistance Profile

As shown in Figure 1, the distribution of MDRO pre-autologous HSCT revealed that MRSA colonization accounted for approximately 55%, followed by ESBLs at 42%, and CRE at 3%. In the pre-allogeneic HSCT group, MRSA colonization

	Surveillance MDRO culture- samples Recovered from Multi-body sites ^A							Viral/parasite serological assessment ^B positive samples		
	rectum	nasal	axilla	groin	eye	sputum	urine	vagina		
CRE	4	Not Tested Not Tested			0	I	2	0	Cytomegalovirus (lgM)	3
ESBLs	17				0	0	4	2	Epstein Barr virus (IgM)	2
MRSA	Not Tested	8	16	19	2	0	0	I	Toxoplasma gondii (lgM)	I

Table I Data on MDRO Recovered from Surveillance Cultures and Viral/Parasitic Serostatus of Patients Admitted to the HSCT Unit

Notes: ^AMulti-body sites tested included the eye, ear, sputum, urine, blood, anus, and vagina. ^BVirology screening included HSV-1, HSV-II, CMV, EBV, HBsAg, HBcAb, HCV, HIV, Toxoplasma, and rapid testing for COVID-19.

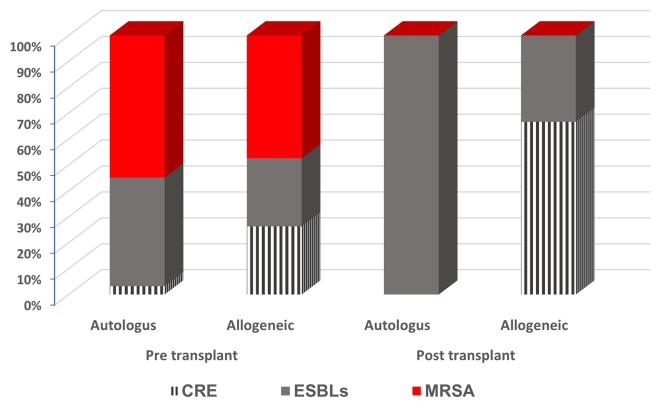


Figure I Distribution of Multidrug-resistance organisms (MDRO) pre- and post-HSCT.

accounted for approximately 47.4%, followed by ESBLs and CRE (26.3%). In post-autologous HSCT, 100% of the recovered MDROs were ESBLs, whereas in post-allogeneic HSCT, ESBLs accounted for 33% and CRE for approximately 67%.

The antibiotic resistance pattern of CRE post-HSCT showed reduced susceptibility to carbapenems (75–100%), cephalosporins (100%), aminoglycosides (50–75%), quinolones (100%), and trimethoprim/sulfamethoxazole (100%). For β -lactam/ β -lactamase inhibitors, 100% of the recovered CRE were resistant to ampicillin/sulbactam and 50% were resistant to ceftazidime/avibactam. On the other hand, the antibiotic resistance pattern of ESBLs post-HSCT revealed a high resistance pattern towards cephalosporins (80–100%), quinolones (100%), and trimethoprim/sulfamethoxazole (100%). In contrast, the recovered ESBLs were 100% sensitive to carbapenems and aminoglycosides (Figure 2).

Statistical Association Between Colonization State and Pre-Graft Infections

As shown in Table 2, before transplantation, there was no statistical difference between colonized patients with MDRO and non-colonized patients with regard to age, sex, underlying conditions, and type of transplant. However, there was a statistically significant difference between the type of transplant and pre-engraftment infections caused by MDRO after transplantation (p = 0.04), whereas the other three factors were not statistically significant.

Pre-Transplant Colonization Rate by MDR Pathogens, Relevant Viruses, and Parasites

The pre-transplant colonization rates of MRSA, ESBLs, and CRE were 26 (65%), 18 (45%), and 6 (15%), respectively, as shown in Table 3. Among all the colonized patients in the cohort, only one patient who was previously colonized by MDR *E. coli* expressing the ESBL phenotype and another patient colonized by *Acinetobacter baumannii* subsequently developed BSIs with the same isolate during the pre-engraftment period. Notably, the majority of patients colonized by MDRO prior to transplantation and after applying infection control interventions did not develop subsequent infections post-HSCT.

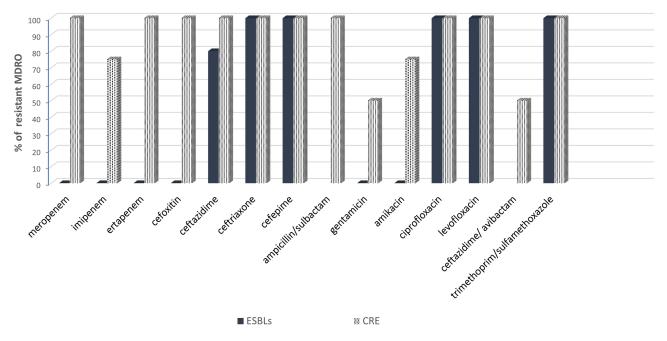


Figure 2 Antimicrobial resistance profile of CRE and ESBLs post-HSCT.

Regarding viral and parasitic serostatus, only one allogeneic HSCT recipient had active CMV infection during the pre-engraftment phase and was negative prior to transplantation. The overall mortality rate during the pre-engraftment phase was 10%, and four patients who died had undergone allogeneic HSCT (p=0.006), which could be related to central line-associated BSIs, as confirmed in one patient. The microbiological profiles of these 4 patients revealed that one patient had CLA BSI caused by MDR *E. coli*. In contrast, one patient was infected with MDR *A. baumannii*, another patient was infected with non-MDR *K. pneumoniae* and the last patient had coagulase-negative staphylococci. Notably, these organisms did not lead to HAIs as defined by the CDC.

Patient demographics/	Pre-tran	splant colonization	Post-transplant (pre-engraftment period)			
clinical characteristics	Group I: Colonized by MDRO (N=32)	Group 2: Not colonized by MDRO (N= 8)	p-value	Group I: Infected by MDRO (N= 9)	Group 2: Not infected by MDRO (N=31)	p-value
Gender						
Male	18	7	0.10	6	19	0.76
Female	14	I		3	12	
Age						
>30 years	6	0	0.18	2	6	0.84
<31 years	26	8		7	25	
Underlying disease						
Multiple myeloma	8	3	0.77	I	10	0.36
R-Hodgkin lymphoma	9	I		2	8	
Acute myeloid leukemia	5	I		I	5	
Others ^A	10	3		5	8	
Type of transplant						
Autologous	19	6	0.41	3	22	0.04*
Allogeneic	13	2		6	9	

Table 2 Demographic Data of Enrolled Patients Pre- and Post-HSCT

Notes: Others ^Achronic myeloid leukemia, acute myeloid leukemia, and relapsed/refractory diffuse large B-cell lymphoma. *means significant (p value< 0.05).

Microbiological profile	N (%) of patients colonized by surveillance culture/other body sites	N (%) of patients colonized and subsequent infection with the same isolate type post-HSCT	Type of hospital- acquired infection	N (%) patients colonized and after applying infection control interventions they did not develop subsequent infections post-HSCT	
		Gram-negative bacilli			
ESBLs: E. coli and K. pneumoniae.	18 (45%)	I (5.5%)	BSI caused by E. coli	17 (94.5%)	
CRE: E. coli and K. pneumoniae	6 (15%)	0	0	6 (100%)	
A. baumannii	3(7.5%)	l (33.3%)	BSI	2(66.7%)	
Pseudomonas spp.	3(7.5%)	0	0	3(100%)	
		Gram-positive cocci			
MRSA	26 (65%)	0	0	26 (100%)	
Staphylococcus aureus I (2.5%)		0	0	I(100%)	
Coagulase-negative staphylococci	11(27.5%)	0	0	11(100%)	

 Table 3 Relationship Between Pre-Transplant Screening and Post-Transplant Infections Among HSCT Recipients

Notes: Staphylococcus aureus and methicillin resistant Staphylococcus aureus by oxacillin-resistant screening agar base medium (Oxoid, UK) and a cefoxitin disk (30 µg) as recommended by clinical laboratory standard institute.

Correlation Between Colonization State and Pre-Engraftment Infections

Table 4 shows the correlation between colonization state by MDRO prior to transplantation and infection by the same MDRO post-transplantation, and their significant association. Statistically significant results (p = 0.037) were observed between the pre-transplant colonization state with certain MDRO (ESBLs, CRE, MRSA) and the development of subsequent infection during the pre-engraftment period. Notably, prior colonization with CRE showed a significant association (p=0.000) with CRE post-transplant infections.

Discussion

Infections caused by MDRO, especially gram-negative bacteria, in addition to reactivation of *Herpesviridae* family, have a significant impact on the prognosis of immunocompromised patients post-HSCT. For the management of such infections, understanding microbial colonization prior to transplant through surveillance programs, along with serological

Groups	Number of transplant by	P value		
Number of patients infected post-transplant by MDRO as ESBLs, CRE, and MRSA		Yes	No	0.037*
	Yes	5	4	
	No	27	4	
Number of patients infected post-transplant by CRE		Yes	No	0.000*
	Yes	3	I	
	No	3	33	
Number of patients infected post-transplant by ESBLs		Yes	No	0.810
	Yes	2	3	
	No	16	19	
Number of patients infected post-transplant by MRSA		Yes	No	1.00
	Yes	0	0	
	No	26	14	

Table 4 Association Between Colonization State by MDRO Prior to Transplant and Infection by Same MDRO Post-Transplant

Note: *means significant (p value< 0.05).

investigation of certain viruses, might be crucial to initiate preventive strategies among HSCT recipients.³⁰ In this study, we used an extensive screening strategy to investigate the rate of colonization by MDRO from multi-body sites as well as to analyze the association between colonization and subsequent infection by the same pathogen post-HSCT.

Pre-transplant screening revealed a high prevalence of MRSA colonization (65%) among the admitted patients, indicating that its spread within community settings should not be underestimated. To increase the sensitivity of MRSA screening, groin and axillary swabs were screened in addition to nares in our cohort. Among the 26 patients colonized by MRSA, 73%, 61.5%, and 30.7% were identified using groin, axilla, and nasal swabs, respectively. In contrast to our findings, other studies have shown that the anterior nares are the sites most frequently colonized by MRSA, ^{31,32} reflecting disagreement over the ideal anatomical site for MRSA sampling. Several factors, including the nature of swabs, decolonization protocol, and laboratory testing tools, can contribute to the variation in MRSA colonization detection rate among HSCT patients.

The rising threat of infections caused by MDR Gram-negative bacteria among HSCT recipients has been documented in the centers, but data on the consequences of these infections remain limited in our region. The prevalence rates of CRE and ESBLs colonization in our study cohort were 15% and 45%, respectively. The rate of colonization by CRE was higher than that recently reported from a center in Turkey,³³ while rate of ESBLs was in tune with a study conducted to predict bacteremia among hematological patients.³⁴ Such variations in colonization rates among HSCT recipient assure that understanding local epidemiology particularities may have a significant impact on outcomes.

Serological testing was used as a pre-transplant screening indicator to determine the serostatus of certain viral/parasitic infections. Our results revealed that three, two, and one HSCT patients had CMV, EBV and Toxoplasmosis active or reactivation of latent infection, respectively. The presence of only one case of active CMV infection during the pre-engraftment period and the absence of other viral infections, despite the COVID-19 pandemic, indicate the effectiveness of prophylactic antiviral therapy and the enhanced pre-transplant serological clearing that covered a wide array of viruses.

In our cohort, the type of transplant was associated with an increased risk of infection during the pre-engraftment phase (p = 0.04), whereas other factors, including age, sex, and underlying conditions, were not statistically significant. This could be attributed to the rate of immune reconstitution, which is typically faster in autologous HSCT recipients than in allogeneic HSCT recipients, who continue to have measurable deficits in cell-mediated and humoral immunity, even after engraftment, as reported in other studies.^{35,36}

Prospective monitoring for subsequent infections with the same MDRO post-transplant revealed the absence of MRSA and CRE and low rates of ESBLs (5.5%) in our center. Although there is limited evidence that pre-transplant MRSA or ESBLs colonization predicts subsequent infection among adult recipients post-HSCT, there is still a significant risk for infections among patients colonized by CRE.³⁷ The success of the management strategy for preventing MRSA infections in our cohort could be attributed to early screening of MRSA, contact precautions, and decolonization protocol as reported in other studies.^{38,39} In the context of infections by ESBLs and CRE, their management was even more complicated due to disruption of the gut microbiome leading to predominance of gram-negative MDRO⁴⁰ and neutropenia that predispose to bacterial infections⁴¹ and invasive fungal infections during the pre-engraftment period.⁴² However, the low rate of ESBLs and absence of CRE infections post-HSCT supports the effectiveness of multimodal infection control interventions in our cohort, in addition to the tailoring of empirical antibiotic treatment depending on the previously known colonization state.

Within 30 days after HSCT, BSIs were documented in 2 patients (5%), and the isolated microorganisms were ESBLproducing *E. coli* and non-MDR *Acinetobacter baumannii*. Our findings were similar to those reported by the Tunisian BMT center (5.9%).⁴³ In total, four patients (10%) died during the pre-engraftment period after allogeneic HSCT, and mortality was related to BSIs caused by ESBL-producing *E. coli* in one patient. Our results are similar to those of other studies that reported infection-related mortality as a major hurdle among allogeneic HSCT recipients, particularly during the early period of transplantation.^{44,45}

A recent study conducted in 2023 showed that the emergence of MDR pathogens as well as patient characteristics should be considered for better management of bacteremia in patients receiving allogeneic HSCT.⁴⁶ Another study revealed that a short course of antibiotics for 7 days might be sufficient in patients with Cancer and Hematopoietic Stem Cell Transplantation suffering from infections with gram-negative bacteria.⁴⁷ Our findings were in accordance with those

of Goloshchapov et al, who demonstrated the correlations between colonization of EBV positivity in the colon and delayed hematopoietic reconstitution.⁴⁸

In previous studies, pre-screening and multimodal infection control approaches for the highly priority pathogens have been confirmed to avoid and combated hospital-acquired infection caused by CRE and ESBLs producers.^{18,49} Moreover, it is recommended to use advanced and accurate methods such as Biofire FilmArray panel or Vitek-system for identifying the colonized pathogens in order to define the proper preventive measure for each pathogen as previously reported.^{49,50.} The relatively small sample size and analysis of single-center data might limit the generalizability of our study. However, this is one of the few studies in regions that has determined the impact of MDRO screening prior to transplantation on subsequent infections with the same isolate during the pre-engraftment period.

Conclusion

This study highlights the effectiveness of pre-transplant screening for high-priority multidrug-resistant pathogens and the application of infection control interventions after HSCT. The majority of patients colonized by MDRO before transplantation and after applying infection control interventions did not develop subsequent infections post-HSCT. Prior to transplantation, there was no statistical difference between colonized patients with MDRO and non-colonized patients with regard to age, sex, underlying conditions, or transplant type. However, there was a statistically significant difference between the types of transplant and pre-engraftment infection. Therefore, understanding the colonization state is critical for guiding infection control measures in HSCT recipients.

Abbreviations

BSIs, bloodstream infections; BMT, Bone marrow transplant; CRE, Carbapenem-resistant Enterobacterales; CDC, Centers for Disease Control and Prevention; COVID-19, Coronavirus disease 2019; CMV, Cytomegalovirus; EBV, Epstein–Barr virus; ESBLs, extended-spectrum β-lactamase; HBV, Hepatitis B virus; HCV, Hepatitis C virus; HIV, Human immunodeficiency virus; HAIs, Hospital-acquired infections; IMC, International medical center; MRSA, methicillin-resistant *Staphylococcus aureus*; MDR, multidrug-resistant; MDRO, multidrug-resistant organism; HSCT, posthematopoietic stem cell transplant; PCR, Polymerase chain reaction.

Data Sharing Statement

All data generated or analyzed during this study are included in this published article in the main manuscript.

Ethics Approval and Consent to Participate

This study was conducted in accordance with the Declaration of Helsinki, and was reviewed and approved by the Faculty of Pharmacy, Ain Shams University Research Ethics Committee (ACUC-FP-ASU RHDIRB2020110301 REC #72). Informed consent was obtained from all subjects and/or their legal guardians (s).

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Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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Disclosure

The authors declare that they have no competing interests in this work.

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