

Study of Species Distribution and Antifungal Susceptibility of *Candida* Isolates from Bloodstream Infection in the ICU of a Tertiary Care Hospital of Rajasthan

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Abstract:

Introduction: *Candida* bloodstream infections (candidemia) are an important cause of healthcare-associated infections in intensive care units (ICUs), contributing to significant morbidity, mortality, prolonged hospitalization, and increased healthcare costs. The increasing predominance of non-*Candida albicans* species and the emergence of antifungal resistance have made early species identification and antifungal susceptibility testing essential for effective patient management.

Materials and Methods: A cross-sectional study was conducted in which 240 blood culture samples were collected from patients admitted to various ICUs over an 8 month period (from October 2025 to May 2026), were processed for isolation and identification of *Candida* species using conventional microbiological methods, CHROMagar *Candida*, cornmeal agar morphology, germ tube test, and confirmation by the VITEK 2 Compact system. Antifungal susceptibility testing was performed by the Kirby–Bauer disk diffusion method following CLSI guidelines.

Results: Among the 240 blood culture samples, 27 (11.3%) yielded *Candida* species. Non-*Candida albicans* species accounted for 77.7% of isolates, while *Candida albicans* comprised 22.3%. Amphotericin B demonstrated the highest susceptibility (81.4%), followed by nystatin (62.96%), whereas fluconazole showed the lowest susceptibility (33.33%).

Conclusion: The study demonstrates a shift towards predominance of non-*albicans* *Candida* species with increasing resistance to azole antifungal agents among ICU patients. Early diagnosis, species-level identification, and routine antifungal susceptibility testing are essential for timely initiation of appropriate therapy, optimizing patient outcomes, and reducing the morbidity and mortality associated with candidemia.

Keywords: *Candida* bloodstream infection; candidemia; *Candida tropicalis*; antifungal susceptibility; fluconazole resistance; non-*Candida albicans*.

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Introduction

Candida species are commensal yeasts that colonize the skin, gastrointestinal tract, genitourinary tract, and mucosal surfaces of healthy individuals. Under conditions of impaired host immunity or disruption of normal microbial flora, these opportunistic fungi can invade the bloodstream and disseminate to multiple organs, resulting in invasive candidiasis. The incidence of candidemia has increased steadily over the past few decades and currently represents one of the most common causes of nosocomial bloodstream infections worldwide. [1,2]

Although *Candida albicans* has traditionally been the predominant cause of invasive *Candida* infections. There has been a gradual epidemiological

shift toward non-*albicans* *Candida* (NAC) species in recent years. This changing epidemiology is clinically important because different *Candida* species exhibit varying antifungal susceptibility profiles. [3,4] Several non-*Candida albicans* species possess intrinsic or acquired resistance to commonly used azole antifungal agents. [5]

The risk factors for *Candida* bloodstream infection (BSI) include ICU admission with prolonged hospital stay, central venous line insertion, urinary catheterization, antibiotic overuse, diabetes mellitus, immunosuppression. [6,7]

There are different laboratory methods for identification of the various *Candida* species. Conventional identification methods, including germ tube testing, CHROMagar *Candida*, cornmeal agar morphology, carbohydrate assimilation tests, and automated identification systems such as the VITEK 2 Compact, improve diagnostic accuracy and facilitate timely clinical management.

Materials and Methods

Study Design and Setting: A cross-sectional observational study was performed in the Department of Microbiology, S.P. Medical College, Bikaner, Rajasthan, over a period of eight months (from October 2025 to May 2026). Under aseptic conditions, Blood culture samples were collected from patients admitted to various ICUs.

Identification of *Candida* Species: Blood samples were inoculated into BacT/ALERT blood culture bottles and incubated at 37°C. Positive blood culture bottles were subcultured onto Blood agar and MacConkey agar using the streak plate method followed by incubation at 37°C for 24 hours. Gram staining was performed directly from the positive blood culture broth and from the isolated colonies.

Species identification was performed using a combination of conventional methods and an automated identification system. Identification included Gram staining, germ tube test, colony morphology on CHROMagar *Candida*, and microscopic morphology on cornmeal agar with Tween 80. Final species confirmation was carried out using the VITEK 2 Compact automated identification system.

Antifungal Susceptibility Testing: Antifungal susceptibility testing was performed by the Kirby–Bauer disk diffusion method on Mueller–Hinton agar supplemented with 2% glucose and 0.5 µg/mL methylene blue in accordance with Clinical and Laboratory Standards Institute (CLSI) guidelines. The antifungal agents tested included fluconazole, ketoconazole, itraconazole, amphotericin B, and

nystatin. Zone diameters were interpreted according to CLSI breakpoints.

Results

A total of 240 blood culture samples from ICU patients with suspected bloodstream infection were processed during the study period. *Candida* species were isolated from 27 (11.3%) samples, while the remaining 213 (88.7%) samples were negative for *Candida*.

Out of the 27 patients with candidemia, 17 (62.96%) were males and 10 (37.04%) were females. The highest number of cases was observed in the 0–10 years age group (37.0%), followed by the 51–60 years age group, indicating that both paediatric and elderly ICU patients were at increased risk of candidemia.

Among the 27 *Candida* isolates, *Candida tropicalis* 11 (40.7%) was the most common species, followed by *Candida albicans* 6 (22.2%), *Candida krusei* 5 (18.5%), *Candida parapsilosis* 3 (11.1%), and *Candida glabrata* 2 (7.4%). (Table 1 and Fig :1). Non-*Candida albicans* species predominated, accounting for 21 (77.8%) isolates, whereas *Candida albicans* constituted 6 (22.2%) isolates. Thus, *Candida tropicalis* was the predominant bloodstream isolate among ICU patients.

The most frequently identified risk factors were prior use of broad-spectrum antibiotics (85.2%) and urinary catheterization (85.2%), followed by prolonged ICU stay (77.8%), recent surgery (59.3%), central venous catheterization (48.1%), corticosteroid therapy (44.4%), diabetes mellitus (33.3%), and total parenteral nutrition (25.9%).

Antifungal susceptibility testing demonstrated that amphotericin B showed the highest in vitro susceptibility (81.4%), followed by nystatin (62.96%). In contrast, fluconazole exhibited the lowest susceptibility (33.33%), indicating reduced susceptibility among *Candida* bloodstream isolates. Variable susceptibility was observed with itraconazole and ketoconazole. (Table 2 and Fig :2)

Table 1: Distribution of *Candida* species isolated from bloodstream infections (n=27)

Candida Species	Number of Isolates	Percentage
<i>Candida albicans</i>	06	22.2%
<i>Candida tropicalis</i>	11	40.7%
<i>Candida krusei</i>	05	18.5%
<i>Candida parapsilosis</i>	03	11.1%
<i>Candida glabrata</i>	02	7.4%

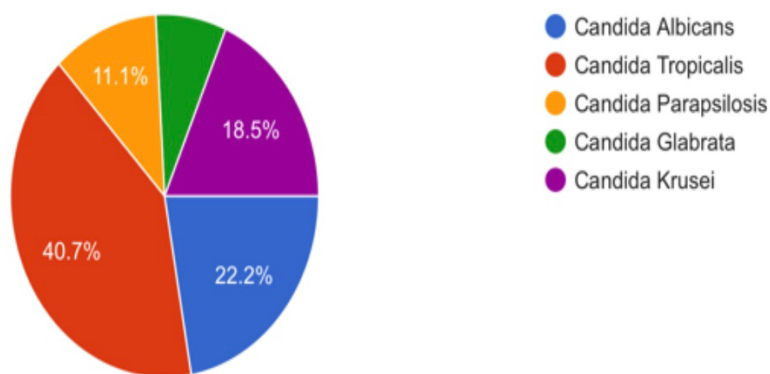


Figure 1: Candida Species Distribution

Table 2: Antifungal Susceptibility pattern among Candida Species isolates by Kirby-Bauer disc diffusion method

Drugs	Sensitivity (%)	Intermediate(%)	Resistance(%)
Fluconazole	33.33%	18.51%	48.14%
Ketoconazole	55.55%	25.92%	18.51%
Itraconazole	37.03%	48.14%	14.81%
Amphotericin B	81.4%	7.4%	11.11%
Nystatin	62.96%	25.92%	11.11%

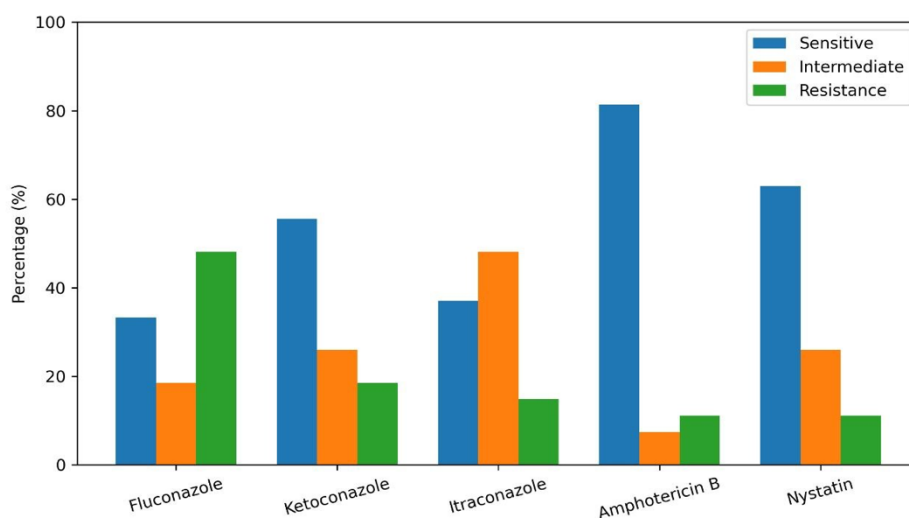


Figure 2: Antifungal Susceptibility pattern among Candida Species

Discussion

Candidemia is a hematogenous infection by the Candida species that predominantly impacts individuals with compromised immune systems, such as those treated with chemotherapy or those diagnosed with HIV/AIDS. Early identification is crucial for proper management and better patient outcomes [8,9]

In the present study, the prevalence of candidemia among blood culture samples was 11.25% (27/240), which is comparable to previous Indian studies Mamta Lamba et al. [16] and Jagdish Chander et al.

[15] who found a prevalence rate of 9.94% and 10.06% respectively.

A male predominance (62.1%) was observed, due to increased exposure of male patients to healthcare-associated risk factors such as prolonged ICU stay, invasive devices, broad-spectrum antibiotic therapy, and underlying comorbidities. [14]

The study demonstrated a clear predominance of non-albicans Candida (77.8%) over Candida albicans (22.2%). Among the non-albicans species, Candida tropicalis was the most frequently isolated species (40.7%). These findings are comparable to Jagdish Chander et al. [15], Tejan et al. [12], R.

Adhikary et al.¹³ and Mamta Lamba et al. [16] which found 40.8%, 42.1%, 39.7% and 43.75% respectively, followed by *Candida krusei*, *Candida parapsilosis*, and *Candida glabrata*.

Several Indian studies have reported a shift toward non-albicans *Candida* species. The increasing predominance of *C. tropicalis* may be associated with prolonged hospitalization, extensive use of broad-spectrum antibiotics, immunosuppressive therapy, invasive procedures, and inadequate infection control practices. [10,11]

The major risk factors identified were urinary catheterization, broad-spectrum antibiotic use, central venous catheterization, steroid therapy, diabetes mellitus. These findings indicate that invasive devices and underlying comorbidities are major predisposing factors for the development of Candidemia. [17]

Antifungal susceptibility testing showed that Amphotericin B (81.4%) was the most effective antifungal agent, whereas Fluconazole exhibited the lowest susceptibility (33.33%). Similar patterns have been reported in R. Adhikary et al. [13], Mamta Lamba et al. [16] reflect the widespread empirical and prolonged use of fluconazole. The increasing resistance among non-albicans *Candida* species highlights the importance of routine antifungal susceptibility testing to guide appropriate therapy. Intrinsic resistance of *Candida krusei* to fluconazole and the reduced azole susceptibility of *Candida glabrata* emphasize the importance of species-level identification before initiating antifungal therapy. [16]

Overall, the findings of the present study confirm the changing epidemiology of candidemia, characterized by the predominance of non-albicans *Candida* species and increasing azole resistance. These observations emphasize the need for early diagnosis, species-level identification, routine antifungal susceptibility testing, and strict infection control practices to improve patient outcomes and optimize antifungal therapy in ICU settings.

Conclusion

The present study demonstrates that non-albicans *Candida* species have emerged as the predominant cause of candidemia, with *Candida tropicalis* being the most frequently isolated species. This changing epidemiological trend highlights the need for routine species-level identification, as different *Candida* species exhibit distinct antifungal susceptibility profiles that significantly influence therapeutic decisions. Amphotericin B showed the highest susceptibility, while Fluconazole exhibited the lowest susceptibility among *Candida* species isolates.

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